

METHODS AND COMPOSITIONS FOR PREDICTING AND/OR MONITORING DIABETES AND TREATMENTS THEREFOR

CROSS REFERENCE TO RELATED APPLICATIONS

5 This application claims the benefit of priority under 35 U.S.C. § 119(e) to U.S. Application No. 63/074,836 filed September 4, 2020. This document is incorporated by reference herein in its entirety.

FEDERALLY SPONSORED RESEARCH OR DEVELOPMENT

10 This invention was made with government support under R01DA037648 and R44DA041014 awarded by the National Institutes of Health. The government has certain rights in the invention.

TECHNICAL FIELD

15 This disclosure generally relates to methods and compositions related to predicting and/or monitoring diabetes in an individual.

BACKGROUND

20 Diabetes (e.g., type 1, type 2, adult onset) is characterized by insulin resistance, and, in some cases, reduced insulin secretion. Many instances of diabetes can be attributed to lifestyle factors, such as obesity (defined by a body mass index of greater than 30), lack of physical activity, poor diet, stress, and urbanization, in combination with genetics.

 While many risk factors for diabetes have been identified, a reliable method of predicting and/or monitoring diabetes is desirable.

SUMMARY

25 Methods and compositions for predicting the incidence of, and/or monitoring the presence or progression of, diabetes are described herein. Specifically, methods and compositions for determining the methylation status of at least one CpG locus and at least one single nucleotide polymorphism (SNP) are described.

In one aspect, kits for determining methylation status of at least one CpG dinucleotide and a genotype of at least one single-nucleotide polymorphism (SNP) are provided. Such a kit generally includes at least one first nucleic acid primer at least 8 nucleotides in length that is complementary to a bisulfite-converted nucleic acid sequence comprising a first CpG dinucleotide at a GC locus selected from the group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312 or a second CpG dinucleotide in linkage disequilibrium with the first CpG dinucleotide, wherein the linkage disequilibrium has a value of $R > 0.3$, wherein the at least one first nucleic acid primer detects a methylated or unmethylated CpG dinucleotide, and at least one second nucleic acid primer at least 8 nucleotides in length that is complementary to a DNA sequence or a bisulfite-converted DNA sequence of a first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855 or a second SNP in linkage disequilibrium with the first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855, wherein the linkage disequilibrium has a value of $R > 0.3$.

In some embodiments, the at least one first nucleic acid primer detects the unmethylated CpG dinucleotide. In some embodiments, the at least one first nucleic acid primer detects the methylated CpG dinucleotide.

In some embodiments, the at least one first nucleic acid primer comprises one or more nucleotide analogs. In some embodiments, the at least one first nucleic acid primer comprises one or more synthetic or non-natural nucleotides.

In some embodiments, the methods described herein further include at least a third nucleic acid primer at least 8 nucleotides in length that is complementary to a nucleic acid sequence upstream of the CpG dinucleotide.

In some embodiments, the kits described herein can further include a solid substrate to which the at least one first nucleic acid primer is bound. In some embodiments, the substrate is a polymer, glass, semiconductor, paper, metal, gel or hydrogel. In some embodiments, the solid substrate is a microarray or microfluidics card.

In some embodiments, the kits described herein further include a detectable label. In some embodiments, the kits described herein further include at least a third nucleic acid primer at least 8 nucleotides in length that is complementary to a nucleic acid sequence downstream of the CpG dinucleotide.

In another aspect, methods of determining the presence of biomarkers associated with diabetes in a biological sample from a patient are provided. Such a method typically includes (a) providing a first portion of the biological sample and a second portion of the biological sample, wherein the nucleic acid from at least the first portion is bisulfite converted; (b) contacting the first portion of the biological sample with a first oligonucleotide primer at least 8 nucleotides in length that is complementary to a sequence that comprises a first CpG dinucleotide at a GC locus selected from the group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312, or a second CpG dinucleotide in linkage disequilibrium with the first CpG dinucleotide at a GC locus selected from the group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312, wherein the linkage disequilibrium has a value of $R > 0.3$, wherein the first nucleic acid primer detects a methylated or unmethylated CpG dinucleotide, wherein the linkage disequilibrium has a value of $R > 0.3$; and (c) contacting the second portion of the biological sample with a nucleic acid primer at least 8 nucleotides in length that is complementary to a DNA sequence or a bisulfite-converted DNA sequence of a first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855 or a second SNP in linkage disequilibrium with the first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855, wherein the linkage disequilibrium has a value of $R > 0.3$. Generally, the percentage of methylation of the CpG dinucleotide at the GC locus selected from the group consisting of cg06500161, cg21139312, cg19693031, cg17901584, and cg13375427 or a second CpG dinucleotide in linkage equilibrium with the first CpG dinucleotide and the identity of the nucleotide in the first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855 or the second SNP in linkage disequilibrium with the first SNP are biomarkers associated with diabetes in the patient.

In still another aspect, methods of determining the presence of a biomarker associated with diabetes in a patient sample are provided. Such methods typically include (a) isolating nucleic acid sample from the patient sample, (b) performing a genotyping assay on a first portion of the nucleic acid sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or a second SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain genotype data; and/or (c) bisulfite converting the nucleic acid in a second portion of the nucleic

acid and performing methylation assessment on a second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R>0.3$) with a CpG from Appendix A to obtain methylation data; and (d) inputting the genotype data from step (b) and/or methylation data from step (c) into an algorithm that accounts for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect, wherein the algorithm is a machine learning algorithm capable of accounting for linear and non-linear effects.

In some embodiments, the at least one interaction effect is selected from the group consisting of a gene-environment interaction (SNPxCpG) effect, a gene-gene interaction (SNPxSNP) effect, and an environment-environment interaction (CpGxCpG) effect. In some embodiments, the at least one interaction effect is a gene-environment interaction effect (SNPxCpG) between a CpG site from Appendix A or a CpG site that is collinear ($R>0.3$) with a CpG site from Appendix A and a SNP from Appendix C or a SNP within moderate linkage disequilibrium ($R>0.3$) from a SNP from Appendix C.

In some embodiments, the at least one interaction effect is an environment-environment interaction effect (CpGxCpG) between at least two CpG sites from Appendix A. In some embodiments, one or both of the at least two CpG sites are collinear ($R>0.3$) with one or both of the at least two CpG sites from Appendix A. In some embodiments, the at least one interaction effect is a gene-gene interaction effect (SNPxSNP) between at least two SNPs from Appendix C. In some embodiments, one or both of the at least two SNPs are collinear ($R>0.3$) with one or both of the at least two SNPs from Appendix C.

In some embodiments, the biological sample is a saliva sample.

In still another aspect, systems for determining methylation status of at least one CpG dinucleotide and a genotype of at least one single-nucleotide polymorphism (SNP) are provided. Such systems typically include a nucleic acid isolation module configured to isolate a nucleic acid sample from a subject sample; a genotyping assay module configured to perform a genotyping assay on a first portion of the nucleic acid sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage disequilibrium ($R>0.3$) with a first SNP from Appendix C to obtain genotype data; a methylation assay module configured to bisulfite convert the nucleic acid in a second portion of the nucleic acid and perform a methylation assessment on a second portion of the nucleic acid sample to detect

methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data; and an identification system configured to account for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on the genotype data from step (b) and/or methylation data from step (c).

In some embodiments, the system further includes an output module configured to provide an output based on an identification by the identification system, wherein the identification accounts for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on the genotype data from step (b) and/or methylation data from step (c).

In some embodiments, the algorithm is a machine learning algorithm capable of accounting for linear and non-linear effects.

In still another aspect, non-transitory computer-readable media storing instructions executable by a processing device to perform operations are provided. Such operations typically include accounting for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on genotype data and/or methylation data, wherein: (i) the genotype data is based on a genotyping assay on a first portion of a nucleic acid sample isolated from a subject sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain the genotype data; and (ii) the methylation data is based on a methylation assay on a bisulfite converted nucleic acid in a second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data.

In some embodiments, the operations further include providing an output based on the accounting. Representative outputs include, without limitation, one or more of storing a report based on the accounting to another non-transitory computer-readable medium, modifying a display based on the accounting, triggering an audible alert based on the accounting, triggering a haptic or vibratory alert based on the accounting, triggering the printing of a report based on the accounting, or triggering the delivery of a therapeutic based on the accounting.

Unless otherwise defined, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which the

methods and compositions of matter belong. Although methods and materials similar or equivalent to those described herein can be used in the practice or testing of the methods and compositions of matter, suitable methods and materials are described below. In addition, the materials, methods, and examples are illustrative only and not intended to be limiting. All publications, patent applications, patents, and other references mentioned
5 herein are incorporated by reference in their entirety.

DESCRIPTION OF DRAWINGS

FIG. 1 is a ROC curve of main effects diabetes classification model.
FIG. 2 is a ROC curve of interaction effects diabetes classification model.
10 FIG. 3 is a block diagram of an exemplary diabetes classification system.
FIG. 4 is a flow diagram of an exemplary process for diabetes classification.
FIG. 5 is a block diagram of exemplary computing devices.

DETAILED DESCRIPTION

The present disclosure provides methods and kits for determining whether a
15 subject has a predisposition to, or likelihood of having or developing diabetes. As shown herein, the methylation status of one or more CpG dinucleotides, alone, or in combination with the genotype and/or the interaction between the genotype and the methylation status (e.g., CH3xSNP), is associated with diabetes.

As described herein, biomarkers described herein can be used in the diagnosis and
20 prognosis of cardiovascular diseases and events. The terms “marker” and “biomarker” can be used interchangeably. As used herein, a biomarker generally refers to a measurable or detectable biological moiety (e.g., the presence or amount of a protein, a genetic and/or histological component). As described in more detail below, the biomarkers used herein typically are associated with diabetes.

25

DNA Methylation

DNA does not exist as naked molecules in the cell. For example, DNA is associated with proteins called histones to form a complex substance known as chromatin. Chemical modifications of the DNA or the histones alter the structure of the
30 chromatin without changing the nucleotide sequence of the DNA. Such modifications are described as “epigenetic” modifications of the DNA. Changes to the structure of the chromatin can have a profound influence on gene expression. If the chromatin is

condensed, factors involved in gene expression may not have access to the DNA, and the genes will be switched off. Conversely, if the chromatin is “open,” the genes can be switched on. Some important forms of epigenetic modification are DNA methylation and histone deacetylation.

5 DNA methylation is a chemical modification of the DNA molecule itself and is carried out by an enzyme called DNA methyltransferase. Methylation can directly switch off gene expression by preventing transcription factors binding to promoters. A more general effect is the attraction of methyl-binding domain (MBD) proteins. These are associated with further enzymes called histone deacetylases (HDACs), which function to
10 chemically modify histones and change chromatin structure. Chromatin-containing acetylated histones are open and accessible to transcription factors, and the genes are potentially active. Histone deacetylation causes the condensation of chromatin, making it inaccessible to transcription factors and causing the silencing of genes.

CpG islands are short stretches of DNA in which the frequency of the CpG
15 sequence is higher than other regions. The “p” in the term CpG indicates that cytosine (“C”) and guanine (“G”) are connected by a phosphodiester bond. CpG islands are often located around promoters of housekeeping genes and many regulated genes. At these locations, the CG sequence is not methylated. By contrast, the CG sequences in inactive genes are usually methylated to suppress their expression.

20 As used herein, the term “methylation status” means the determination whether a certain target DNA, such as a CpG dinucleotide, is methylated or is unmethylated. As used herein, the term “CpG dinucleotide repeat motif” means a series of two or more CpG dinucleotides positioned in a DNA sequence.

About 56% of human genes and 47% of mouse genes are associated with CpG
25 islands. Often, CpG islands overlap the promoter and extend about 1000 base pairs downstream into the transcription unit. Identification of potential CpG islands during sequence analysis helps to define the extreme 5' ends of genes, something that is notoriously difficult with cDNA-based approaches. The methylation of a CpG island can be determined by a skilled artisan using any method suitable to determine such
30 methylation. For example, the skilled artisan can use a bisulfite reaction-based method for determining such methylation.

It would be understood that linkage equilibrium can be quantitated as linkage disequilibrium, using, for example, the Pearson correlation (R) or co-inheritance of alleles

(D'). For example, a low level of linkage can be reflected in a correlation (e.g., R value) of about 0.1 or less, a moderate level of linkage is reflected in a R value of about 0.3, while a high level of linkage is reflected in a R value of 0.5 or greater. It also would be understood that, when referring to methylation (i.e. CpGs), collinearity (with an R value) is used as a determination of the linear strength of the association between two CpGs (e.g., a low level of collinearity can be reflected by an R value of about 0.1 or less; a moderate level of collinearity can be reflected by an R value of about 0.3; and a high level of collinearity can be reflected by an R value of about 0.5 or greater).

The present disclosure provides methods to determine the nucleic acid methylation of one or more loci in a subject in order to predict the clinical course and eventual outcome of subjects suspected of being predisposed or of having diabetes.

Genetic screening (also called genotyping or molecular screening) can be broadly defined as testing to determine if a subject has a genetic marker that either causes a disease state or is "linked" to the genetic component causing the disease state. Linkage refers to the phenomenon that DNA sequences which are close together in the genome have a tendency to be inherited together. Two sequences may be linked because of some selective advantage of co-inheritance. More typically, however, two polymorphic sequences are co-inherited because of the relative infrequency with which meiotic recombination events occur within the region between the two polymorphisms. The co-inherited polymorphic alleles are said to be in "linkage disequilibrium" with one another because, in a given population, they tend to either both occur together or else not occur at all in any particular member of the population. Indeed, where multiple polymorphisms in a given chromosomal region are found to be in linkage disequilibrium with one another, they define a quasi-stable genetic "haplotype." In contrast, recombination events occurring between two polymorphic loci cause them to become separated onto distinct homologous chromosomes. If meiotic recombination between two physically linked polymorphisms occurs frequently enough, the two polymorphisms will appear to segregate independently and are said to be in linkage equilibrium.

It would be understood that linkage disequilibrium can be quantitated (using, for example, the Pearson correlation (R) or co-inheritance of alleles (D')). For example, a low level of linkage can be reflected in a correlation (e.g., R value) of about 0.1 or less, a moderate level of linkage is reflected in a R value of about 0.3, while a high level of linkage is reflected in a R value of 0.5 or greater. It also would be understood that, when

referring to methylation (i.e. CpGs), collinearity (with an R value) is used as a determination of the linear strength of the association between two CpGs (e.g., a low level of collinearity can be reflected by an R value of about 0.1 or less; a moderate level of collinearity can be reflected by an R value of about 0.3; and a high level of collinearity can be reflected by an R value of about 0.5 or greater).

In particular, in certain embodiments of the disclosure, the methods may be practiced as follows. A sample, such as a blood sample, is taken from a subject. In certain embodiments, a single cell type, e.g., lymphocytes, basophils, or monocytes isolated from the blood, may be isolated for further testing. The DNA is harvested from the sample and examined to determine the methylation of one or more loci. For example, the DNA of interest can be treated with bisulfite to deaminate unmethylated cytosine residues to uracil. Since uracil base pairs with adenosine, thymidines are incorporated into subsequent DNA strands in the place of unmethylated cytosine residues during subsequent PCR amplifications. Next, the target sequence is amplified by PCR, and probed with a loci-specific probe. Depending on the particular sequence of the probe used, only the methylated or unmethylated DNA will bind to the probe.

Methods of determining the nucleic acid profile for a subject are well known to a skilled artisan and include any of the well-known detection methods. Various PCR methods are described, for example, in *PCR Primer: A Laboratory Manual*, Dieffenbach 7 Dveksler, Eds., Cold Spring Harbor Laboratory Press, 1995. Other methods include, but are not limited to, nucleic acid quantification, restriction enzyme digestion, DNA sequencing, hybridization technologies, such as Southern Blotting, amplification methods such as Ligase Chain Reaction (LCR), Nucleic Acid Sequence Based Amplification (NASBA), Self-sustained Sequence Replication (SSR or 3SR), Strand Displacement Amplification (SDA), and Transcription Mediated Amplification (TMA), Quantitative PCR (qPCR), or other DNA analyses, as well as RT-PCR, *in vitro* translation, Northern blotting, and other RNA analyses. In another embodiment, hybridization on a microarray is used.

Single Nucleotide Polymorphism (SNP) Genotyping

Traditional methods for the screening of heritable diseases have depended on either the identification of abnormal gene products (e.g., sickle cell anemia) or an abnormal phenotype (e.g., mental retardation). With the development of simple and

inexpensive genetic screening methodology, it is now possible to identify polymorphisms that indicate a propensity to develop disease, even when the disease is of polygenic origin.

Single nucleotide polymorphism (SNP) genotyping measures genetic variations of SNPs between members of a species. A SNP is a single base pair change at a specific locus, usually consisting of two alleles (where the rare allele frequency is >1%). SNPs are very common. Because SNPs are conserved during evolution, they have been proposed as markers for use in quantitative trait loci (QTL) analysis and in association studies in place of microsatellites. Many different SNP genotyping methods are known, including hybridization-based methods (such as Dynamic allele-specific hybridization, molecular beacons, and SNP microarrays) enzyme-based methods (including restriction fragment length polymorphism, PCR-based methods, flap endonuclease, primer extension, 5'-nuclease, and oligonucleotide ligation assay), other post-amplification methods based on physical properties of DNA (such as single strand conformation polymorphism, temperature gradient gel electrophoresis, denaturing high performance liquid chromatography, high-resolution melting of the entire amplicon, use of DNA mismatch-binding proteins, SNPLEX and surveyor nuclease assay), and sequencing (such as "next generation" sequencing). See, e.g., US Patent No. 7,972,779.

A plurality of alleles at a locus can arise from one or more polymorphisms in a region of a gene that encodes a polypeptide or in a regulatory control sequence that affects expression of the polypeptide, such as a promoter or polyadenylation sequence. Alternatively, alleles can arise from one or more polymorphisms at a locus distal to a gene that encodes a polypeptide or in a regulatory control sequence. A polymorphism can affect a polypeptide at a transcriptional or a translational level (e.g., a polypeptide's transcription rate, translation rate, degradation rate, and/or activity). Allelic differences can be characterized in a sample from a single subject or from a plurality of subjects using methods that are known to a skilled artisan. Such methods can include, but are not limited to, measuring the potential for a polynucleotide sequence to be expressed and/or measuring an amount of an encoded polypeptide. Methods are available that can detect proteins or nucleic acids directly or indirectly, and assay methods are specifically contemplated to include screening for the presence of particular sequences or structures of nucleic acids or polypeptides using, e.g., any of various known microarray technologies.

It will be fully appreciated by the skilled artisan that the allele need not have previously been shown to have had any link or association with the disorder phenotype. Instead, an allele and a pathogenic environmental risk factor can interact to predict a predisposition to a disorder phenotype even when neither the allele nor the risk factor bears any direct relation to the disorder phenotype.

Genetic screening (also called genotyping or molecular screening), can be broadly defined as testing to determine if a subject has mutations (or alleles or polymorphisms) that either cause a disease state or are “linked” to the mutation causing a disease state. Linkage refers to the phenomenon that DNA sequences which are close together in the genome have a tendency to be inherited together. Two sequences may be linked because of some selective advantage of co-inheritance. More typically, however, two polymorphic sequences are co-inherited because of the relative infrequency with which meiotic recombination events occur within the region between the two polymorphisms. The co-inherited polymorphic alleles are said to be in “linkage disequilibrium” with one another because, in a given population, they tend to either both occur together or else not occur at all in any particular member of the population. Indeed, where multiple polymorphisms in a given chromosomal region are found to be in linkage disequilibrium with one another, they define a quasi-stable genetic “haplotype.” In contrast, recombination events occurring between two polymorphic loci cause them to become separated onto distinct homologous chromosomes. If meiotic recombination between two physically linked polymorphisms occurs frequently enough, the two polymorphisms will appear to segregate independently and are said to be in linkage equilibrium.

It would be understood that linkage equilibrium / disequilibrium can be quantitated (using, for example, the Pearson correlation (R) or co-inheritance of alleles (D')). For example, a low level of linkage can be reflected in a correlation (e.g., R value) of about 0.1 or less, a moderate level of linkage is reflected in a R value of about 0.3, while a high level of linkage is reflected in a R value of 0.5 or greater.

While the frequency of meiotic recombination between two markers is generally proportional to the physical distance between them on the chromosome, the occurrence of “hot spots” as well as regions of repressed chromosomal recombination can result in discrepancies between the physical and recombinatorial distance between two markers. Thus, in certain chromosomal regions, multiple polymorphic loci spanning a broad chromosomal domain may be in linkage disequilibrium with one another, and thereby

define a broad-spanning genetic haplotype. Furthermore, where a disease-causing mutation is found within or in linkage with this haplotype, one or more polymorphic alleles of the haplotype can be used as a diagnostic or prognostic indicator of the likelihood of developing the disease. This association between otherwise benign polymorphisms and a disease-causing polymorphism occurs if the disease mutation arose in the recent past, so that sufficient time has not elapsed for equilibrium to be achieved through recombination events. Therefore, identification of a haplotype that spans or is linked to a disease-causing mutational change serves as a predictive measure of an individual's likelihood of having inherited that disease-causing mutation. Such prognostic or diagnostic procedures can be utilized without necessitating the identification and isolation of the actual disease-causing lesion. This is significant because the precise determination of the molecular defect involved in a disease process can be difficult and laborious, especially in the case of multifactorial diseases.

The statistical correlation between a disorder and a polymorphism does not necessarily indicate that the polymorphism directly causes the disorder. Rather the correlated polymorphism may be a benign allelic variant which is linked to (i.e., in linkage disequilibrium with) a disorder-causing mutation that has occurred in the recent evolutionary past, so that sufficient time has not elapsed for equilibrium to be achieved through recombination events in the intervening chromosomal segment. Thus, for the purposes of diagnostic and prognostic assays for a particular disease, detection of a polymorphic allele associated with that disease can be utilized without consideration of whether the polymorphism is directly involved in the etiology of the disease. Furthermore, where a given benign polymorphic locus is in linkage disequilibrium with an apparent disease-causing polymorphic locus, still other polymorphic loci which are in linkage disequilibrium with the benign polymorphic locus are also likely to be in linkage disequilibrium with the disease-causing polymorphic locus. Thus, these other polymorphic loci will also be prognostic or diagnostic of the likelihood of having inherited the disease-causing polymorphic locus. A broad-spanning haplotype (describing the typical pattern of co-inheritance of alleles of a set of linked polymorphic markers) can be targeted for diagnostic purposes once an association has been drawn between a particular disease or condition and a corresponding haplotype. Thus, the determination of an individual's likelihood for developing a particular disease or condition can be made by characterizing one or more disease-associated polymorphic

alleles (or even one or more disease-associated haplotypes) without necessarily determining or characterizing the causative genetic variation.

Many methods are available for detecting specific alleles at polymorphic loci. Certain methods for detecting a specific polymorphic allele will depend, in part, upon the molecular nature of the polymorphism. For example, the various allelic forms of the polymorphic locus may differ by a single base-pair of the DNA. Such single nucleotide polymorphisms (or SNPs) are major contributors to genetic variation, comprising some 80% of all known polymorphisms, and their density in the genome is estimated to be on average 1 per 1,000 base pairs. SNPs are most frequently bi-allelic, or occurring in only two different forms (although up to four different forms of an SNP, corresponding to the four different nucleotide bases occurring in DNA, are theoretically possible). Nevertheless, SNPs are mutationally more stable than other polymorphisms, making them suitable for association studies in which linkage disequilibrium between markers and an unknown variant is used to map disease-causing mutations. In addition, because SNPs typically have only two alleles, they can be genotyped by a simple plus / minus assay rather than a length measurement, making them more amenable to automation.

In one embodiment, allelic profiling can be accomplished using a nucleic acid microarray. The genetic testing field is rapidly evolving and, as such, the skilled artisan will appreciate that a wide range of profiling tests exist, and will be developed, to determine the allelic profile of individuals in accord with the disclosure.

Nucleic Acids and Polypeptides

The term “nucleic acid” refers to deoxyribonucleotides or ribonucleotides and polymers thereof in either single- or double-stranded form, made of monomers (nucleotides) containing a sugar, phosphate and a base that is either a purine or pyrimidine. Unless specifically limited, the term encompasses nucleic acids containing known analogs of natural nucleotides that have similar binding properties as the reference nucleic acid and are metabolized in a manner similar to naturally occurring nucleotides. Unless otherwise indicated, a particular nucleic acid sequence also encompasses conservatively modified variants thereof (*e.g.*, degenerate codon substitutions) and complementary sequences, as well as the sequence explicitly indicated. Specifically, degenerate codon substitutions may be achieved by generating sequences in which the third position of one or more selected (or all) codons is substituted with mixed-base

and/or deoxyinosine residues. The terms “nucleic acid,” “nucleic acid molecule,” or “polynucleotide” are used interchangeably and may also be used interchangeably with gene, cDNA, DNA and/or RNA encoded by a gene.

5 The term “nucleotide sequence” refers to a polymer of DNA or RNA which can be single-stranded or double-stranded, optionally containing synthetic, non-natural or altered nucleotide bases capable of incorporation into DNA or RNA polymers. A DNA molecule or polynucleotide is a polymer of deoxyribonucleotides (A, G, C, and T), and an RNA molecule or polynucleotide is a polymer of ribonucleotides (A, G, C and U).

10 A “gene,” for the purposes of the present disclosure, includes a DNA region encoding a gene product, as well as all DNA regions, which regulate the production of the gene product, whether or not such regulatory sequences are adjacent to coding and/or transcribed sequences. The term “gene” is used broadly to refer to any segment of nucleic acid associated with a biological function. Genes include coding sequences and/or the regulatory sequences required for their expression. Accordingly, a gene
15 includes, but is not necessarily limited to, promoter sequences, terminators, translational regulatory sequences such as ribosome binding sites and internal ribosome entry sites, enhancers, silencers, insulators, boundary elements, replication origins, matrix attachment sites and locus control regions. For example, “gene” refers to a nucleic acid fragment that expresses mRNA, functional RNA, or specific protein, including regulatory sequences.
20 “Functional RNA” refers to sense RNA, antisense RNA, ribozyme RNA, siRNA, or other RNA that may not be translated but yet has an effect on at least one cellular process. “Genes” also include non-expressed DNA segments that, for example, form recognition sequences for other proteins. “Genes” can be obtained from a variety of sources, including cloning from a source of interest or synthesizing from known or predicted
25 sequence information, and may include sequences designed to have desired parameters.

“Gene expression” refers to the conversion of the information, contained in a gene, into a gene product. It refers to the transcription and/or translation of an endogenous gene, heterologous gene or nucleic acid segment, or a transgene in cells. In addition, expression refers to the transcription and stable accumulation of sense (mRNA)
30 or functional RNA. Expression may also refer to the production of protein. The term “altered level of expression” refers to the level of expression in transgenic cells or organisms that differs from that of normal or untransformed cells or organisms.

A gene product can be the transcriptional product of a gene (*e.g.*, mRNA, tRNA, rRNA, antisense RNA, ribozyme, structural RNA or any other type of RNA) or a protein produced by translation of an mRNA. Gene products also include RNAs that are modified, by processes such as capping, polyadenylation, methylation, and editing, and proteins modified by, for example, methylation, acetylation, phosphorylation, ubiquitination, ADP-ribosylation, myristilation, and glycosylation. The term “RNA transcript” refers to the product resulting from RNA polymerase-catalyzed transcription of a DNA sequence. When the RNA transcript is a complementary copy of the DNA sequence, it is referred to as the primary transcript; a RNA sequence derived from post-transcriptional processing of the primary transcript is referred to as the mature RNA. “Messenger RNA” (mRNA) refers to the RNA that lacks introns and that can be translated into protein by the cell. “cDNA” refers to a single- or a double-stranded DNA that is complementary to and derived from mRNA. “Functional RNA” refers to sense RNA, antisense RNA, ribozyme RNA, siRNA, or other RNA that may not be translated but yet has an effect on at least one cellular process.

A “coding sequence” or a sequence that “encodes” a polypeptide is a nucleic acid molecule that is transcribed (in the case of DNA) and/or translated (in the case of mRNA) into a polypeptide *in vivo* when placed under the control of appropriate regulatory sequences. The boundaries of the coding sequence are determined by a start codon at the 5′ (amino) terminus and a translation stop codon at the 3′ (carboxy) terminus. A coding sequence can include, but is not limited to, cDNA from viral, prokaryotic or eukaryotic mRNA, genomic DNA sequences from viral (*e.g.*, DNA viruses and retroviruses) or prokaryotic DNA, and synthetic DNA sequences. A transcription termination sequence can be located 3′ to the coding sequence.

“Regulatory sequences” and “suitable regulatory sequences” each refer to nucleotide sequences located upstream (5′ non-coding sequences), within, or downstream (3′ non-coding sequences) of a coding sequence, and which influence the transcription, RNA processing or stability, or translation of the associated coding sequence. Regulatory sequences include enhancers, promoters, translation leader sequences, introns, and polyadenylation signal sequences. They include natural and synthetic sequences as well as sequences that may be a combination of synthetic and natural sequences.

Certain embodiments of the disclosure encompass isolated or substantially purified nucleic acid compositions. In the context of the present disclosure, an “isolated”

or “purified” DNA molecule or RNA molecule is a DNA molecule or RNA molecule that exists apart from its native environment and is, therefore, not a product of nature. An isolated DNA molecule or RNA molecule may exist in a purified form or may exist in a non-native environment such as, for example, a transgenic host cell. For example, an
5 “isolated” or “purified” nucleic acid molecule is substantially free of other cellular material, or culture medium when produced by recombinant techniques, or substantially free of chemical precursors or other chemicals when chemically synthesized. In one embodiment, an “isolated” nucleic acid is free of sequences that naturally flank the nucleic acid (i.e., sequences located at the 5' and 3' ends of the nucleic acid) in the
10 genomic DNA of the organism from which the nucleic acid is derived.

By “fragment” is intended a polypeptide consisting of only a part of the intact full-length polypeptide sequence and structure. The fragment can include a C-terminal deletion, an N-terminal deletion, and/or an internal deletion of the native polypeptide. A fragment of a protein will generally include at least about 5-100 contiguous amino acid
15 residues of the full-length molecule (e.g., at least about 15-25 contiguous amino acid residues of the full-length molecule, at least about 20-50 or more contiguous amino acid residues of the full-length molecule, or any integer between 5 amino acids and the full-length sequence).

“Naturally occurring” is used to describe a composition that can be found in
20 nature as distinct from being artificially produced. For example, a nucleotide sequence present in an organism, which can be isolated from a source in nature and which has not been intentionally modified by a person in the laboratory, is naturally occurring.

A “5' non-coding sequence” refers to a nucleotide sequence located 5' (upstream) to the coding sequence. 5' non-coding sequences are present in the fully processed
25 mRNA upstream of the initiation codon and may affect processing of the primary transcript to mRNA, mRNA stability or translation efficiency. A “3' non-coding sequence” refers to nucleotide sequences located 3' (downstream) to a coding sequence and may include polyadenylation signal sequences and other sequences encoding regulatory signals capable of affecting mRNA processing or gene expression.

30 A “promoter” refers to a nucleotide sequence, usually upstream (5') to its coding sequence, which directs and/or controls the expression of the coding sequence by providing the recognition for RNA polymerase and other factors required for proper transcription. “Promoter” can include a minimal promoter that is a short DNA sequence

comprised of a TATA-box and other sequences that serve to specify the site of transcription initiation, to which regulatory elements are added for control of expression. “Promoter” also can refer to a nucleotide sequence that includes a minimal promoter plus one or more regulatory elements (e.g., enhancers) that are capable of controlling the expression of a coding sequence or functional RNA. Promoters may be derived in their entirety from a native sequence, or be composed of different elements derived from different promoters found in nature, or even be comprised of synthetic DNA sequences. A promoter may also contain DNA sequences that are involved in the binding of protein factors that control the effectiveness of transcription initiation in response to physiological or developmental conditions. “Constitutive expression” refers to expression using a constitutive promoter. “Conditional” and “regulated expression” refer to expression controlled by a regulated promoter.

An “enhancer” is a DNA sequence that can stimulate promoter activity. An enhancer may be an innate element of the promoter or a heterologous element inserted to enhance the level or tissue specificity of a promoter. Enhancers often are capable of operating in both orientations, and are capable of functioning even when moved either upstream or downstream from the promoter. Both enhancers and other regulatory elements within a promoter bind sequence-specific DNA-binding proteins that mediate their effects.

“Operably-linked” refers to the association of nucleic acid sequences on a single nucleic acid fragment so that the function of one of the sequences is affected by another. For example, a regulatory DNA sequence is said to be “operably linked to” or “associated with” a DNA sequence that codes for an RNA or a polypeptide if the two sequences are situated such that the regulatory DNA sequence affects expression of the coding DNA sequence (i.e., that the coding sequence or functional RNA is under the transcriptional control of the promoter). Coding sequences can be operably-linked to regulatory sequences in sense or antisense orientation.

“Expression” refers to the transcription and/or translation of an endogenous gene, heterologous gene or nucleic acid segment, or a transgene in cells. In addition, expression refers to the transcription and stable accumulation of sense (mRNA) or functional RNA. Expression may also refer to the production of protein. The term “altered level of expression” refers to a level of expression in cells or organisms that differs from that of normal cells or organisms.

For sequence comparison, typically one sequence acts as a reference sequence to which test sequences are compared. When using a sequence comparison algorithm, test and reference sequences are input into a computer, and sequence algorithm program parameters are designated. The sequence comparison algorithm then calculates the percent sequence identity for the test sequence(s) relative to the reference sequence, based on the designated algorithm parameters.

The following terms are used to describe the sequence relationships between two or more nucleic acids or polynucleotides: (a) "reference sequence," (b) "comparison window," (c) "sequence identity," (d) "percentage of sequence identity," and (e) "substantial identity." As used herein, "reference sequence" is a defined sequence used as a basis for sequence comparison. A reference sequence may be a subset or the entirety of a specified sequence; for example, as a segment of a full-length cDNA or gene sequence, or the complete cDNA or gene sequence. As used herein, "comparison window" makes reference to a contiguous and specified segment of a polynucleotide sequence, wherein the polynucleotide sequence in the comparison window may comprise additions or deletions (i.e., gaps) compared to the reference sequence (which does not comprise additions or deletions) for optimal alignment of the two sequences. Generally, the comparison window is at least 20 contiguous nucleotides in length, and optionally can be 30, 40, 50, 100, or longer. Those of skill in the art understand that, to avoid a high similarity to a reference sequence due to inclusion of gaps in the polynucleotide sequence, a gap penalty is typically introduced and is subtracted from the number of matches.

Methods of alignment of sequences for comparison are well-known in the art. Thus, the determination of percent identity between any two sequences can be accomplished using a mathematical algorithm. Non-limiting examples of such mathematical algorithms are the algorithm of Myers and Miller (Myers and Miller, CABIOS, 4, 11 (1988)); the local homology algorithm of Smith et al. (Smith et al., Adv. Appl. Math., 2, 482 (1981)); the homology alignment algorithm of Needleman and Wunsch (Needleman and Wunsch, JMB, 48, 443 (1970)); the search-for-similarity-method of Pearson and Lipman (Pearson and Lipman, Proc. Natl. Acad. Sci. USA, 85, 2444 (1988)); the algorithm of Karlin and Altschul (Karlin and Altschul, Proc. Natl. Acad. Sci. USA, 87, 2264 (1990)), modified as in Karlin and Altschul (Karlin and Altschul, Proc. Natl. Acad. Sci. USA 90, 5873 (1993)).

Computer implementations of these mathematical algorithms can be utilized for comparison of sequences to determine sequence identity. Such implementations include, but are not limited to: CLUSTAL in the PC/Gene program (available from Intelligenetics, Mountain View, Calif.); the ALIGN program (Version 2.0) and GAP, BESTFIT, BLAST, FASTA, and TFASTA in the Wisconsin Genetics Software Package, Version 8 (available from Genetics Computer Group (GCG), 575 Science Drive, Madison, Wis., USA). Alignments using these programs can be performed using the default parameters. The CLUSTAL program is well described by Higgins et al. (Higgins et al., CABIOS, 5, 151 (1989)); Corpet et al. (Corpet et al., Nucl. Acids Res., 16, 10881 (1988)); Huang et al. (Huang et al., CABIOS, 8, 155 (1992)); and Pearson et al. (Pearson et al., Meth. Mol. Biol., 24, 307 (1994)). The ALIGN program is based on the algorithm of Myers and Miller, supra. The BLAST programs of Altschul et al. (Altschul et al., J. Mol. Biol., 215, 403 (1990)) are based on the algorithm of Karlin and Altschul, supra.

Software for performing BLAST analyses is publicly available through the National Center for Biotechnology Information. This algorithm involves first identifying high scoring sequence pairs (HSPs) by identifying short words of length “W” in the query sequence, which either match or satisfy some positive-valued threshold score T when aligned with a word of the same length in a database sequence. “T” is referred to as the neighborhood word score threshold. These initial neighborhood word hits act as seeds for initiating searches to find longer HSPs containing them. The word hits are then extended in both directions along each sequence for as far as the cumulative alignment score can be increased. Cumulative scores are calculated using, for nucleotide sequences, the parameters “M” (reward score for a pair of matching residues; always >0) and “N” (penalty score for mismatching residues; always <0), and for amino acid sequences, a scoring matrix is used to calculate the cumulative score. Extension of the word hits in each direction are halted when the cumulative alignment score falls off by the quantity “X” from its maximum achieved value, the cumulative score goes to zero or below due to the accumulation of one or more negative-scoring residue alignments, or the end of either sequence is reached.

In addition to calculating percent sequence identity, the BLAST algorithm also performs a statistical analysis of the similarity between two sequences. One measure of similarity provided by the BLAST algorithm is the smallest sum probability (P(N)), which provides an indication of the probability by which a match between two nucleotide

or amino acid sequences would occur by chance. For example, a test nucleic acid sequence is considered similar to a reference sequence if the smallest sum probability in a comparison of the test nucleic acid sequence to the reference nucleic acid sequence is less than about 0.1, less than about 0.01, or even less than about 0.001.

5 To obtain gapped alignments for comparison purposes, Gapped BLAST (in BLAST 2.0) can be utilized. Alternatively, PSI-BLAST (in BLAST 2.0) can be used to perform an iterated search that detects distant relationships between molecules. When utilizing BLAST, Gapped BLAST, PSI-BLAST, the default parameters of the respective programs (e.g., BLASTN for nucleotide sequences, BLASTX for proteins) can be used.

10 The BLASTN program (for nucleotide sequences) uses as defaults a wordlength (W) of 11, an expectation (E) of 10, a cutoff of 100, M=5, N=-4, and a comparison of both strands. For amino acid sequences, the BLASTP program uses as defaults a wordlength (W) of 3, an expectation (E) of 10, and the BLOSUM62 scoring matrix. Alignment may also be performed manually by inspection.

15 For purposes of the present disclosure, comparison of nucleotide sequences for determination of percent sequence identity to the promoter sequences disclosed herein may be made using the BlastN program (version 1.4.7 or later) with its default parameters or any equivalent program. By “equivalent program” is intended any sequence comparison program that, for any two sequences in question, generates an alignment

20 having identical nucleotide or amino acid residue matches and an identical percent sequence identity when compared to the corresponding alignment generated by the program.

As used herein, “sequence identity” or “identity” in the context of two nucleic acid or polypeptide sequences makes reference to a specified percentage of residues in the

25 two sequences that are the same when aligned for maximum correspondence over a specified comparison window, as measured by sequence comparison algorithms or by visual inspection. When percentage of sequence identity is used in reference to proteins it is recognized that residue positions which are not identical often differ by conservative amino acid substitutions, where amino acid residues are substituted for other amino acid

30 residues with similar chemical properties (e.g., charge or hydrophobicity) and, therefore, do not change the functional properties of the molecule. When sequences differ in conservative substitutions, the percent sequence identity may be adjusted upwards to correct for the conservative nature of the substitution. Sequences that differ by such

conservative substitutions are said to have “sequence similarity” or “similarity.” Means for making this adjustment are well known to those of skill in the art. Typically, this involves scoring a conservative substitution as a partial rather than a full mismatch, thereby increasing the percentage sequence identity. Thus, for example, where an identical amino acid is given a score of 1 and a non-conservative substitution is given a score of zero, a conservative substitution is given a score between zero and 1. The scoring of conservative substitutions is calculated, e.g., as implemented in the program PC/GENE (Intelligenetics, Mountain View, Calif).

As used herein, “percent sequence identity” means the value determined by comparing two optimally aligned sequences over a comparison window, wherein the portion of the polynucleotide sequence in the comparison window may comprise additions or deletions (i.e., gaps) as compared to the reference sequence (which does not comprise additions or deletions) for optimal alignment of the two sequences. The percentage is calculated by determining the number of positions at which the identical nucleic acid base or amino acid residue occurs in both sequences to yield the number of matched positions, dividing the number of matched positions by the total number of positions in the window of comparison, and multiplying the result by 100 to yield the percentage of sequence identity.

The term “substantial identity” of polynucleotide sequences means that a polynucleotide comprises a sequence that has at least 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, or 94%, or even at least 95%, 96%, 97%, 98%, 99% or 100% sequence identity, compared to a reference sequence using one of the alignment programs described herein using standard parameters. One of skill in the art will recognize that these values can be appropriately adjusted to determine corresponding identity of proteins encoded by two nucleotide sequences by taking into account codon degeneracy, amino acid similarity, reading frame positioning, and the like. Substantial identity of amino acid sequences for these purposes normally means sequence identity of at least 70% (e.g., 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%), at least 80% (e.g., 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%), at least 90% (e.g., 91%, 92%, 93%, or 94%), or even at least 95% (e.g., 96%, 97%, 98%, 99%, or 100%).

The term “substantial identity” in the context of a peptide indicates that a peptide comprises a sequence with at least 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%,

79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, or 94%, or even 95%, 96%, 97%, 98% or 99%, sequence identity to the reference sequence over a specified comparison window. In certain embodiments, optimal alignment is conducted using the homology alignment algorithm of Needleman and Wunsch (Needleman and Wunsch, J. Mol. Biol., 48, 443 (1970)). An indication that two peptide sequences are substantially identical is that one peptide is immunologically reactive with antibodies raised against the second peptide. Thus, a peptide is substantially identical to a second peptide, for example, where the two peptides differ only by a conservative substitution. Thus, the disclosure also provides nucleic acid molecules and peptides that are substantially identical to the nucleic acid molecules and peptides presented herein.

Another indication that nucleotide sequences are substantially identical is if two molecules hybridize to each other under stringent conditions. Hybridization of nucleic acids is discussed in more detail below.

Oligonucleotide Primers and Probes

As used herein, “primer,” “probe,” and “oligonucleotide” are used interchangeably. The term “nucleic acid probe” or a “probe specific for” a nucleic acid refers to a nucleic acid sequence that has at least about 80%, e.g., at least about 90%, e.g., at least about 95% contiguous sequence identity or homology to the nucleic acid sequence encoding the targeted sequence of interest. A probe (or oligonucleotide or primer) of the disclosure is at least about 8 nucleotides in length (e.g., at least about 8-50 nucleotides in length, e.g., at least about 10-40, e.g., at least about 15-35 nucleotides in length). The oligonucleotide probes or primers of the disclosure may comprise at least about eight nucleotides at the 3' of the oligonucleotide that have at least about 80%, e.g., at least about 85%, e.g., at least about 90%, e.g., at least about 95% contiguous identity to the targeted sequence of interest.

Primer pairs are useful for determination of the nucleotide sequence of a particular SNP using PCR. The pairs of single-stranded DNA primers can be annealed to sequences within or surrounding the SNP in order to prime amplifying DNA synthesis of the SNP itself. The first step of the process involves contacting a biological sample obtained from a subject, which sample contains nucleic acid, with at least one primer to form a hybridized DNA. The oligonucleotide primers that are useful in the methods of the present disclosure can be any primer comprised of about 8 bases up to about 80 or 100

bases or more. In one embodiment of the present disclosure, the primers are between about 10 and about 20 bases.

The primers themselves can be synthesized using techniques that are well known in the art. Generally, the primers can be made using oligonucleotide synthesizing machines that are commercially available.

The primers or probes of the present disclosure can be labeled using techniques known to those of skill in the art. For example, the labels used in the assays of disclosure can be primary labels (where the label comprises an element that is detected directly) or secondary labels (where the detected label binds to a primary label, e.g., as is common in immunological labeling). An introduction to labels (also called “tags”), tagging or labeling procedures, and detection of labels is found in Polak and Van Noorden (1997) Introduction to Immunocytochemistry, second edition, Springer Verlag, N.Y. and in Haugland (1996) Handbook of Fluorescent Probes and Research Chemicals, a combined handbook and catalogue Published by Molecular Probes, Inc., Eugene, Oreg. Primary and secondary labels can include undetected elements as well as detected elements. Useful primary and secondary labels in the present disclosure can include spectral labels such as fluorescent dyes (e.g., fluorescein and derivatives such as fluorescein isothiocyanate (FITC) and Oregon Green™, rhodamine and derivatives (e.g., Texas red, tetramethylrhodamine isothiocyanate (TRITC), etc.), digoxigenin, biotin, phycoerythrin, AMCA, CyDyes™, and the like), radiolabels (e.g., ³H, ¹²⁵I, ³⁵S, ¹⁴C, ³²P, ³³P), enzymes (e.g., horse-radish peroxidase, alkaline phosphatase) spectral colorimetric labels such as colloidal gold or colored glass or plastic (e.g., polystyrene, polypropylene, latex) beads. The label may be coupled directly or indirectly to a component of the detection assay (e.g., the labeled nucleic acid) according to methods well known in the art. As indicated above, a wide variety of labels may be used, with the choice of label depending on sensitivity required, ease of conjugation with the compound, stability requirements, available instrumentation, and disposal provisions.

In general, a detector that monitors a probe-substrate nucleic acid hybridization is adapted to the particular label that is used. Typical detectors include spectrophotometers, phototubes and photodiodes, microscopes, scintillation counters, cameras, film and the like, as well as combinations thereof. Examples of suitable detectors are widely available from a variety of commercial sources known to persons of skill. Commonly, an optical

image of a substrate comprising bound labeled nucleic acids is digitized for subsequent computer analysis.

Labels include those that use (1) chemiluminescence (using Horseradish Peroxidase and/or Alkaline Phosphatase with substrates that produce photons as breakdown products) with kits being available, *e.g.*, from Molecular Probes, Amersham, 5 Boehringer-Mannheim, and Life Technologies/Gibco BRL; (2) color production (using both Horseradish Peroxidase and/or Alkaline Phosphatase with substrates that produce a colored precipitate) (kits available from Life Technologies/Gibco BRL, and Boehringer-Mannheim); (3) hemifluorescence using, *e.g.*, Alkaline Phosphatase and the substrate 10 AttoPhos (Amersham) or other substrates that produce fluorescent products, (4) fluorescence (*e.g.*, using Cy-5 (Amersham), fluorescein, and other fluorescent labels); (5) radioactivity using kinase enzymes or other end-labeling approaches, nick translation, random priming, or PCR to incorporate radioactive molecules into the labeled nucleic acid. Other methods for labeling and detection will be readily apparent to one skilled in 15 the art.

Fluorescent labels can be used and have the advantage of requiring fewer precautions in handling, and being amendable to high-throughput visualization techniques (optical analysis including digitization of the image for analysis in an integrated system comprising a computer). Preferred labels are typically characterized by one or more of 20 the following: high sensitivity, high stability, low background, low environmental sensitivity and high specificity in labeling. Fluorescent moieties, which can be incorporated into a label, generally are known including Texas red, dioxigenin, biotin, 1- and 2-aminonaphthalene, p,p'-diaminostilbenes, pyrenes, quaternary phenanthridine salts, 9-aminoacridines, p,p'-diaminobenzophenone imines, anthracenes, oxacarbocyanine, 25 merocyanine, 3-aminoequilenin, perylene, bis-benzoxazole, bis-p-oxazolyl benzene, 1,2-benzophenazin, retinol, bis-3-aminopyridinium salts, hellebrigenin, tetracycline, sterophenol, benzimidazolylphenylamine, 2-oxo-3-chromen, indole, xanthen, 7-hydroxycoumarin, phenoxazine, calicylate, strophanthidin, porphyrins, triarylmethanes, flavin and many others. Many fluorescent labels are commercially available from the 30 SIGMA Chemical Company (Saint Louis, MO), Molecular Probes, R&D systems (Minneapolis, MN), Pharmacia LKB Biotechnology (Piscataway, NJ), CLONTECH Laboratories, Inc. (Palo Alto, CA), Chem Genes Corp., Aldrich Chemical Company (Milwaukee, WI), Glen Research, Inc., GIBCO BRL Life Technologies, Inc.

(Gaithersburg, MD), Fluka ChemicaBiochemika Analytika (Fluka Chemie AG, Buchs, Switzerland), and Applied Biosystems™ (Foster City, CA), as well as many other commercial sources known to one of skill.

Means of detecting and quantifying labels are well known to those of skill in the art. Thus, for example, when the label is a radioactive label, means for detection include a scintillation counter or photographic film as in autoradiography; and when the label is optically detectable, typical detectors include microscopes, cameras, phototubes, photodiodes and many other detection systems that are widely available.

Oligonucleotide primers or probes may be prepared having any of a wide variety of base sequences according to techniques that are well known in the art. Suitable bases for preparing an oligonucleotide primer or probe may be selected from naturally occurring nucleotide bases such as adenine, cytosine, guanine, uracil, and thymine; and non-naturally occurring or "synthetic" nucleotide bases such as 7-deaza-guanine 8-oxo-guanine, 6-mercaptoguanine, 4-acetylcytidine, 5-(carboxyhydroxyethyl)uridine, 2'-O-methylcytidine, 5-carboxymethylamino-methyl-2-thioridine, 5-carboxymethylaminomethyluridine, dihydrouridine, 2'-O-methylpseudouridine, β ,D-galactosylqueosine, 2'-O-methylguanosine, inosine, N6-isopentenyladenosine, 1-methyladenosine, 1-methylpseudouridine, 1-methylguanosine, 1-methylinosine, 2,2-dimethylguanosine, 2-methyladenosine, 2-methylguanosine, 3-methylcytidine, 5-methylcytidine, N6-methyladenosine, 7-methylguanosine, 5-methylaminomethyluridine, 5-methoxyaminomethyl-2-thiouridine, β ,D-mannosylqueosine, 5-methoxycarbonylmethyluridine, 5-methoxyuridine, 2-methylthio-N6-isopentenyladenosine, N-((9- β -D-ribofuranosyl-2-methylthiopurine-6-yl)carbamoyl)threonine, N-((9- β -D-ribofuranosylpurine-6-yl)N-methylcarbamoyl)threonine, uridine-5-oxyacetic acid methylester, uridine-5-oxyacetic acid, wybutoxosine, pseudouridine, queosine, 2-thiocytidine, 5-methyl-2-thiouridine, 2-thiouridine, 2-thiouridine, 5-Methyluridine, N-((9-beta-D-ribofuranosylpurine-6-yl)carbamoyl)threonine, 2'-O-methyl-5-methyluridine, 2'-O-methyluridine, wybutosine, and 3-(3-amino-3-carboxypropyl)uridine. Any oligonucleotide backbone may be employed, including DNA, RNA (although RNA is less preferred than DNA), modified sugars such as carbocycles, and sugars containing 2' substitutions such as fluoro and methoxy. The oligonucleotides may be oligonucleotides wherein at least one, or all, of the internucleotide bridging phosphate residues are modified phosphates, such as methyl

phosphonates, methyl phosphonotioates, phosphorinopholidates,
phosphoropiperazidates and phosphoramidates (for example, every other one of the
internucleotide bridging phosphate residues may be modified as described). The
oligonucleotide may be a "peptide nucleic acid" such as described in Nielsen *et al.*,
5 Science, 254:1497-1500 (1991).

As used herein, a "single base pair extension probe" is a nucleic acid that
selectively recognizes a single nucleotide polymorphism (i.e., either the A or the G of an
A/G polymorphism). Generally, these probes take the form of a DNA primer (e.g., as in
PCR primers) that are modified so that incorporation of the primer releases a fluorophore.
10 One example of this is a Taqman[®] probe that uses the 5' exonuclease activity of the
enzyme Taq Polymerase for measuring the amount of target sequences in the samples.
TaqMan[®] probes consist of a 18-22 bp oligonucleotide probe, which is labeled with a
reporter fluorophore at the 5' end, and a quencher fluorophore at the 3' end. Incorporation
of the probe molecule into a PCR chain (which occurs because the probe set is contained
15 in a mixture of PCR primers) liberates the reporter fluorophore from the effects of the
quencher. The primer must be able to recognize the target binding site. Some primer
extension probes can be "activated" directly by DNA polymerase without a full PCR
extension cycle.

The only requirement is that the oligonucleotide probe should possess a sequence
20 at least a portion of which is capable of binding to a known portion of the sequence of the
DNA sample. The nucleic acid probes provided by the present disclosure are useful for a
number of purposes.

Methods of Detecting Nucleic Acids

A. Amplification

According to the methods of the present disclosure, the amplification of DNA
present in a biological sample may be carried out by any means known to the art.
Examples of suitable amplification techniques include, but are not limited to, polymerase
chain reaction (including, for RNA amplification, reverse-transcriptase polymerase chain
reaction), ligase chain reaction, strand displacement amplification, transcription-based
30 amplification, self-sustained sequence replication (or "3SR"), the Qbeta replicase system,
nucleic acid sequence-based amplification (or "NASBA"), the repair chain reaction (or
"RCR"), and boomerang DNA amplification (or "BDA").

The bases incorporated into the amplification product can be natural or modified bases (modified before or after amplification), and the bases can be selected to optimize subsequent electrochemical detection steps.

5 Polymerase chain reaction (PCR) can be carried out in accordance with known techniques. See, e.g., U.S. Patent Numbers 4,683,195; 4,683,202; 4,800,159; and 4,965,188. In general, PCR involves, first, treating a nucleic acid sample (e.g., in the presence of a heat stable DNA polymerase) with one oligonucleotide primer for each strand of the specific sequence to be detected under hybridizing conditions so that an extension product of each primer is synthesized that is complementary to each nucleic acid strand, with the primers sufficiently complementary to each strand of the specific sequence to hybridize therewith so that the extension product synthesized from each primer, when it is separated from its complement, can serve as a template for synthesis of the extension product of the other primer, and then treating the sample under denaturing conditions to separate the primer extension products from their templates if the sequence or sequences to be detected are present. These steps are cyclically repeated until the desired degree of amplification is obtained. Detection of the amplified sequence may be carried out by adding, to the reaction product, an oligonucleotide probe capable of hybridizing to the reaction product (e.g., an oligonucleotide primer or probe of the present disclosure), the probe carrying a detectable label, and then detecting the label in accordance with known techniques. Various labels that can be incorporated into or operably linked to nucleic acids are well known in the art, such as radioactive, enzymatic, and florescent labels. Where the nucleic acid to be amplified is RNA, amplification may be carried out by initial conversion to DNA by reverse transcriptase in accordance with known techniques.

25 Strand displacement amplification (SDA) can be carried out in accordance with known techniques. For example, SDA can be carried out with a single amplification primer or a pair of amplification primers, with exponential amplification being achieved with the latter. In general, SDA amplification primers comprise, in the 5' to 3' direction, a flanking sequence (the DNA sequence of which is noncritical), a restriction site for the restriction enzyme employed in the reaction, and an oligonucleotide sequence (e.g., an oligonucleotide primer or probe as described herein) that hybridizes to the target sequence to be amplified and/or detected. The flanking sequence, which serves to facilitate binding of the restriction enzyme to the recognition site and provides a DNA polymerase priming

site after the restriction site has been nicked, can be about 15 to 20 nucleotides in length. The restriction site is functional in the SDA reaction. For example, the oligonucleotide primer or probe portion can be about 13 to 15 nucleotides in length.

5 Ligase chain reaction (LCR) also can be carried out in accordance with known techniques. In general, the reaction is carried out with two pairs of oligonucleotide probes: one pair binds to one strand of the sequence to be detected; the other pair binds to the other strand of the sequence to be detected. Each pair together completely overlaps the strand to which it corresponds. The reaction is carried out by, first, denaturing (e.g., separating) the strands of the sequence to be detected, then reacting the strands with the
10 two pairs of oligonucleotide probes in the presence of a heat stable ligase so that each pair of oligonucleotide probes is ligated together, then separating the reaction product, and then cyclically repeating the process until the sequence has been amplified to the desired degree. Detection then can be carried out in like manner as described above with respect to PCR.

15 According to the methods described herein, a particular SNP at a particular locus can be detected. Techniques that are useful in the methods described herein include, but are not limited to, direct DNA sequencing, PFGE analysis, allele-specific oligonucleotide (ASO), dot blot analysis and denaturing gradient gel electrophoresis, and are well known to a skilled artisan.

20 There are several methods that can be used to detect DNA sequence variation. Direct DNA sequencing, either manual sequencing or automated fluorescent sequencing can detect sequence variation. Another approach is the single-stranded conformation polymorphism assay (SSCA). This method does not detect all sequence changes, especially if the DNA fragment size is greater than 200 bp, but can be optimized to detect
25 most DNA sequence variation. The reduced detection sensitivity is a disadvantage, but the increased throughput possible with SSCA makes it an attractive, viable alternative to direct sequencing for mutation detection on a research basis. The fragments that have shifted mobility on SSCA gels then can be sequenced to determine the exact nature of the DNA sequence variation. Other approaches based on the detection of mismatches
30 between the two complementary DNA strands include clamped denaturing gel electrophoresis (CDGE), heteroduplex analysis (HA) and chemical mismatch cleavage (CMC). Once a sequence change has been identified, an allele specific detection approach such as allele specific oligonucleotide (ASO) hybridization can be utilized to

rapidly screen large numbers of other samples for that same sequence change (e.g., mutation, polymorphism). Such a technique can utilize probes that are labeled with gold nanoparticles to yield a visual color result.

5 Detection of SNPs can be accomplished by sequencing the desired target region using techniques well known in the art. Alternatively, sequences can be amplified directly from a genomic DNA preparation from tissue from a subject using known techniques. The DNA sequence of the amplified sequences then can be determined.

There are several well known methods for a more complete, yet still indirect, test for confirming the presence of a mutant allele: 1) single stranded conformation analysis (SSCA); 2) denaturing gradient gel electrophoresis (DGGE); 3) RNase protection assays; 10 4) allele-specific oligonucleotides (ASOs); 5) the use of proteins which recognize nucleotide mismatches, such as the *E. coli* mutS protein; and/or 6) allele-specific PCR. For allele-specific PCR, primers are used that hybridize at their 3' ends to a particular allele. If the particular mutation is not present, an amplification product is not observed. 15 Amplification Refractory Mutation System (ARMS) can also be used. Insertions and deletions of genes can also be detected by cloning, sequencing and amplification. In addition, restriction fragment length polymorphism (RFLP) probes for the gene or surrounding marker genes can be used to score alteration of an allele or an insertion in a polymorphic fragment. Other techniques for detecting insertions and deletions as known 20 in the art can be used.

In the first three methods (SSCA, DGGE and RNase protection assay), a new electrophoretic band appears. SSCA detects a band that migrates differentially because the sequence change causes a difference in single-strand, intramolecular base pairing. RNase protection involves cleavage of the mutant polynucleotide into two or more 25 smaller fragments. DGGE detects differences in migration rates of mutant sequences compared to wild-type sequences, using a denaturing gradient gel. In an allele-specific oligonucleotide assay, an oligonucleotide is designed which detects a specific sequence, and the assay is performed by detecting the presence or absence of a hybridization signal. In the mutS assay, the protein binds only to sequences that contain a nucleotide mismatch 30 in a heteroduplex between mutant and wild-type sequences.

Mismatches, according to the present disclosure, are hybridized nucleic acid duplexes in which the two strands are not 100% complementary. Lack of total homology may be due to deletions, insertions, inversions or substitutions. Mismatch detection can

be used to detect point mutations in the gene or in its mRNA product. While these techniques are less sensitive than sequencing, they are simpler to perform on a large number of samples. An example of a mismatch cleavage technique is the RNase protection method. The riboprobe and either mRNA or DNA isolated from the tumor
5 tissue are annealed (hybridized) together and subsequently digested with the enzyme RNase A that is able to detect some mismatches in a duplex RNA structure. If a mismatch is detected by RNase A, it cleaves at the site of the mismatch. Thus, when the annealed RNA preparation is separated on an electrophoretic gel matrix, if a mismatch has been detected and cleaved by RNase A, an RNA product will be seen which is
10 smaller than the full length duplex RNA for the riboprobe and the mRNA or DNA. The riboprobe need not be the full length of the mRNA or gene but can be a segment of either. If the riboprobe includes only a segment of the mRNA or gene, it will be desirable to use a number of these probes to screen the whole mRNA sequence for mismatches.

In similar fashion, DNA probes can be used to detect mismatches, through
15 enzymatic or chemical cleavage. Alternatively, mismatches can be detected by shifts in the electrophoretic mobility of mismatched duplexes relative to matched duplexes. With either riboprobes or DNA probes, the cellular mRNA or DNA that might contain a mutation can be amplified using PCR before hybridization.

B. Hybridization

20 The phrase "hybridizing specifically to" refers to the binding, duplexing, or hybridizing of a molecule only to a particular nucleotide sequence under stringent conditions when that sequence is present in a complex mixture (e.g., total cellular) DNA or RNA. "Bind(s) substantially" refers to complementary hybridization between a primer or probe nucleic acid and a target nucleic acid and embraces minor mismatches that can
25 be accommodated by reducing the stringency of the hybridization media to achieve the desired detection of the target nucleic acid sequence.

Generally, stringent conditions are selected to be about 5°C lower than the thermal melting point (T_m) for the specific sequence at a defined ionic strength and pH. However, stringent conditions encompass temperatures in the range of about 1°C to about 20°C,
30 depending upon the desired degree of stringency as otherwise qualified herein. Nucleic acids that do not hybridize to each other under stringent conditions are still substantially identical if the polypeptides they encode are substantially identical. This may occur, e.g., when a copy of a nucleic acid is created using the maximum codon degeneracy permitted

by the genetic code. One indication that two nucleic acid sequences are substantially identical is when the polypeptide encoded by the first nucleic acid is immunologically cross reactive with the polypeptide encoded by the second nucleic acid.

“Stringent conditions” are those that (1) employ low ionic strength and high
5 temperature for washing, for example, 0.015 M NaCl / 0.0015 M sodium citrate (SSC);
0.1% sodium lauryl sulfate (SDS) at 50°C, or (2) employ a denaturing agent such as
formamide during hybridization, e.g., 50% formamide with 0.1% bovine serum albumin /
0.1% Ficoll / 0.1% polyvinylpyrrolidone / 50 mM sodium phosphate buffer at pH 6.5
with 750 mM NaCl, 75 mM sodium citrate at 42°C. Another example is use of 50%
10 formamide, 5 x SSC (0.75 M NaCl, 0.075 M sodium citrate), 50 mM sodium phosphate
(pH 6.8), 0.1% sodium pyrophosphate, 5x Denhardt's solution, sonicated salmon sperm
DNA (50 µg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2
x SSC and 0.1% SDS. Other examples of stringent conditions are well known in the art.

“Stringent hybridization conditions” and “stringent hybridization wash
15 conditions” in the context of nucleic acid hybridization experiments such as Southern and
Northern hybridizations are sequence dependent, and are different under different
environmental parameters. Longer sequences hybridize specifically at higher
temperatures. The thermal melting point (T_m) is the temperature (under defined ionic
strength and pH) at which 50% of the target sequence hybridizes to a perfectly matched
20 primer or probe sequence. Specificity is typically the function of post-hybridization
washes, the critical factors being the ionic strength and temperature of the final wash
solution. For DNA-DNA hybrids, the T_m can be approximated from the equation of
Meinkoth and Wahl (1984); $T_m = 81.5^\circ\text{C} + 16.6 (\log M) + 0.41 (\%GC) - 0.61 (\% \text{ form}) -$
500/L; where M is the molarity of monovalent cations, %GC is the percentage of
25 guanosine and cytosine nucleotides in the DNA, % form is the percentage of formamide
in the hybridization solution, and L is the length of the hybrid in base pairs. T_m is
reduced by about 1°C for each 1% of mismatching; thus, T_m , hybridization, and/or wash
conditions can be adjusted to hybridize to sequences of the desired identity. For example,
if sequences with >90% identity are sought, the T_m can be decreased 10°C. Generally,
30 stringent conditions are selected to be about 5°C lower than the T_m for the specific
sequence and its complement at a defined ionic strength and pH. However, severely
stringent conditions can utilize a hybridization and/or wash at 1, 2, 3, or 4°C lower than
the T_m ; moderately stringent conditions can utilize a hybridization and/or wash at 6, 7, 8,

9, or 10°C lower than the T_m ; low stringency conditions can utilize a hybridization and/or wash at 11, 12, 13, 14, 15, or 20°C lower than the T_m . Using the equation, hybridization and wash compositions, and desired temperature, those of ordinary skill will understand that variations in the stringency of hybridization and/or wash solutions are inherently described. If the desired degree of mismatching results in a temperature of less than 45°C (aqueous solution) or 32°C (formamide solution), the SSC concentration can be increased so that a higher temperature can be used. Generally, highly stringent hybridization and wash conditions are selected to be about 5°C lower than the T_m for the specific sequence at a defined ionic strength and pH.

An example of highly stringent wash conditions is 0.15 M NaCl at 72°C for about 15 minutes. An example of stringent wash conditions is a 0.2 x SSC wash at 65°C for 15 minutes. Often, a high stringency wash is preceded by a low stringency wash to remove background signal. An example of a medium stringency wash for a duplex of, e.g., more than 100 nucleotides, is 1 x SSC at 45°C for 15 minutes. For short nucleotide sequences (e.g., about 10 to 50 nucleotides), stringent conditions typically involve salt concentrations of less than about 1.5 M, less than about 0.01 to 1.0 M, Na ion concentration (or other salts) at pH 7.0 to 8.3, and the temperature is typically at least about 30°C and at least about 60°C for long oligonucleotides (e.g., >50 nucleotides). Stringent conditions also can be achieved by the addition of destabilizing agents such as formamide. In general, a signal to noise ratio of 2x (or higher) than that observed for an unrelated oligonucleotide in the particular hybridization assay indicates detection of a specific hybridization. Nucleic acids that do not hybridize to each other under stringent conditions are still substantially identical if the proteins that they encode are substantially identical. This can occur, e.g., when a copy of a nucleic acid is created using the maximum codon degeneracy permitted by the genetic code.

Very stringent conditions can be equal to the T_m for a particular oligonucleotide. An example of stringent conditions for hybridization of complementary nucleic acids that have more than 100 complementary residues on a filter in a Southern or Northern blot is 50% formamide, e.g., hybridization in 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1 x SSC at 60 to 65°C. Exemplary low stringency conditions include hybridization with a buffer solution of 30 to 35% formamide, 1 M NaCl, 1% SDS (sodium dodecyl sulphate) at 37°C, and a wash in 1x to 2x SSC (20x SSC = 3.0 M NaCl/0.3 M trisodium citrate) at 50 to 55°C. Exemplary moderate stringency conditions

include hybridization in 40 to 45% formamide, 1.0 M NaCl, 1% SDS at 37°C, and a wash in 0.5x to 1x SSC at 55 to 60°C.

“Northern analysis” or “Northern blotting” is a method used to identify RNA sequences that hybridize to a known probe such as an oligonucleotide, DNA fragment, cDNA or fragment thereof, or RNA fragment. The probe can be labeled with a radioisotope such as ³²P, by biotinylation or with an enzyme. The RNA to be analyzed can be usually electrophoretically separated on an agarose or polyacrylamide gel, transferred to nitrocellulose, nylon, or other suitable membrane, and hybridized with the probe, using standard techniques well known in the art.

Nucleic acid sample may be contacted with an oligonucleotide in any suitable manner known to those skilled in the art. For example, the DNA sample may be solubilized in solution, and contacted with the oligonucleotide by solubilizing the oligonucleotide in solution with the DNA sample under conditions that permit hybridization. Suitable conditions are well known to those skilled in the art.

Alternatively, the DNA sample may be solubilized in solution with the oligonucleotide immobilized on a solid support, whereby the DNA sample may be contacted with the oligonucleotide by immersing the solid support having the oligonucleotide immobilized thereon in the solution containing the DNA sample.

The term “substrate” refers to any solid support to which an oligonucleotide may be attached. The substrate material may be modified, covalently or otherwise, with coatings or functional groups to facilitate binding of oligonucleotides. Suitable substrate materials include polymers, glasses, semiconductors, papers, metals, gels and hydrogels among others. Substrates may have any physical shape or size, e.g., plates, strips, or microparticles. The term “spot” refers to a distinct location on a substrate to which oligonucleotides of known sequence are attached. A spot may be an area on a planar substrate, or it may be, for example, a microparticle distinguishable from other microparticles. The term “bound” means affixed to the solid substrate. A spot is “bound” to the solid substrate when it is affixed in a particular location on the substrate for purposes of the screening assay.

In certain embodiments of the present disclosure, the substrate is a polymer, glass, semiconductor, paper, metal, gel or hydrogel. In certain embodiments of the present disclosure, a kit can further include a solid substrate and at least one control

oligonucleotide, wherein the at least one control oligonucleotide is bound onto the substrate in a distinct spot.

In certain embodiments of the present disclosure, the solid substrate is a microarray. An “array” or “microarray” is used synonymously herein to refer to a plurality of primers or probes attached to one or more distinguishable spots on a substrate. A microarray may include a single substrate or a plurality of substrates, for example a plurality of beads or microspheres. A “copy” of a microarray contains the same types and arrangements of primer or probes.

Methods for Detecting Diabetes

The present disclosure provides a method for determining whether a subject has the likelihood of having diabetes by determining methylation status of a CpG dinucleotide repeat or CpG dinucleotide repeat motif region, where the methylation status of the CpG dinucleotide is associated with diabetes. In certain embodiments, the method determines the methylation status of a plurality (e.g., any integer between 1 and 10,000, such as at least 100) of CpG dinucleotide repeat motif regions.

As used herein, a “biological sample” encompasses essentially any sample type obtained from a subject that can be used in a diagnostic or prognostic method described herein. The biological sample may be any bodily fluid, tissue or any other sample from which clinically relevant biomarkers may be determined. “Biological samples” also can encompass cells in culture, cell supernatants, cell lysates, blood, serum, plasma, urine, cerebral spinal fluid, biological fluid, and tissue samples. In some embodiments, blood samples, or samples derived from blood, e.g. plasma, circulating, peripheral, lymphocytes, etc., are assayed for the presence of one or more SNPs and/or the methylation status of one or more CpG dinucleotides. A biological sample also can be saliva. Typically, a biological sample that contains nucleic acids is provided and tested. Biological samples can be obtained from subjects using well known techniques such as venipuncture, lumbar puncture, fluid sample such as saliva or urine, or the like.

As used herein, the term “healthy” means that a subject does not manifest a particular condition, and is no more likely than at random to be susceptible to a particular condition.

As used herein, the term “predisposition” is defined as a tendency or susceptibility for a subject to manifest a condition. For example, a predisposed subject is more likely to manifest a condition than is a control subject or a healthy subject.

Prevalence is defined by the American Psychological Association (APA) as the
5 “the total number or percentage of cases (e.g., of a disease or disorder) existing in a
population” (APA Dictionary of Psychology, (American Psychological Association,
Washington, DC, 2007)). In some instances, point prevalence is used to describe the
prevalence of cases at a discrete point of time, and period prevalence is used to describe
the number of cases that exist for a period of time (e.g., a month, a year). Prevalence
10 typically is expressed as a rate per population unit (e.g., number of cases per 100,000
people) instead of an absolute number or a percent.

Similarly, incidence is defined by the APA as “the rate of occurrence of new cases
of a given event or condition (e.g., a disorder, disease, symptom, or injury) in a particular
population in a given period” of time (APA Dictionary of Psychology, (American
15 Psychological Association, Washington, DC, 2007)). In some instances, the period of
time can be a year or less than a year; in some instances, the period of time can be longer
than a year (e.g., two years, five years, ten years).

Diagnosis is defined by the APA as the “process of identifying and determining
the nature of a disease or disorder by its signs and symptoms, through the use of
20 assessment techniques (e.g., tests and examinations) and other available evidence” (APA
Dictionary of Psychology, (American Psychological Association, Washington, DC,
2007)). A diagnosis can refer to the present time period, or to a time period in the past or
the future.

Likewise, prognosis is defined by the APA as “a prediction of the course,
25 duration, severity, and outcome of a condition, disease, or disorder” (APA Dictionary of
Psychology, (American Psychological Association, Washington, DC, 2007)). A
prognosis can be made, for example, over a period of one month, six months, one year,
five years, ten years, or longer.

Risk assessment is defined as a “study of a subject done for the purpose of trying
30 to determine the probability that that person will develop a particular disease or, if the
disease is already present, the probability that the person will suffer exacerbation of it or
death from it” (Youngson, 2005, Collins Dictionary of Medicine). In some instances, risk

assessment is based on conditions or events and not directly on disease. In some instances, a risk assessment is determined over a period of time (e.g., months, years).

Biomarkers are described herein that can be used in methods of detecting whether or not a subject is predisposed to diabetes. The methods described herein can be used to predict the likelihood for an individual to develop diabetes and/or prognosticate as to the severity of the disease or the outcome for the individual. Such methods typically include providing a biological sample from the subject; contacting DNA from the biological sample with bisulfite under alkaline conditions; contacting the bisulfite-treated DNA with at least one first oligonucleotide primer at least 8 nucleotides in length that is complementary to a sequence that comprises a CpG dinucleotide (e.g., at a GC locus referred to as cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312); and determining the methylation status of the CpG dinucleotide. It would be understood that the at least one first oligonucleotide probe can detect either the unmethylated CpG dinucleotide or the methylated CpG dinucleotide. Such a method using genomic or bisulfite-treated DNA can further include determining the genotype of a single nucleotide polymorphism (SNP) (e.g., rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855) or a second SNP in linkage disequilibrium with the first SNP. As described herein, methylation of one or more particular CpG dinucleotides and the presence of one or more particular SNPs can be used to predict whether or not the subject has diabetes or has an increased likelihood of developing diabetes.

In some embodiments, the method further comprises contacting the bisulfite-treated DNA with at least one second oligonucleotide probe at least 8 nucleotides in length that is complementary to a sequence that comprises a CpG dinucleotide, where the at least one second oligonucleotide probe detects either the unmethylated CpG dinucleotide or the methylated CpG dinucleotide, whichever is not detected by the at least one first oligonucleotide probe.

In some embodiments, the ratio of methylated CpG dinucleotides to unmethylated CpG dinucleotides in the biological sample can be determined as a part of the methods described herein. Determining the ratio of methylated CpG dinucleotides to unmethylated CpG dinucleotides at a particular locus or genome-wide can be used, for example, to assess the level of transcriptional activity and/or gene expression locally or globally.

It would be appreciated that determining the methylation status of the one or more CpG dinucleotides and determining the presence (or absence) of a SNP can utilize any number of techniques, such as, for example, amplifying and/or sequencing steps. Amplifying and sequencing are well known techniques in the art and are used routinely to
5 determine both the methylation status of a particular sequence and the presence / absence of a SNP.

Methods of determining the presence of biomarkers associated with diabetes in a biological sample from a subject are provided. Such methods typically include providing a first portion of the biological sample and contacting DNA from the first portion with
10 bisulfite under alkaline conditions. The bisulfite-treated first portion can be contacted with a first oligonucleotide probe that is at least 8 nucleotides in length and that is complementary to a sequence that comprises a CpG dinucleotide (detected, e.g., at a CG locus referred to as cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312), and a second portion of the biological sample can be contacted with a
15 nucleic acid probe at least 8 nucleotides in length that is complementary to a SNP (e.g., rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855).

As described herein, the percentage of methylation of the CpG dinucleotide at one or more of the GC loci designated cg06500161, cg21139312, cg19693031, cg17901584, or cg13375427 (or at a CpG that is in linkage disequilibrium with one or more of such
20 CpGs) and the identity of the nucleotide at one or more SNPs designated rs41480347, rs2141638, rs6684261, and rs1423855 (or at a SNP that is in linkage disequilibrium with one or more of such SNPs) are biomarkers associated with diabetes and can be used to predict the likelihood that an individual will develop diabetes and/or prognosticate as to the severity of the disease or the outcome for the individual.

25 While the effects of the indicated loci on whether or not an individual has or will develop diabetes is a complex relationship, the following trends can be associated with diabetes: increasing methylation of the CpG dinucleotide at loci cg06500161 and cg21139312; decreasing methylation of the CpG dinucleotide at loci cg19693031, cg17901584, and cg13375427; the presence of a G nucleotide at SNP rs1834967, the
30 presence of an A nucleotide at SNP rs41480347, the presence of a G nucleotide at SNP rs2141638, the presence of a C nucleotide at SNP rs6684261, and the presence of a C nucleotide at SNP rs1423855.

Kits for Detecting Diabetes

In a further embodiment of the disclosure, articles of manufacture and kits containing probes, oligonucleotides or antibodies are provided. Such articles of manufacture can be used in the methods described herein. An article of manufacture can include one or more containers with, for example, a label. Suitable containers include, for example, bottles, vials, and test tubes. The containers can be formed from a variety of materials such as glass or plastic. The container can hold a composition that includes one or more agents that are effective for practicing the methods described herein. The label on the container indicates that the composition can be used for a specific application. The kit of the disclosure will typically comprise the container described above and one or more other containers comprising materials desirable from a commercial and user standpoint, including buffers, diluents, filters and package inserts with instructions for use.

In certain embodiments, the present disclosure provides a kit for determining the methylation status of at least one CpG dinucleotide and the presence of at least one single-nucleotide polymorphism (SNP). In certain embodiments, a kit as described herein may contain a number of primers that is any integer between 1 and 10,000, such as 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, . . . 9997, 9998, 9999, 10,000. As used herein, the term “nucleic acid primer” or “nucleic acid probes” or “oligonucleotide” encompasses both DNA and RNA sequences. In certain embodiments, the primers or probes may be physically located on a single solid substrate or on multiple substrates.

A kit as described herein can include at least one first nucleic acid primer (e.g., at least 8 nucleotides in length) that is complementary to a bisulfite-converted nucleic acid sequence comprising a CpG dinucleotide (detected, e.g., at a GC locus referred to as cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312), and at least one second nucleic acid primer (e.g., at least 8 nucleotides in length) that is complementary to a SNP (e.g., rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855). The at least one first nucleic acid primer can detect the methylated or unmethylated CpG dinucleotide.

It would be appreciated that any of the nucleic acid primers, probes or oligonucleotides described herein can include one or more nucleotide analogs and/or one or more synthetic or non-natural nucleotides.

It also would be appreciated that the kits described herein can include a solid substrate. In some embodiments, one or more of the nucleic acid primers can be bound to

the solid support. Examples of solid supports include, without limitation, polymers, glass, semiconductors, papers, metals, gels or hydrogels. Additional examples of solid supports include, without limitation, microarrays or microfluidics cards.

5 It also would be appreciated that any of the kits described herein can include one or more detectable labels. In some embodiments, one or more of the nucleic acid primers can be labeled with the one or more detectable labels. Representative detectable labels include, without limitation, an enzyme label, a fluorescent label, and a colorimetric label.

Algorithm for Predicting Diabetes

10 Any number of algorithms that can capture linear effects (e.g., linear regression) or both linear and non-linear effects (e.g., Random Forest, Gradient Boosting, Neural Networks (e.g., deep neural network, extreme learning machine (ELM)), Support Vector Machine, Hidden Markov model) can be used in the methods described herein. See, for example, McKinney et al., 2011, Appl. Bioinform., 5(2):77-88; Gunther et al., 2012, 15 BMC Genet., 13:37; and Ogutu et al., 2011, BMC Proceedings, 5(Suppl 3):S11. Any type of machine learning algorithm or deep learning neural network algorithm (tuned or non-tuned) capable of capturing linear and/or non-linear contribution of traits for the prediction can be used. In some instances, a combination of algorithms (e.g., a combination or ensemble of multiple algorithms that capture linear and/or non-linear 20 contributions of traits) is used.

Simply by way of example, Random Forest™ is a popular machine learning algorithm created by Breiman & Cutler for generating “classification trees” (see, for example, “stat.berkeley.edu/~breiman/RandomForests/cc_home.htm” on the World Wide Web). Using standard machine learning and predictive modeling techniques, a diagnostic 25 classifier algorithm was written to be implemented in R and Python programming languages (though it can be implemented in many other programming languages), according to well described guidelines by Breiman & Cutler. A diagnostic classifier algorithm was generated using data from at least two traits (T) and the diagnosis of interest from that population. To determine the output (e.g., diagnosis) for a new 30 individual, one simply determines values for the at least two traits (T) and inputs that information into an algorithm (e.g., the diagnostic classifier algorithm described herein or another algorithm discussed above) that is capable of capturing the linear and non-linear contributions of the traits.

As described herein, the inputs are at least one genotype (e.g., SNP) and the methylation status of at least one CpG dinucleotide, and the outcome can represent a positive or a negative probability (e.g., prediction or diagnosis) for diabetes. The Traits (T) used to determine the outcome can represent the methylation status of at least one CpG dinucleotide or at least one genotype (e.g., of a SNP), but Traits (T) also can correspond to at least one interaction (e.g., between methylation status and genotype (CpGxSNP), between the methylation status of two different sites (CpGxCpG) or between two different genotypes (SNPxSNP)). It would be appreciated that any such interactions can be visualized using partial dependence plots.

FIG. 3 is a block diagram of an example diabetes classification system 300. In some embodiments, the system 300 can perform monitoring and/or prediction of diabetes. For example, the system 300 can be used to perform one or more of the example processes described herein.

In the illustrated example, a subject 301 provides a subject sample 302. In some embodiments, the subject sample 302 can be, e.g., a blood sample, a saliva sample, or any other appropriate biological sample from the subject 301. In some embodiments, medical personnel 303 (e.g., a doctor, a nurse, a lab technician, a caregiver) may assist the subject 301 with obtaining the subject sample 302. In some embodiments, the subject 301 may obtain the subject sample 302 from herself or himself (e.g., by using a portable blood sampling device or a home collection kit).

A nucleic acid isolation module 310 isolates a nucleic acid sample 312 from the subject sample 302. In some embodiments, the nucleic acid isolation module 310 can be a manual, semi-automated, or automatic process that perform or more of cell lysis, removal of contaminating proteins, deactivating DNAases and/or RNAases, and recovery of DNA and/or RNA. For example, the nucleic acid isolation module 310 can be a part of an automated process or analysis device configured to isolate the nucleic acid sample 312 from the subject sample 302. In another example, the nucleic acid isolation module 310 can be part of one or more of the example kits described in this document, to be used by a human user such as the medical personnel 303.

A genotyping assay module 320 receives a portion 314a of the nucleic acid sample 312. The genotyping assay module 320 is configured to perform a genotyping assay on the portion 314a of the nucleic acid sample 312 to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second

SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to determine, identify, or otherwise obtain a collection of genotype data 322. In some embodiments, the genotyping assay module 320 can be a manual, semi-automated, or automatic process.

5 For example, genotyping assay module 320 can be a part of an automated process or analysis device configured to perform a genotyping assay on the portion 314a. In another example, the genotyping assay module 320 can be part of one or more of the example kits described in this document, to be used by a human user such as the medical personnel 303 or a laboratory technician.

10 A methylation assay module 330 receives a portion 314b of the nucleic acid sample 312. The methylation assay module 330 is configured to bisulfite convert the nucleic acid in the portion 314b of the nucleic acid sample 312 and perform methylation assessment on the portion 314b of the nucleic acid sample 312 to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to determine, identify, or otherwise obtain a collection of
15 methylation data 332.

An identification system 340 is configured to receive the collection of genotype data 322 and the collection of methylation data 332, and identify one or more predetermined traits or characteristics of the subject 301 based on a diagnostic classifier algorithm module 342. The diagnostic classifier algorithm module 342 is configured to
20 account for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect. In some embodiments, the diagnostic classifier algorithm module 342 can perform one or more of the algorithms described herein that may indicate the presence of disease (e.g., diagnostic indicators) or a propensity to develop disease (e.g., predict). For example, the identification system may be configured to identify
25 genetic and/or environmental characteristics that determines the presence of or the likelihood of a subject developing disease (e.g., diabetes), even when the disease is of polygenic origin. In some implementations, the diagnostic classifier algorithm module 342 can be a machine learning algorithm capable of accounting for linear and non-linear effects.

30 The identification system 340 provides an output 350 based on the diagnostic and/or prognostic indicators provided by the diagnostic classifier algorithm module 342. In some embodiments, the identification system 340 can include an output module configured to provide the output 350. In some implementations, the output 350 can be an

identification of one or more diseases that the subject 301 may already have. For example, the output 350 may indicate that traits that are indicative of the presence of diabetes were found in the subject 301. In some implementations, the output 350 can be an indication of a likelihood that the subject 301 may develop a disease within a
5 predetermined time frame (e.g., the subject 301 may have a 43% chance of developing diabetes within 3 years, the subject 301 may have a 77% of having a heart attack within 2 years). In some implementations, the output 350 can include therapeutic and/or preventative recommendations based on the diagnostic and/or prognostic indicators provided by the diagnostic classifier algorithm module 342. For example, in response to
10 an identification or prediction of a diabetic condition in the subject 301, the output 350 may include a recommendation to consult with the medical personnel 303, identify possible dietary or lifestyle changes by the subject 301 to address or avoid the condition, identify potential treatments and/or remedies for the subject 301 to consider in consultation with the medical personnel 303, or combinations of these and/or any other
15 appropriate information based on the output of the algorithm(s) of the diagnostic classifier algorithm module 342.

In the illustrated example, the output 350 is provided in various formats. The information provided by the output 350 can be formatted into a message 360 that is provided to the subject 301 and/or to the medical personnel 303. In some
20 implementations, the message 360 can be formatted as a report (e.g., a word processing file, a portable document format file) that is at least temporarily stored on a non-transitory storage medium (e.g., a hard drive, a FLASH memory), where it can be retrieved by the subject 301 and/or the medical personnel 303 for review. In some implementations, the message 360 can be formatted as an electronic message (e.g., an email, a text message, an
25 instant message) that is transmitted to the subject 301 and/or the medical personnel 303 for review. In some implementations, the message 360 can be a printed report. For example, the output 350 can be provided to a printing system that is configured to generate a hard copy report based on the output 350. Subsequent automated or manual processing systems can package the report as a letter or other parcel that can be sent for
30 physical delivery to the subject 301 and/or to the medical personnel 303 (e.g., the system 300 can create a paper printout of the results and mail them through postal mail).

A treatment device 370 can be configured to receive the diagnostic and/or prognostic indicators provided by the output 350 and provide therapy and/or treatment

based on the diagnostic and/or prognostic indicators. For example, the output 350 may indicate that the subject 301 has diabetes, and the treatment device 370 may be insulin or another drug delivered e.g., directly (e.g., tablets or capsules) or via a syringe or a wearable insulin pump that reacts by identifying or receiving configuration settings for an appropriate dosage of insulin to treat the condition and/or activating a pump or plunger or other mechanism to administer the dosage to the subject 301. In another example, the output 350 may indicate that the subject 301 has a high likelihood of the onset of diabetes within the next two years, three years or five years, and the treatment device 370 may be a continuous glucose monitoring system that reacts by identifying or by receiving configuration settings for an appropriate dose of insulin or any other appropriate therapeutic and/or preventative substances. In some embodiments, the treatment device 370 can be configured to also include one or more of the nucleic acid isolation module 310, the genotyping assay module 320, the methylation assay module 330, and/or the identification system 340.

A storage system 380 is configured to store the output 350. For example, the information included in the output 350 can be stored temporarily, for a predetermined period of time, or substantially permanently in a database, in a file, or as any other appropriate collection of data. In some embodiments, the storage system 380 can store the output 350 in a non-transitory storage medium (e.g., a hard drive, a FLASH memory). For example, the output 350 may include some or all of the collection of genotype data 322, the collection of methylation data 332, and/or the output 350 in personal health record that the subject 301 can store or carry with them. In some embodiments, the storage system 380 can store the output 350 as a physical medium, for example, the storage system 380 can include a printer that can generate a paper report based on the output 350, and/or store the report as a hard copy that can be physically filed away for later retrieval.

An input/output device 382 is physical device configured to display or otherwise present an output that is perceptible to humans (e.g., the subject 301, the medical personnel 303). For example, the input/output device 382 may be an electronic display device in a doctor's office. The system 300 may process the subject sample 302, and then alter the configuration of pixels onscreen to modify the information displayed by the input/output device 382 based on the output 350 (e.g., a screen can be updated to display an identified diagnosis and/or prognosis for the subject 301 to the medical personnel). In another example, the input/output device 382 can be configured to provide audible (e.g.,

spoken output) and/or tactile (e.g., braille, haptic, vibratory) output that modifies or otherwise transforms the output 350 into a physical and/or tangible output (e.g., to convey the diagnostic and/or prognostic indicators in a manner that is perceptible to a user who is sight-challenged). In another example, the input/output device 382 can be configured to
5 alter, transform, or modify a physical characteristic of a physical structure or medium based on the output 350.

A user device 384 (e.g., a computer, a smartphone, a tablet computer, a computerized terminal) is configured to display, emit, or otherwise present one or more outputs that are perceptible to a human user, such as the subject 301 and/or the medical
10 personnel 303. For example, the user device 384 can receive the output 350 (e.g., as data, as the message 360) and provide an alert to the user and/or provide an output (e.g., display a report, read a report aloud) based on the output 350. In some embodiments, the user device 384 can include one or more of the storage device 380 or the input/output device 382. In some embodiments, the user device 382 can be part of the treatment device
15 370. In some embodiments, the user device 384 can be configured to include one or more of the nucleic acid isolation module 310, the genotyping assay module 320, the methylation assay module 330, or the identification system 340.

In some implementations, some or all of the system 300 may be reused to provide additional information. For example, the system 300 may be used to gather an initial set
20 of health information for the subject 301 and/or identify information that can assist the medical personnel 303 with an initial diagnosis/prognosis. Later, the patient 301 may be re-examined using the system 300, for example, to determine the effectiveness of prescribed medical and/or lifestyle strategies over time. Since the collection of genotype data 322 does not change over time for an individual person, the system 300 may refrain
25 from performing the functions of the genotyping assay module 320 again. In such examples, the methylation assay module 330 may be used to generate an updated version of the collection of methylation data 332, and the updated collection of methylation data 332 can be provided to the identification system 340 for processing along with the collection of genotype data 322 that was previously generated. In some implementations,
30 the subject sample 302 can be collected on a periodic basis and processed based on the existing collection of genotype data 322 and updated collections of methylation data 332 to produce updated outputs 350 that can be used to provide ongoing monitoring of one or more conditions identified for the subject 301.

FIG. 4 is a flow diagram of an example process 400 for diabetes classification. In some implementations, the process 400 can be some or all of the example processes described above. In some implementations, the process 400 can be the process performed by some or all of the example system 300 of FIG. 3.

5 At 410, a nucleic acid sample is isolated from a subject sample. For example, the example nucleic acid isolation module 310 can be configured to isolate and/or substantially purify nucleic acid compositions from the example subject sample 302 to produce the example nucleic acid sample 312.

10 At 420, a genotyping assay is performed on a first portion of the nucleic acid sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain genotype data. For example, the example genotyping assay module 320 could be used to analyze the example portion 314a of the nucleic acid sample 312 to produce the example collection of genotype data 322.

15 At 430, a second portion of the nucleic acid sample is bisulfite converted, and a methylation assessment is performed on the second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data. For example, the example methylation assay module 330 can be used to process the portion 314b of the nucleic acid sample 312 to produce the example collection of methylation data 332.

20 At 440, the genotype data from step 420 and/or methylation data from step 430 is input into an algorithm. For example, the example collection of genotype data 322 and the example collection of methylation data 332 are input into the example identification system 340 and processed using the example diagnostic classifier algorithm module 342.

25 At 450, at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect are accounted for. For example, the example diagnostic classifier algorithm module 342 can be configured to account for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect. In some implementations, the diagnostic classifier algorithm module 342, can be a machine learning algorithm capable of accounting for linear and non-linear effects.

30 At 460, an output is provided. For example, the example identification system 340 can provide the example output 350.

At 470 another nucleic acid sample is isolated from another sample from the subject. For example, the example nucleic acid isolation module 310 can be configured to isolate and/or substantially purify nucleic acid compositions from another sample to produce another example nucleic acid sample. Since the collection of genotype data 322 from a subject does not change over time, the newly-produced nucleic acid sample can be used to obtain methylation data 332, which is used along with the existing collection of genotype data 322 to provide an updated output (e.g., to perform a checkup on the subject 301 at a later point in time). In some implementations, this abbreviated process can be performed on a periodic or semi-periodic basis to provide ongoing monitoring of one or more medical conditions identified for the subject 301.

FIG. 5 is a block diagram of example computing devices 500, 550 that may be used to implement the systems and methods described in this document, either as a client or as a server or plurality of servers. Computing device 500 is intended to represent various forms of digital computers, such as laptops, desktops, workstations, personal digital assistants, servers, blade servers, mainframes, and other appropriate computers. Computing device 500 can also represent all or parts of various forms of computerized devices, such as embedded digital controllers, media bridges, modems, network routers, network access points, network repeaters, and network interface devices including mesh network communication interfaces. Computing device 550 is intended to represent various forms of mobile devices, such as personal digital assistants, cellular telephones, smartphones, and other similar computing devices. The components shown here, their connections and relationships, and their functions, are meant to be exemplary only, and are not meant to limit implementations of the inventions described and/or claimed in this document.

Computing device 500 includes a processor 502, a memory 504, a storage device 506, a high-speed interface 508 connecting to memory 504 and high-speed expansion ports 510, and a low speed interface 512 connecting to a low speed bus 514 and storage device 506. Each of the components 502, 504, 506, 508, 510, and 512, are interconnected using various busses, and may be mounted on a common motherboard or in other manners as appropriate. The processor 502 can process instructions for execution within the computing device 500, including instructions stored in the memory 504 or on the storage device 506 to display graphical information for a GUI on an external input/output device, such as display 516 coupled to high speed interface 508. In other

implementations, multiple processors and/or multiple buses may be used, as appropriate, along with multiple memories and types of memory. Also, multiple computing devices 500 may be connected, with each device providing portions of the necessary operations (e.g., as a server bank, a group of blade servers, or a multi-processor system).

5 The memory 504 stores information within the computing device 500. In one implementation, the memory 504 is a computer-readable medium. In one implementation, the memory 504 is a volatile memory unit or units. In another implementation, the memory 504 is a non-volatile memory unit or units.

10 The storage device 506 is capable of providing mass storage for the computing device 500. In one implementation, the storage device 506 is a computer-readable medium. In various different implementations, the storage device 506 may be a floppy disk device, a hard disk device, an optical disk device, or a tape device, a flash memory or other similar solid state memory device, or an array of devices, including devices in a storage area network or other configurations. In one implementation, a computer program
15 product is tangibly embodied in an information carrier. The computer program product contains instructions that, when executed, perform one or more methods, such as those described above. The information carrier is a computer- or machine-readable medium, such as the memory 504, the storage device 506, or memory on processor 502.

20 The high speed controller 508 manages bandwidth-intensive operations for the computing device 500, while the low speed controller 512 manages lower bandwidth-intensive operations. Such allocation of duties is exemplary only. In one implementation, the high-speed controller 508 is coupled to memory 504, display 516 (e.g., through a graphics processor or accelerator), and to high-speed expansion ports 510, which may accept various expansion cards (not shown). In the implementation, low-speed controller
25 512 is coupled to storage device 506 and low-speed expansion port 517 through the low-speed bus 514. The low-speed expansion port, which may include various communication ports (e.g., Universal Serial Bus (USB), BLUETOOTH, BLUETOOTH Low Energy (BLE), Ethernet, wireless Ethernet (WiFi), High-Definition Multimedia Interface (HDMI), ZIGBEE, visible or infrared transceivers, Infrared Data Association (IrDA),
30 fiber optic, laser, sonic, ultrasonic) may be coupled to one or more input/output devices, such as a keyboard, a pointing device, a scanner, a networking device such as a gateway, modem, switch, or router, e.g., through a network adapter 513.

Peripheral devices can communicate with the high speed controller 508 through one or more peripheral interfaces of the low speed controller 512, including but not limited to a USB stack, an Ethernet stack, a WiFi radio, a BLUETOOTH Low Energy (BLE) radio, a ZIGBEE radio, an HDMI stack, and a BLUETOOTH radio, as is
5 appropriate for the configuration of the particular sensor. For example, a sensor that outputs a reading over a USB cable can communicate through a USB stack.

The network adapter 513 can communicate with a network 515. Computer networks typically have one or more gateways, modems, routers, media interfaces, media bridges, repeaters, switches, hubs, Domain Name Servers (DNS), and Dynamic Host
10 Configuration Protocol (DHCP) servers that allow communication between devices on the network and devices on other networks (e.g. the Internet). One such gateway can be a network gateway that routes network communication traffic among devices within the network and devices outside of the network. One common type of network communication traffic that is routed through a network gateway is a Domain Name
15 Server (DNS) request, which is a request to the DNS to resolve a uniform resource locator (URL) or uniform resource indicated (URI) to an associated Internet Protocol (IP) address.

The network 515 can include one or more networks. The network(s) may provide for communications under various modes or protocols, such as Global System for Mobile
20 communication (GSM) voice calls, Short Message Service (SMS), Enhanced Messaging Service (EMS), or Multimedia Messaging Service (MMS) messaging, Code Division Multiple Access (CDMA), Time Division Multiple Access (TDMA), Personal Digital Cellular (PDC), Wideband Code Division Multiple Access (WCDMA), CDMA2000, General Packet Radio System (GPRS), or one or more television or cable networks,
25 among others. For example, the communication may occur through a radio-frequency transceiver. In addition, short-range communication may occur, such as using a BLUETOOTH, BLE, ZIGBEE, WiFi, IrDA, or other such transceiver.

In some embodiments, the network 515 can have a hub-and-spoke network configuration. A hub-and-spoke network configuration can allow for an extensible
30 network that can accommodate components being added, removed, failing, and replaced. This can allow, for example, more, fewer, or different devices on the network 515. For example, if a device fails or is deprecated by a newer version of the device, the network

515 can be configured such that network adapter 513 can to be updated about the replacement device.

In some embodiments, the network 515 can have a mesh network configuration (e.g., ZIGBEE). Mesh configurations may be contrasted with conventional star/tree network configurations in which the networked devices are directly linked to only a small subset of other network devices (e.g., bridges/switches), and the links between these devices are hierarchical. A mesh network configuration can allow infrastructure nodes (e.g., bridges, switches and other infrastructure devices) to connect directly and non-hierarchically to other nodes. The connections can be dynamically self-organize and self-configure to route data. By not relying on a central coordinator, multiple nodes can participate in the relay of information. In the event of a failure of one or more of the nodes or the communication links between them, the mesh network can self-configure to dynamically redistribute workloads and provide fault-tolerance and network robustness.

The computing device 500 may be implemented in a number of different forms, as shown in the figure. For example, it may be implemented as a standard server 520, or multiple times in a group of such servers. It may also be implemented as part of a rack server system 524. It may also be implemented as part of network device such a modem, gateway, router, access point, repeater, mesh node, switch, hub, or security device (e.g., camera server). In addition, it may be implemented in a personal computer such as a laptop computer 522. Alternatively, components from computing device 500 may be combined with other components in a mobile device (not shown), such as device 550. In some embodiments, the device 550 can be a mobile telephone (e.g., a smartphone), a handheld computer, a tablet computer, a network appliance, a camera, an enhanced general packet radio service (EGPRS) mobile phone, a media player, a navigation device, an email device, a game console, an interactive or so-called “smart” television, a media streaming device, or a combination of any two or more of these data processing devices or other data processing devices. In some implementations, the device 550 can be included as part of a motor vehicle (e.g., an automobile, an emergency vehicle (e.g., fire truck, ambulance), a bus). Each of such devices may contain one or more of computing device 500, 550, and an entire system may be made up of multiple computing devices 500, 550 communicating with each other through a low speed bus or a wired or wireless network.

Computing device 550 includes a processor 552, memory 564, an input/output device such as a display 554, a communication interface 566, and a transceiver 568, among other components. The device 550 may also be provided with a storage device, such as a microdrive or other device, to provide additional storage. Each of the
5 components 550, 552, 564, 554, 566, and 568, are interconnected using various buses, and several of the components may be mounted on a common motherboard or in other manners as appropriate.

The processor 552 can process instructions for execution within the computing device 550, including instructions stored in the memory 564. The processor may also
10 include separate analog and digital processors. The processor may provide, for example, for coordination of the other components of the device 550, such as control of user interfaces, applications run by device 550, and wireless communication by device 550.

Processor 552 may communicate with a user through control interface 558 and display interface 556 coupled to a display 554. The display 554 may be, for example, a
15 TFT LCD display or an OLED display, or other appropriate display technology. The display interface 556 may comprise appropriate circuitry for driving the display 554 to present graphical and other information to a user. The control interface 558 may receive commands from a user and convert them for submission to the processor 552. In addition, an external interface 562 may be provide in communication with processor 552, so as to
20 enable near area communication of device 550 with other devices. External interface 562 may provide, for example, for wired communication (e.g., via a docking procedure) or for wireless communication (e.g., via Bluetooth or other such technologies).

The memory 564 stores information within the computing device 550. In one implementation, the memory 564 is a computer-readable medium. In one implementation,
25 the memory 564 is a volatile memory unit or units. In another implementation, the memory 564 is a non-volatile memory unit or units. Expansion memory 574 may also be provided and connected to device 550 through expansion interface 572, which may include, for example, a SIMM card interface. Such expansion memory 574 may provide extra storage space for device 550, or may also store applications or other information for
30 device 550. Specifically, expansion memory 574 may include instructions to carry out or supplement the processes described above, and may include secure information also. Thus, for example, expansion memory 574 may be provide as a security module for device 550, and may be programmed with instructions that permit secure use of device

550. In addition, secure applications may be provided via the SIMM cards, along with additional information, such as placing identifying information on the SIMM card in a non-hackable manner.

The memory may include for example, flash memory and/or MRAM memory, as discussed below. In one implementation, a computer program product is tangibly embodied in an information carrier. The computer program product contains instructions that, when executed, perform one or more methods, such as those described above. The information carrier is a computer- or machine-readable medium, such as the memory 564, expansion memory 574, or memory on processor 552.

Device 550 may communicate wirelessly through communication interface 566, which may include digital signal processing circuitry where necessary. Communication interface 566 may provide for communications under various modes or protocols, such as GSM voice calls, Voice Over LTE (VOLTE) calls, SMS, EMS, or MMS messaging, CDMA, TDMA, PDC, WCDMA, CDMA2000, GPRS, WiMAX, LTE, 5G, among others. Such communication may occur, for example, through radio-frequency transceiver 568. In addition, short-range communication may occur, such as using a Bluetooth, WiFi, or other such transceiver (not shown) configured to provide uplink and/or downlink portions of data communication. In addition, GPS receiver module 570 may provide additional wireless data to device 550, which may be used as appropriate by applications running on device 550.

Device 550 may also communicate audibly using audio codec 560, which may receive spoken information from a user and convert it to usable digital information. Audio codec 560 may likewise generate audible sound for a user, such as through a speaker, e.g., in a handset of device 550. Such sound may include sound from voice telephone calls, may include recorded sound (e.g., voice messages, music files, etc.) and may also include sound generated by applications operating on device 550.

The computing device 550 may be implemented in a number of different forms, as shown in the figure. For example, it may be implemented as a cellular telephone 580. It may also be implemented as part of a smartphone 582, personal digital assistant, or other similar mobile device.

Various implementations of the systems and techniques described here can be realized in digital electronic circuitry, integrated circuitry, specially designed ASICs (application specific integrated circuits), computer hardware, firmware, software, and/or

combinations thereof. These various implementations can include implementation in one or more computer programs that are executable and/or interpretable on a programmable system including at least one programmable processor, which may be special or general purpose, coupled to receive data and instructions from, and to transmit data and instructions to, a storage system, at least one input device, and at least one output device.

These computer programs (also known as programs, software, software applications or code) include machine instructions for a programmable processor, and can be implemented in a high-level procedural and/or object-oriented programming language, and/or in assembly/machine language. As used herein, the terms “machine-readable medium” “computer-readable medium” refers to any computer program product, apparatus and/or device (e.g., magnetic discs, optical disks, memory, Programmable Logic Devices (PLDs)) used to provide machine instructions and/or data to a programmable processor, including a machine-readable medium that receives machine instructions as a machine-readable signal. The term “machine-readable signal” refers to any signal used to provide machine instructions and/or data to a programmable processor.

To provide for interaction with a user, the systems and techniques described here can be implemented on a computer having a display device (e.g., a CRT (cathode ray tube) or LCD (liquid crystal display) monitor) for displaying information to the user and a keyboard and a pointing device (e.g., a mouse or a trackball) by which the user can provide input to the computer. Other kinds of devices can be used to provide for interaction with a user as well; for example, feedback provided to the user can be any form of sensory feedback (e.g., visual feedback, auditory feedback, or tactile feedback); and input from the user can be received in any form, including acoustic, speech, or tactile input.

The systems and techniques described here can be implemented in a computing system that includes a back end component (e.g., as a data server), or that includes a middleware component (e.g., an application server), or that includes a front end component (e.g., a client computer having a graphical user interface or a Web browser through which a user can interact with an implementation of the systems and techniques described here), or any combination of such back end, middleware, or front end components. The components of the system can be interconnected by any form or medium of digital data communication (e.g., a communication network). Examples of

communication networks include a local area network (“LAN”), a wide area network (“WAN”), and the Internet.

Some communication networks can be configured to carry power as well as information on the same physical media. This allows a single cable to provide both data
5 connection and electric power to devices. Examples of such shared media include power over network configurations in which power is provided over media that is primarily or previously used for communications. One specific embodiment of power over network is Power Over Ethernet (POE) which pass electric power along with data on twisted pair Ethernet cabling. Examples of such shared media also include network over power
10 configurations in which communication is performed over media that is primarily or previously used for providing power. One specific embodiment of network over power is Power Line Communication (PLC) (also known as power-line carrier, power-line digital subscriber line (PDSL), mains communication, power-line telecommunications, or power-line networking (PLN), Ethernet-Over-Power (EOP)) in which data is carried on a
15 conductor that is also used simultaneously for AC electric power transmission.

The computing system can include clients and servers. A client and server are generally remote from each other and typically interact through a communication network. The relationship of client and server arises by virtue of computer programs running on the respective computers and having a client-server relationship to each other.

20 The computing system can include routers, gateways, modems, switches, hub, bridges, and repeaters. A router is a networking device that forwards data packets between computer networks and performs traffic directing functions. A network switch is a networking device that connects networked devices together by performing packet switching to receive, process, and forward data to destination devices. A gateway is a
25 network device that allows data to flow from one discrete network to another. Some gateways can be distinct from routers or switches in that they can communicate using more than one protocol and can operate at one or more of the seven layers of the open systems interconnection model (OSI). A media bridge is a network device that converts data between transmission media so that it can be transmitted from computer to computer.
30 A modem is a type of media bridge, typically used to connect a local area network to a wide area network such as a telecommunications network. A network repeater is a network device that receives a signal and retransmits it to extend transmissions and allow the signal can cover longer distances or overcome a communications obstruction.

It will be apparent that the present disclosure provides a skilled artisan the ability to construct a matrix in which the methylation status of one or more CpG dinucleotides and one or more genotypes (e.g., SNPs; e.g., at one or more alleles) can be evaluated as described herein, typically using a computer, to identify interactions and allow for prediction of diabetes. Although such an analysis is complex, no undue experimentation is required as all necessary information is either readily available to the skilled artisan or can be acquired by experimentation as described herein.

Methods of Treating Diabetes

As described herein, compositions and methods are provided for detecting whether or not a subject is predisposed to diabetes, to predict the likelihood for an individual to develop diabetes, and/or prognosticate as to the severity of the disease or the outcome for the individual. The information obtained from the compositions and methods provided herein can educate the appropriate treatment strategies.

Treatments for diabetes are known in the art and typically depend on the type of diabetes (e.g., type I, type II, adult onset). With respect to type I diabetes, treatments include, for example, insulin delivery and/or cell therapy (e.g., stem cell therapy, beta cell transplantation). It would be appreciated that insulin can be delivered via injection into soft tissues or implantation of an insulin pump, and that the insulin can be a short-acting, rapid-acting, or long-acting insulin. With respect to type II diabetes, treatments include, for example, lifestyle changes (e.g., diet (e.g., low carbohydrate diet), weight loss, exercise, reduction or cessation in smoking and/or drinking), pharmaceuticals (e.g., insulin, alpha-glucosidase inhibitors, biguanides, dopamine agonist, dipeptidyl peptidase-4 (DPP-4) inhibitors, glucagon-like peptide (GLP)-1 receptor agonists, sodium-glucose transporter (SGLT)-2 inhibitors, sulfonylureas, thiazolidinediones) and/or mechanical devices (e.g., insulin pumps, continuous glucose monitor (CGM) systems).

In accordance with the present disclosure, there may be employed conventional molecular biology, microbiology, biochemical, and recombinant DNA techniques within the skill of the art. Such techniques are explained fully in the literature. The invention will be further described in the following examples, which do not limit the scope of the methods and compositions of matter described in the claims.

EXAMPLES

Example 1—Framingham Heart Study

The data used in this study is derived from participants in the Framingham Heart Study (FHS) (Dawber et al., 1963, Ann. N.Y. Acad. Sci., 107:539-56). FHS is a longitudinal study aimed at understanding the risks of cardiovascular disease (CVD) and consists of several cohorts including the Original Cohort, Offspring Cohort, Omni Cohort, Third Generation Cohort, New Offspring Spouse Cohort and Second Generation Omni Cohort. Specifically, the Offspring Cohort, initiated in 1971, consisting of the offspring of the Original Cohort and their spouses was used in this study. This cohort consists of 2,483 males and 2,641 females, for a total of 5,124 individuals (Mahmood et al., 2014, The Lancet, 383:999-1008). The specific analyses described herein were approved by the University of Iowa Institutional Review Board.

Example 2—Genome-Wide DNA Methylation

Of the 5,124 individuals in the Offspring Cohort, only 2,567 individuals (duplicates removed) with DNA methylation data were considered. These individuals were included in the DNA methylation study because they attended the Framingham Offspring 8th exam, provided consent for genetic research, had a buffy coat sample, and had sufficient DNA quantity and quality for methylation profiling. Exam 8 took place between 2005 and 2008. Genomic DNA extracted from their white blood cells was bisulfite converted, then genome-wide DNA methylation was profiled using the Illumina HumanMethylation450 BeadChip (San Diego, CA) at either the University of Minnesota or Johns Hopkins University. The intensity data (IDAT) files of the samples alongside their slide and array information were used to perform the DASEN normalization using the MethyLumi, Watermelon and IlluminaHumanMethylation450k.db R packages (Pidsley et al., 2013, BMC Genomics, 14:1-10). The DASEN normalization performs probe filtering, background correction and adjustment for probe types. Samples were removed if they contained >1% of CpG sites with a detection p-value >0.05. CpG sites were removed if they had a bead count of <3 and/or >1% of samples had a detection p-value >0.05. After DASEN normalization, there were 2,560 samples and 484,241 sites remaining (484,125 CpG sites). CpG sites were grouped by chromosome. Of those CpG sites, 472,822 mapped to autosomes. Methylation beta values were converted to M

values using the beta2m R function in the Lumi package and subsequently converted to z-scores using an R script (Du et al., 2008, Bioinform., 24:1547-8).

Example 3—Genome-Wide Genotype

5 Of the 2,560 remaining individuals after DNA methylation quality control, 2,406 (1,100 males and 1,306 females) had genome-wide genotype data from the Affymetrix GeneChip HumanMapping 500K Array Set (Santa Clara, CA). This array is capable of profiling 500,568 SNPs in the genome. Quality control was performed at both the sample and SNP probe levels in PLINK. The initial quality control step involved identifying
10 individuals with discordant sex information. None were identified. Next, individuals with a heterozygosity rate of greater or smaller than the mean $\pm$ 2SD and with a proportion of missing SNPs >0.03 were excluded. Related individuals were also excluded if the identity by descent value was >0.185 (halfway between second and third degree relatives). After performing these sample level quality control steps, 1,599
15 individuals remained (722 males and 877 females). On the probe level, SNPs with a minor allele frequency $>1\%$, Hardy-Weinberg equilibrium p-value $>10^{-5}$ and SNP missing rate of $<5\%$ were retained. A total of 403,192 SNPs remained after these quality control steps. Using the recode option in PLINK (Purcell et al., 2007, Am. J. Hum. Gen., 81:559-75), genotypes were coded as 0, 1 or 2 per minor allele frequency.

Example 4—Phenotype

For individuals in this study, their percent hemoglobin A1c (HbA1c) at the eighth examination cycle was extracted. Individuals were classified as non-diabetic if their percent HbA1c was $<5.7\%$ and as diabetic if their percent HbA1c was $>6.4\%$. Based on
25 this criterion, 896 were classified as non-diabetic and 130 as diabetic. The demographics of these individuals are summarized in Table 1.

Table 1. Demographics of the 1,026 individuals in the diabetes dataset

	Diabetic	Non-diabetic
n		
Male	70	417
Female	60	479
Age		
Male	67.5 $\pm$ 7.5	66.4 $\pm$ 8.5
Female	69.9 $\pm$ 8.8	65.7 $\pm$ 8.3

Example 5—Variable Reduction

The total number of genetic (SNP) and epigenetic (DNA methylation) probes remaining after quality control measures were 403,192 and 472,822, respectively. Due to the large number of variables (876,014 total, excluding possible interaction between SNPs and DNA methylation sites), and to avoid collinearity, variable reduction was performed.

Linkage disequilibrium based SNP pruning was performed in PLINK (Purcell et al., 2007, Am. J. Hum. Gen., 81:559-75) with a window size of 50 SNPs, window shift of 5 SNPs and a pairwise SNP-SNP LD threshold of 0.5. This reduced the number of SNPs from 403,192 to 161,474. To further reduce the number of SNPs, the chi-squared p-value was calculated between the remaining 161,474 SNPs and CHF and stroke status. Those with a chi-squared p-value <0.1 were retained for classification analyses, resulting in 15,687 SNPs for diabetes.

To reduce the number of DNA methylation loci, first, the point bi-serial correlation was calculated between the 472,822 CpG sites and CHF and stroke status. CpG sites were retained if the point bi-serial correlation was at least 0.1. A total of 24,014 CpG sites remained for diabetes. Subsequently, Pearson correlations between sites were calculated independently for each illness. If the Pearson correlation between two loci were at least 0.8, the loci with a smaller point bi-serial correlation was discarded. In the end, 12,161 DNA methylation loci remained for the classification analyses of diabetes.

Example 6—Receiver Operating Characteristic Curve

A receiver operating characteristic (ROC) curve provides a graphical representation of binary classification performance with varying discrimination thresholds. Therefore, to assess the capability of DNA methylation and SNPs in classifying diabetes, an R script was written to perform logistic regression of the models shown below and subsequently calculate the area under the curve (AUC) of the ROC curve (Beck et al., 1986, Arch. Pathol. Lab. Med., 110:13-20) using the pROC package in R. This was performed systematically using DNA methylation sites that were ordered in descending order of point bi-serial with respect to diabetes and SNPs that were order in

ascending order of chi-squared p-value with respect to diabetes. In the models listed below, SNP*meth term represents the gene-environment interaction.

$$\text{Diabetes} \sim \text{SNP}_i + \text{meth}_i + \text{SNP}_i * \text{meth}_i$$

5 Example 7—ROC of Diabetes Classification

Using the top five DNA methylation sites (cg19693031, cg17901584, cg06500161, cg13375427, cg21139312) and SNPs (rs1834967, rs41480347, rs2141638, rs6684261, rs1423855), a model incorporating only main effects were fitted for diabetes. The ROC AUC was 0.86 and is shown in FIG. 1. The model parameters are summarized in Table 2.

Table 2. Parameters of the main effects diabetes model

Variable	Estimate	Std. Error	z value	Pr(> z)
cg19693031	-0.5655	0.1126	-5.022	5.11E-07
cg17901584	-0.5634	0.1409	-3.999	6.36E-05
cg06500161	0.8421	0.1364	6.175	6.64E-10
cg13375427	-0.4272	0.1385	-3.085	0.00203
cg21139312	0.2458	0.1295	1.898	0.05766
rs1834967	-1.6477	0.4327	-3.808	0.00014
rs41480347	2.268	0.4624	4.905	9.33E-07
rs2141638	0.4721	0.2411	1.958	0.0502
rs6684261	0.5726	0.1821	3.144	0.00166
rs1423855	0.8748	0.3492	2.505	0.01225

To further demonstrate the importance of incorporating both DNA methylation and SNPs in better predicting diabetes status, interaction terms as depicted in the methods section were included in the diabetes model. The ROC AUC for this model increased from the previous model to 0.87 and is shown in FIG. 2. The model parameters are summarized in Table 3.

Table 3. Parameters of the interaction effects diabetes model

Variable	Estimate	Std. Error	z value	Pr(> z)
cg19693031	-0.896117	0.194595	-4.605	4.12E-06
rs1834967	-2.505628	0.822427	-3.047	0.00231
rs41480347	1.909286	0.709135	2.692	0.00709
rs2141638	0.233778	0.353157	0.662	0.50799
rs6684261	0.64591	0.24402	2.647	0.00812

rs1423855	0.301257	0.653121	0.461	0.64461
cg17901584	-0.400774	0.224852	-1.782	0.07469
cg06500161	0.671997	0.252358	2.663	0.00775
cg13375427	-0.448898	0.256536	-1.75	0.08015
cg21139312	0.012049	0.218659	0.055	0.95606
cg19693031:rs1834967	-0.131394	0.612813	-0.214	0.83023
cg19693031:rs41480347	-0.140471	0.925474	-0.152	0.87936
cg19693031:rs2141638	0.074571	0.249454	0.299	0.76499
cg19693031:rs6684261	0.409676	0.183324	2.235	0.02544
cg19693031:rs1423855	0.597921	0.415749	1.438	0.15038
rs1834967:cg17901584	-0.677134	0.697083	-0.971	0.33136
rs41480347:cg17901584	-1.618487	0.902544	-1.793	0.07293
rs2141638:cg17901584	-0.49346	0.356389	-1.385	0.16617
rs6684261:cg17901584	0.065158	0.224121	0.291	0.77126
rs1423855:cg17901584	-0.786052	0.644765	-1.219	0.2228
rs1834967:cg06500161	0.927205	0.689422	1.345	0.17866
rs41480347:cg06500161	1.097323	0.921671	1.191	0.23382
rs2141638:cg06500161	0.234633	0.320307	0.733	0.46385
rs6684261:cg06500161	-0.036366	0.238139	-0.153	0.87863
rs1423855:cg06500161	0.538688	0.68103	0.791	0.42895
rs1834967:cg13375427	0.035485	0.688161	0.052	0.95888
rs41480347:cg13375427	0.162918	0.617926	0.264	0.79205
rs2141638:cg13375427	-0.200623	0.331071	-0.606	0.54453
rs6684261:cg13375427	-0.007325	0.248728	-0.029	0.97651
rs1423855:cg13375427	0.153555	0.545698	0.281	0.77841
rs1834967:cg21139312	0.279818	0.564912	0.495	0.62037
rs41480347:cg21139312	0.357336	0.684866	0.522	0.60184
rs2141638:cg21139312	-0.017348	0.302024	-0.057	0.9542
rs6684261:cg21139312	0.24886	0.204095	1.219	0.22272
rs1423855:cg21139312	0.488248	0.573736	0.851	0.39477

The two models for diabetes show highly significant CpG sites and SNPs that are highly significant at the 0.05 significance level with respect to diabetes status. The tables and AUC values clearly demonstrate the importance of accounting for both genetic and epigenetic effects.

Example 8—Additional Data

Appendix A shows a list of CpGs whose methylation is associated with diabetes.

Appendix B shows a list of genes whose methylation is associated with diabetes.

Appendix C shows a list of SNPs associated with diabetes. The numerical values provided in Appendix A, B, and C are the mean of 5-fold cross validation scores, AUC

ROC (Area Under The Receiver Operating Characteristic Curve), sensitivity and specificity, which were computed by the diabetes assessment / prediction algorithm described herein. Sensitivity is the true positive rate and specificity is the true negative rate.

5

It is to be understood that, while the methods and compositions of matter have been described herein in conjunction with a number of different aspects, the foregoing description of the various aspects is intended to illustrate and not limit the scope of the methods and compositions of matter. Other aspects, advantages, and modifications are within the scope of the following claims.

10

Disclosed are methods and compositions that can be used for, can be used in conjunction with, can be used in preparation for, or are products of the disclosed methods and compositions. These and other materials are disclosed herein, and it is understood that combinations, subsets, interactions, groups, etc. of these methods and compositions are disclosed. That is, while specific reference to each various individual and collective combinations and permutations of these compositions and methods may not be explicitly disclosed, each is specifically contemplated and described herein. For example, if a particular composition of matter or a particular method is disclosed and discussed and a number of compositions or methods are discussed, each and every combination and permutation of the compositions and the methods are specifically contemplated unless specifically indicated to the contrary. Likewise, any subset or combination of these is also specifically contemplated and disclosed.

15

20

A number of embodiments have been described. Nevertheless, it will be understood that various modifications may be made without departing from the spirit and scope of the invention. For example, various forms of the flows shown above may be used, with steps re-ordered, added, or removed. Also, although several applications of the payment systems and methods have been described, it should be recognized that numerous other applications are contemplated. Accordingly, other embodiments are within the scope of the following claims.

25

30

ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg19693031	0.697863954	0.619076923	0.663911846	TXNIP	1	145441552	cg24277313	0.57020386	0.516	0.573553719	PRKAA2	1	57110878
cg15584070	0.661238398	0.572	0.618732782		2	129439854	cg09376129	0.57019962	0.517846154	0.591735537	TMC5	16	19467042
cg18826637	0.657563043	0.58	0.638567493		2	145116633	cg07571531	0.57017249	0.547384615	0.566391185	TCF21	6	134216031
cg036633120	0.651694003	0.673230769	0.602203857	NUDT1	7	2284600	cg12646747	0.5701691	0.538461538	0.565289256	ACOX1	2	111491205
cg26565838	0.637734266	0.618461538	0.604407713	GALNT10	5	153530764	cg05681560	0.57016656	0.524	0.585123967		16	164288814
cg25158622	0.634487815	0.608615385	0.598347107	BRD2	6	32942429	cg25533897	0.57015808	0.523076923	0.571900826	CI9orf63	19	30984040
cg15111351	0.62392117	0.603076923	0.61046832	MAP2K1	15	66687733	cg19593660	0.57015385	0.494461538	0.613774105	FLJ43860	8	142483022
cg20109054	0.623441407	0.643076923	0.553719008	C6orf48	6	31804109	cg2144389	0.57015385	0.492615385	0.592786501	DUOX1	15	45434285
cg01020748	0.622125026	0.658461538	0.530578512	LARK30	18	7231037	cg26835440	0.57013605	0.548	0.56584022		8	55037773
cg20872778	0.620726425	0.656307692	0.519592229	ARCA13	7	48494156	cg04928049	0.57013266	0.508615385	0.598347107		1	19665240
cg20464368	0.620397542	0.618153846	0.542454545	HEATR1	1	236767266	cg23460057	0.57012926	0.400615385	0.642424242	ADRB3	8	37824340
cg21849844	0.619510998	0.547384615	0.628099174	ZNF267	16	31891978	cg16716189	0.57010299	0.485230769	0.587878788	FAM19A3	1	113261711
cg18155407	0.618125026	0.584923077	0.612121212	TFRD12	19	33210662	cg06292124	0.57010045	0.515076923	0.569146006	SLC25A22	11	794867
cg22993042	0.615617716	0.602769231	0.590082645		10	4093925	cg08738600	0.57002161	0.415692308	0.638567493	GRASP	12	52408520
cg02530375	0.61547616	0.579076923	0.599449036		11	69706831	cg03254538	0.57001822	0.478461538	0.566942149	EFSEC	13	127872841
cg23144681	0.614833346	0.570769231	0.609917355	PAM	10	23488710	cg01061697	0.57000551	0.522461538	0.536088154		6	168387030
cg10640944	0.614797203	0.58	0.591184573	IKZF3	5	102201890	cg04573397	0.56999873	0.586153846	0.52892562	TFAP2D	6	50691286
cg20709994	0.614222505	0.617230769	0.544903581	IKZF3	17	379221198	cg02733945	0.56996737	0.531384615	0.58953168	NRS2	1	200011858
cg26403843	0.613612206	0.568923077	0.59338843	RNF145	5	158634085	cg25509174	0.56993431	0.547692308	0.55261708	PCOLCE2	3	142607511
cg24847846	0.613437593	0.648923077	0.503581267		3	12595473	cg09624870	0.56992922	0.539692308	0.567493113	TACC2	10	123871988
cg03070187	0.613095147	0.600615385	0.568595041	TRIP10	19	6739184	cg03116837	0.56989447	0.5562923077	0.601101928		5	50218625
cg05656566	0.61285018	0.609230769	0.582369146	ADAMT52	5	17859324	cg25681872	0.56989277	0.532	0.602203857		8	216703
cg02424359	0.612780674	0.532307692	0.590633609	ZNF844	19	12175639	cg03661703	0.56988854	0.508923077	0.57650854		16	27862708
cg03444077	0.612569612	0.5	0.62369146		10	45359721	cg16754015	0.56988515	0.540307692	0.590633609	ADAR2	10	1415068
cg21198603	0.612563679	0.608615385	0.577961433	C14orf80	14	105957410	cg19107641	0.56986565	0.524	0.597245179		7	1660443
cg06548615	0.612542488	0.544307692	0.639669421	PPGRIP1	16	53737863	cg10914538	0.56983683	0.533538462	0.55927865	NOS1	12	117797056
cg07715298	0.612431447	0.587076923	0.578512397	HDAC4	2	240404019	cg03552039	0.56981988	0.571076923	0.537190083	FGFR2	10	123243495
cg26385126	0.612167832	0.523692308	0.611570248	NCOR2	12	124912021	cg25786640	0.5698131	0.585846154	0.563636364	MAD1L1	7	1946447
cg09815251	0.611552448	0.486461538	0.617630854		5	113029364	cg0414652	0.56976393	0.572923077	0.540495868	NPA15	16	69597925
cg04816311	0.611299004	0.600923077	0.577410468	C7orf50	7	1066650	cg06232266	0.56974105	0.539076923	0.569696997	PEBP4	8	22722736
cg08964190	0.61121085	0.602153846	0.575757576	LRRCL108	11	61275247	cg26701242	0.56971647	0.538461538	0.573002755	SLM2	21	38077971
cg07661704	0.611175249	0.632	0.54214876	SLC7A13	4	139144433	cg18080528	0.56965544	0.555384615	0.551515152	FKBP1	6	32097687
cg00151370	0.61093452	0.568615385	0.584022039	ATXN1	6	16323285	cg03050854	0.56962746	0.524615385	0.591735537	HTB6	1	20005414
cg05647197	0.610903157	0.525230769	0.620936639	KLF11	2	10182525	cg11287055	0.56962492	0.562461538	0.53719008	DSCA3	21	38630234
cg25232221	0.610559864	0.625538462	0.574104683	H1A-E	6	30457562	cg18396987	0.56962323	0.461846154	0.595392287	STC1	1	115398073
cg16742075	0.610459843	0.530769231	0.603856749	H1A-A	6	29911366	cg09343092	0.56956359	0.454153846	0.643526171	HOXA6	7	27185282
cg01013522	0.610252172	0.462461538	0.588429752	CNTNAP5	2	124971544	cg01415527	0.56956389	0.515076923	0.57630854	SBNO2	4	1110591
cg27115863	0.610128417	0.560615385	0.615426997		22	37921640	cg26238771	0.56955287	0.563076923	0.534453262	HCN2	19	592158
cg07726187	0.610026701	0.577615385	0.59338843	TXNDC6	3	138048101	cg17512187	0.56953253	0.548307692	0.5867686	C2orf89	2	95082845
cg18889183	0.609418097	0.562769231	0.585674931	DNMT3A	2	25564471	cg05985821	0.56951727	0.587692308	0.509090909		8	326586
cg18940258	0.609322314	0.533538462	0.633057851	LOC389033	2	130692257	cg20663376	0.56948421	0.532615385	0.564187328	TRPM1	15	31372997
cg17355400	0.609271456	0.554769231	0.586225895		12	46844645	cg16999594	0.56947997	0.531692308	0.578312397		6	169595716

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg06075652	0.609185844	0.594769231	0.581267218		16	86253508	cg23469448	0.56946896	0.547692308	0.562534435		12	114232746
cg05916906	0.608816275	0.611076923	0.534435262	2-Mir	19	8478510	cg27476810	0.56946472	0.532230769	0.562534435	RB1CC1	8	536277771
cg24517430	0.608794236	0.455394615	0.55922865	MESDC1	15	81294299	cg24600724	0.56946387	0.478461538	0.597796143	UNC5A	5	176289552
cg11237792	0.608446705	0.564615385	0.602754821	TTC30A	2	178484058	cg22037593	0.56946133	0.50120769	0.543801653	ANKRD43	5	132148963
cg14444787	0.607708413	0.532	0.6059366391	EIF3H	8	117768111	cg02589828	0.56945963	0.577558462	0.54214876	SAR1B	5	133967994
cg19084031	0.607577877	0.406461538	0.697520661		15	38361382	cg26714388	0.5694946	0.471384615	0.586225895		12	34488729
cg07792822	0.607482941	0.448	0.603856749	KIRREL	1	157961914	cg02067608	0.56946692	0.484	0.582369146		7	25894793
cg19846609	0.607261708	0.611384615	0.555371901		11	124999394	cg14280546	0.56943359	0.563076923	0.553168044	LOC440173	9	89649348
cg07935255	0.607144734	0.649230769	0.509641873	SUGT11	13	41495616	cg05487589	0.56943505	0.565538462	0.547107438	EBF2	8	25896196
cg21475150	0.607119305	0.572	0.577961433	RPL31	2	101618248	cg04163119	0.56943166	0.594461538	0.554820937	PHCKIE4	6	41754006
cg03841977	0.606438803	0.626461538	0.51184573	CBR1	21	37442777	cg24648119	0.56940453	0.517230769	0.597245179	PCDHAG	5	140243639
cg02565702	0.606325069	0.510769231	0.608815427	TACC1	8	38614515	cg26880735	0.5693986	0.470769231	0.594490358		18	44784846
cg01494017	0.606259801	0.640307692	0.569146006		5	1724892	cg18209895	0.56939521	0.570769231	0.539393939	ZFP41	8	144328965
cg01485326	0.605951422	0.600923077	0.536539118	FAM38A	16	88787689	cg14010914	0.56939352	0.532307692	0.531680441	PPP2R5E	14	64009731
cg03739172	0.605705446	0.602769231	0.562534435	PAX7	1	19046707	cg09233380	0.56937995	0.539384615	0.558126722		6	29720748
cg10411146	0.604908667	0.642153846	0.419283747	ATHL1	11	289366	cg02746752	0.56936131	0.485846154	0.594490358		16	1169415
cg14926439	0.604860352	0.60615385	0.579614325	NF2	22	29999402	cg06488842	0.56933757	0.548	0.56969697	PES1	22	30988475
cg135300892	0.604813732	0.563076923	0.608917355	ASC2	11	22992000	cg05713954	0.56928332	0.432615385	0.616528926		16	49518536
cg06904048	0.604662852	0.521846154	0.591184573	CYP212	1	60392328	cg26954671	0.56927654	0.539394615	0.531129477	SLC39A14	8	22223465
cg25266281	0.604506887	0.626461538	0.54214876		2	242904766	cg0441453	0.56926891	0.508	0.655096419		5	41902083
cg10794991	0.604227167	0.524923077	0.588980716		8	141537675	cg03483707	0.56926637	0.589846154	0.509090909	SNORD115-48	15	25513530
cg09034601	0.603981352	0.578769231	0.559779614		6	28175888	cg01969910	0.56925705	0.545230769	0.537190083	GNEB4	3	179169571
cg03198958	0.603627887	0.547692308	0.585123967	TRIM11	4	189061030	cg02518899	0.5692562	0.594769231	0.550964187	KRT14	17	39742965
cg16563141	0.603595525	0.5	0.625344353	RGPD2	2	88124876	cg21821659	0.56925111	0.547076923	0.558126722	KIAA1310	2	97304298
cg04272694	0.603570248	0.532923077	0.619834711	DAC12	6	168718991	cg09084477	0.56924772	0.501538462	0.588980716	K8TBD7	13	41768978
cg07126216	0.603486332	0.548307692	0.61046832		12	108426797	cg23885108	0.56923755	0.499692308	0.616528926		17	37755750
cg05505680	0.603304937	0.640307692	0.500826446	MIR596	8	1765217	cg03958058	0.56922568	0.537538462	0.554269972	PPP2R2D	10	133761284
cg08158233	0.602949777	0.403076923	0.699173554		8	37351056	cg15627136	0.56921466	0.524923077	0.581818182		13	112986635
cg14585167	0.602921805	0.515692308	0.603305785		2	709329	cg10639425	0.56921212	0.555076923	0.55261708		5	1364079
cg04035728	0.602871795	0.618153846	0.530576512	CHRNA2	8	27323467	cg00009436	0.56921043	0.476923077	0.61322314		4	186402631
cg17441377	0.602638685	0.602769231	0.570738898		17	3906940	cg03586456	0.56920449	0.548923077	0.550413223	PTH2	19	49926070
cg10099757	0.602609027	0.578461538	0.603856749	HRNBP3	17	77207289	cg18296405	0.56919093	0.576307692	0.504683196	SENP7	3	10123232
cg09776757	0.60200551	0.579076923	0.577961433	RAI1	17	17653333	cg212920932	0.56917398	0.493230769	0.612627176	PCDH2A	5	140172958
cg08982961	0.601959737	0.617846154	0.589429752		2	227474635	cg05610131	0.56916131	0.492615385	0.590633609		13	112870576
cg10452666	0.601853783	0.550153846	0.577410488		14	90848892	cg24598948	0.56915194	0.554461538	0.572451791	LOC441666	10	42863539
cg18789932	0.601687646	0.657846154	0.528374656		12	11535700	cg25711997	0.56915109	0.538769231	0.499173554	HPRIP	10	106075710
cg06991380	0.601565586	0.437538462	0.742699725		4	8369129	cg21492832	0.56914431	0.5	0.580165289	TNFRM121	14	105991308
cg1764586	0.601520661	0.593076923	0.571349862		8	94871988	cg01702431	0.56913075	0.571076923	0.515702479	KSR2	12	118406138
cg11482879	0.600992562	0.578461538	0.534435262		17	4480083	cg23821249	0.56912227	0.556615385	0.56584022		22	16404670
cg25746778	0.600508582	0.485846154	0.520661157	EBF3	10	131758331	cg18736950	0.56911719	0.491692308	0.585674931		15	95870440
cg21489999	0.600498411	0.562769231	0.584573003	GRAMD2	15	72476373	cg18133145	0.56911549	0.538769231	0.563085399	PLEKHM2	1	16060689
cg26402169	0.600463658	0.570153846	0.579063361	MAD1L1	7	1883139	cg04227077	0.5691121	0.538769231	0.544903581		16	32252787

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg03163982	0.60940178	0.634461538	0.534435262		13	22058128	cg03103192	0.56910617	0.546769231	0.463360882	SPATA18	4	52917271
cg02040548	0.600197499	0.600923077	0.570247934	IPO9	1	2201798459	cg068883544	0.56008752	0.540307692	0.547655402	NRB	14	77732283
cg04433201	0.599940665	0.652307692	0.445179053	FAM101B	17	296265	cg06808498	0.53908498	0.539692308	0.549311295	SIX6	14	60974525
cg13874287	0.5997847	0.609230769	0.52231405		1	2820170	cg02272859	0.56904344	0.617338462	0.510192837	SLC34A2	4	25056514
cg14022794	0.599539733	0.462153846	0.642424242	CACNA1B	9	141017457	cg08053355	0.5690231	0.54	0.543250689	C2orf63	2	55458963
cg01442142	0.599316804	0.548923077	0.601101928	KY	3	134370862	cg22683388	0.56901293	0.476307692	0.525068871	AGRL5	2	27274757
cg16828293	0.599223564	0.561846154	0.532396694		13	107027718	cg07676517	0.56900021	0.555692308	0.543250689		15	96897414
cg13115661	0.599125238	0.617539462	0.550964187	ASTN1	1	177134454	cg24830524	0.56899174	0.524923077	0.563636364	COL6A3	2	238249152
cg20655119	0.599094723	0.570461538	0.586225895	RBM47	4	40425759	cg19603966	0.56898072	0.561538462	0.56584022	DNAJC1	10	22292833
cg26877594	0.598957406	0.445539462	0.656198347		8	297359	cg19189335	0.56897309	0.517230769	0.584022039	DLL1	6	17059369
cg17030488	0.598919263	0.538461538	0.596694215		12	115372748	cg13377264	0.56897224	0.594769231	0.519359229		6	30419135
cg02379819	0.598504853	0.656923077	0.491460055	SLC32A1	20	37355188	cg25963980	0.5689697	0.586769231	0.480991736	SLC1A2	11	35442225
cg10381071	0.598315851	0.601846154	0.574104683	TLE3	15	70391035	cg01023787	0.5689341	0.524	0.60508642	SLC6A18	5	1243714
cg23000734	0.598735325	0.508307692	0.615426957	TBP2	10	126850823	cg02393609	0.56891884	0.523384615	0.577410468	ALKBH4	7	102105195
cg16860694	0.59862259	0.464307692	0.54600551	FXYD5	19	35643689	cg09254318	0.56890273	0.556615385	0.575615835	BCAN	1	156625372
cg10323116	0.598525568	0.60846154	0.585674931		16	86795398	cg05654780	0.56890019	0.556615385	0.575615835	RASA3	13	114872606
cg08304537	0.598485272	0.577538462	0.591735537	MUC2	11	1095283	cg27086157	0.56889087	0.564615385	0.442424242	LOC728989	1	146515238
cg06120313	0.598386099	0.421538462	0.687052342	OSPL5	11	3174092	cg26810966	0.56887646	0.515076923	0.57630854	LOC728989	9	124924045
cg06012903	0.598377622	0.703692308	0.401101928	PTPRN2	7	157382161	cg14114910	0.56887222	0.432	0.638567493	MORF5	8	1400221
cg04232586	0.598269125	0.562461538	0.585123967	C8orf38	8	96037599	cg20859841	0.56887052	0.628307692	0.404407713		12	114234123
cg22726211	0.598078406	0.594769231	0.581267218		17	46662371	cg24771877	0.56885527	0.561846154	0.539944904	LOHNF2	2	100939477
cg27468857	0.597751218	0.616307692	0.585123967	ARHGAP10	8	1783358	cg21926883	0.5688256	0.532923077	0.546556474	ANKH	5	14870749
cg00327796	0.597745285	0.556	0.554820937		3	23032191	cg02107279	0.56882475	0.541230769	0.575206612		1	221069136
cg24786174	0.597710532	0.587384615	0.540495868	ZNF516	18	74118243	cg26047066	0.56880356	0.563384615	0.539944904	TRIM2	4	154220167
cg22158158	0.597687646	0.502769231	0.644628099	TNXB	6	31978789	cg13467964	0.56879508	0.463076923	0.516253444		11	124932906
cg27554551	0.597224836	0.540307692	0.605509642	VASN	16	4421486	cg01086404	0.56879254	0.532615385	0.561432507	SLC37A2	11	1594312
cg26888225	0.597199407	0.576923077	0.463564738	ZNRFL	16	73032603	cg19048476	0.56879169	0.540923077	0.544903581	HCCA2	11	1594312
cg15310617	0.59702225	0.570153846	0.577410468	ZNF776	19	58257751	cg23125542	0.56876457	0.572	0.519008264	ATP6D1	11	67183148
cg08172637	0.596993431	0.555384615	0.583471074	GNF7	19	2556024	cg15134801	0.56873999	0.500307692	0.575757576	ELOVL7	5	60138838
cg00197058	0.596934096	0.602461538	0.580165289	TBX4	17	59554672	cg25164624	0.56872897	0.547076923	0.597796143	GFR4	20	3641641
cg05989533	0.596567069	0.626461538	0.526721763		6	28710579	cg25353070	0.56872219	0.548	0.548209366	ZNF330	17	33289410
cg12870217	0.596528926	0.484307692	0.652341598		14	26862142	cg07291445	0.56869421	0.540307692	0.562534435		7	39015928
cg14714391	0.596493325	0.538153846	0.580716253	F2R	5	76010444	cg14767165	0.56869167	0.484307692	0.58625895	ACAT2	6	160183909
cg26631034	0.596409409	0.547076923	0.555371901	SLIT2	4	121590210	cg01288811	0.56868235	0.508	0.607713499	IFAD1	7	112062682
cg07790920	0.596248338	0.579692308	0.577961433	SLIT2	4	121590210	cg02285791	0.56868815	0.540615385	0.579063361		16	27550747
cg15612328	0.596133079	0.571384615	0.426997245		13	114051373	cg14487271	0.56867472	0.588615385	0.558677686	WWP1	8	87354422
cg09471658	0.596034753	0.548307692	0.587878788	TNFRSF8	1	12178858	cg06074774	0.56865553	0.555692308	0.549862259	SKPR	11	126138802
cg08886020	0.596023734	0.577846154	0.590082645		16	1080388	cg18937225	0.568635014	0.515692308	0.579614325	TTC39A	1	51793660
cg00509431	0.595944056	0.546461538	0.580165289	PHACTR2	6	143999465	cg27591511	0.56863318	0.530769231	0.569146006	CADPS	3	62571762
cg14836248	0.595932189	0.600923077	0.570798898	CANMT1	1	7209987	cg04994773	0.56863234	0.547692308	0.568959041	KIAA1324	1	109656777
cg05296266	0.595843187	0.579692308	0.550964187		18	15138176	cg200000805	0.5686264	0.547384615	0.551515152		1	97046756
cg08791131	0.59570926	0.532	0.588980716	NDHG4	16	58497801	cg23999405	0.56862386	0.554769231	0.452892562	MGMT	10	131265087
cg26362197	0.595571943	0.445230769	0.651239669	GRIN2D	19	49948005	cg14210765	0.56861538	0.461230769	0.633608815		18	73628830

ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg20227860	0.595425302	0.594769231	0.554820937	RFC1	4	39368720	cg05576795	0.56860606	0.491384615	0.609917355		5	173617174
cg08794954	0.595569358	0.572	0.572451791	DDZ4	11	79124571	cg27553372	0.56858741	0.564	0.571349862		12	120731091
cg10318039	0.595337995	0.516615385	0.583471074	SEMA3D	7	84671308	cg22514736	0.5657046	0.562461538	0.58292011		1	156426017
cg16302655	0.595504937	0.593538462	0.567493113	TTC12	11	11348579	cg01195861	0.56855435	0.570769331	0.547858402	POP5	12	121019729
cg27278519	0.595293918	0.586461538	0.541597796		12	81104111	cg04807108	0.56855266	0.563384615	0.541597796		15	78697933
cg24194674	0.595287137	0.532923077	0.629752065	ADAM12	10	127779842	cg1400650	0.56855012	0.539384615	0.547858402	MIR548H4	15	69328155
cg16160218	0.595193897	0.562461538	0.590082645	CNNM2	10	10480354	cg26707398	0.56854673	0.54	0.534986236	MYH14	19	50713914
cg22482194	0.594946366	0.546769231	0.575757576	MAP3K14	3	24538388	cg18023066	0.56854079	0.531384615	0.571900826	PSIPTE22	22	17090586
cg21658968	0.594845518	0.547692308	0.571349862		11	92754025	cg00697668	0.56853147	0.500923077	0.566942149	JAG2	14	105617575
cg04741728	0.594728544	0.671384615	0.483746556		12	133003907	cg25590734	0.56852384	0.555384615	0.525068871	CCDC124	19	18044416
cg14956836	0.594639542	0.484615385	0.630303033	KCNMA1	10	79201862	cg06088360	0.56850519	0.539384615	0.597796143		6	170538929
cg24307499	0.594574274	0.610153846	0.579063361	NLRP2	19	55494075	cg204927230	0.56849841	0.455692308	0.580716253		14	86577671
cg25428278	0.59428523	0.508923077	0.620936639	PPP2R2C	4	6474782	cg26684319	0.56847552	0.563692308	0.540495868	ERICH1	8	672406
cg07018853	0.594259801	0.548307692	0.552066116	KIAA1310	2	97304450	cg06525145	0.56845772	0.413538462	0.674931129		1	144594085
cg18318878	0.594203009	0.556923077	0.592837466		1	214159037	cg14822696	0.56844162	0.610153846	0.568044077	ASPSR1	17	79935258
cg1714861	0.594060606	0.556	0.570247934	PRKAG2	7	151423659	cg20169823	0.56841195	0.547692308	0.562534435		12	116354837
cg13720737	0.594058063	0.578769231	0.570247934		13	113611494	cg03711791	0.56841111	0.547076923	0.578512397	KCNK1	20	49626347
cg16504398	0.593977538	0.576923077	0.575757576	PIP4K2A	10	22902615	cg08203127	0.56840432	0.477538462	0.575553719	ARHGAP16	16	331727
cg03057712	0.593975669	0.515692308	0.550964187		19	2489637	cg06555490	0.568377211	0.522153846	0.547107438	CSNK1A1	5	148931367
cg00681003	0.593861411	0.470769231	0.630303033	CST9	20	22585647	cg05930207	0.56837127	0.409538462	0.654545455	GID3	17	38518520
cg04452040	0.5938337677	0.555692308	0.568044077		10	102501358	cg18644252	0.56833312	0.454153846	0.62369146	LOC144571	12	9217859
cg24152353	0.593780886	0.571384615	0.586225895	KCNAB2	1	6149107	cg16541222	0.56831871	0.556307692	0.538292011	EPB41A4	5	111541635
cg01067405	0.593718161	0.634461538	0.506887052	LILRA4	19	54850417	cg05946687	0.568306	0.532615385	0.559779614	NKX2-1	14	36989287
cg13668060	0.593706294	0.557230769	0.557575758		9	102131971	cg00689492	0.56830345	0.508615385	0.581818182	MAEA	4	1303491
cg00460690	0.593681712	0.54	0.597245179	PLEKHA7	11	16909174	cg20033239	0.56826701	0.579692308	0.523966942	LOC84856	10	42972782
cg08823894	0.593626616	0.447692308	0.640220386	ASCL4	12	108167969	cg24355809	0.56825599	0.500307692	0.619283747	NXN	17	746574
cg07148341	0.593619835	0.477846154	0.61322314		6	43744275	cg01601518	0.56822293	0.556923077	0.49882259	ZFYVE28	4	2404284
cg04759767	0.593550328	0.612	0.556473829		13	25320824	cg02984188	0.56819326	0.525230769	0.580716253	GRH13	1	24646392
cg02735486	0.593549481	0.594153846	0.544903581	ANK2	4	113970935	cg22001483	0.56818818	0.594153846	0.52112121		6	168109493
cg04221521	0.59353168	0.579076923	0.549862259		5	72597715	cg14238325	0.56816868	0.594461538	0.525068871	CHP3	6	43276781
cg25985778	0.593530833	0.564307692	0.552066116	RPL38	17	72199373	cg14676702	0.56816783	0.535692308	0.534986236	C2orf77	2	170550423
cg21238227	0.593535289	0.585538462	0.552066116	MTT1L	2	2111623	cg21453656	0.56816783	0.539384615	0.569146006	USP13	3	179502686
cg04790977	0.593430962	0.523692308	0.616528976	C1orf96	2	220717120	cg23230834	0.56816275	0.478461538	0.592286501	MACC1	7	20198439
cg20798574	0.593325683	0.608307692	0.543352617	LMO3	12	1678267	cg004119909	0.56815512	0.555692308	0.527272727	DIG5	10	79646104
cg13372309	0.593193473	0.617538462	0.5184573	THRA	17	38244572	cg24504927	0.56814274	0.484923077	0.591184573	OC90	8	133071588
cg03152602	0.593088366	0.570153846	0.574655647		1	38244572	cg24504927	0.56812121	0.508538462	0.575757576		12	85667353
cg10118705	0.593084976	0.610769231	0.495316804	HDAC4	2	239974138	cg13683280	0.56810935	0.650461538	0.446831956	OIFM1	9	137980469
cg00223136	0.593057851	0.523076923	0.535371901		5	3609364	cg00515161	0.56810595	0.581846154	0.510743802	BHP1	17	59940900
cg07078095	0.592951897	0.610153846	0.548760331	ZNF398	16	2054141	cg00575066	0.56810595	0.581846154	0.510743802		6	10390039
cg06728493	0.592410256	0.555692308	0.55261708	SOX2OT	3	181417613	cg21617694	0.56810426	0.575846154	0.560881543		6	28715138
cg16481556	0.592273787	0.585846154	0.553168044		8	2196764	cg27054655	0.56810087	0.515384615	0.563636364	ADAMT52	5	178722969
cg19480425	0.592194109	0.548	0.574655647		22	22339538	cg09019134	0.56805949	0.604	0.479338843		8	19616280
cg09654046	0.592104259	0.483692308	0.652892562	FRM4	7	99517255	cg26262786	0.56803221	0.576923077	0.5158126722	SPG7	16	89590583

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg22523072	0.592094935	0.546461538	0.585123967		1	248300255	cg10372920	0.5680178	0.463076923	0.609917355		17	5800501
cg03987842	0.591978809	0.632615385	0.492561983	SLC7A6	16	68321858	cg14013597	0.56801102	0.547384615	0.541040832		2	131046122
cg24596023	0.591972028	0.578153846	0.567493113	RHOA	3	4949349	cg267223707	0.5680037	0.587692308	0.523415978		7	142045216
cg19003626	0.59190676	0.438461538	0.583471074	KIF26A	14	104640199	cg25664021	0.56800085	0.517846154	0.612872176	LRP1	12	57595357
cg15897613	0.591848273	0.570461538	0.553719008	ILUZF1	1	23418405	cg18642499	0.56799746	0.507384615	0.566942149	FAM150B	2	289550
cg03884405	0.591808434	0.539076923	0.562534435	PHF20	20	34359544	cg13468457	0.56799661	0.539384615	0.561432507		9	110501986
cg01256539	0.591789786	0.593230769	0.530578512	PRR16	5	119802022	cg26826330	0.56797372	0.547692308	0.527272727		7	155592408
cg18296334	0.591788091	0.523384615	0.551515152	TNXL4A	18	77748704	cg13871565	0.56792541	0.587692308	0.511294766		1	244973585
cg08096062	0.591760966	0.555692308	0.583471074	CCDC85A	2	56410509	cg072227033	0.56792371	0.567692308	0.539944904	ARL4A	7	12725802
cg0279549	0.591643231	0.507384615	0.604407713	CGORF89	6	36887307	cg016488609	0.56792371	0.516923077	0.577961433		4	2049057
cg07422940	0.591641449	0.54	0.601101928		16	54546809	cg04191954	0.567911	0.564307692	0.501926375	CACNG2	22	37099064
cg0677570	0.591579572	0.509538462	0.643526171		11	15962841	cg14728235	0.56786662	0.514461538	0.573302755	GNASAS	20	57415177
cg03015758	0.59155499	0.562153846	0.581267218	CCDC64B	16	3083902	cg16058990	0.5678576	0.549230769	0.532782369		6	30359364
cg09024430	0.591531257	0.608		MED18	1	28655366	cg07774884	0.56784573	0.525538462	0.592833466	STC1	8	23712364
cg18291703	0.59144395	0.623692308	0.572451791		7	2518684	cg05325887	0.56784573	0.525538462	0.592833466	C16orf11	16	611985
cg06019853	0.591393092	0.626769231	0.548209366	ESRP1	8	95651895	cg03099683	0.56784064	0.517538462	0.592786501		4	132661931
cg23180593	0.591353253	0.586461538	0.563636364	CARS2	13	111301487	cg18215878	0.5678386	0.516	0.563085399	PLEKHA5	12	19459468
cg19999035	0.591328671	0.531384615	0.586255895	SHROOM3	4	77355817	cg16661464	0.56781945	0.54	0.577410468		18	61037759
cg24760414	0.591301547	0.562153846	0.550964187	OSBP15	11	3167563	cg17034181	0.56781182	0.539384615	0.571900826	SLC30A7	1	101362442
cg16470793	0.591294766	0.587384615	0.507988981		16	1330147	cg16675034	0.56777961	0.484	0.624793388	STK10	5	171616574
cg05223487	0.591215936	0.580307692	0.579063361		16	1080540	cg01298579	0.56777114	0.523076923	0.587678788	NO10	2	10630716
cg01251646	0.591172706	0.570153846	0.537741047		5	172719820	cg26509801	0.5677491	0.499692308	0.582369146	KEAP1	19	10613855
cg14644378	0.591172706	0.469846154	0.643526171	DNAJB5	9	34988588	cg16907514	0.56772197	0.515692308	0.581267218	KTNP	14	56046098
cg27432604	0.591092181	0.556615385	0.570798998		11	27239162	cg03395651	0.56771774	0.514153846	0.552066116	BANP	16	88107091
cg13456086	0.59105997	0.406769231	0.721763085	GOLSN	8	110704757	cg11393479	0.56771265	0.50120769	0.580165289	RGL3	19	11529564
cg06842946	0.590968769	0.54	0.582369146		11	49070005	cg27351780	0.5677118	0.563384615	0.564187328	FIX1	11	35639995
cg19443257	0.590967578	0.538769231	0.582369146	SPSB4	3	140770549	cg26174215	0.56768976	0.535692308	0.551515152	IFT140	16	1583899
cg25921813	0.590834499	0.579076923	0.591184573	C3orf71	3	194797138	cg07341220	0.56768468	0.508	0.554269972		8	
cg20402074	0.590612418	0.547384615	0.571349862	SLAE	11	124543718	cg07633618	0.56767535	0.509846154	0.487603306		3	185000208
cg01200150	0.590542064	0.516307692	0.585123967		18	76322683	cg18009000	0.56765925	0.537846154	0.58677686	H2AFY2	10	71811927
cg00861497	0.590531045	0.501538462	0.603305785		7	131514718	cg03304378	0.5676406	0.564615385	0.507438017		2	5847191
cg26914865	0.590517483	0.438153846	0.613774105		19	13735016	cg22111318	0.56761263	0.540307692	0.547107438	SDHAF2	11	61196590
cg02633380	0.590493749	0.564	0.56584022	ODF2L	1	86859488	cg10700718	0.5676067	0.595692308	0.54214876	MYOM2	8	19927719
cg04117433	0.590472558	0.437846154	0.660606061		10	783591	cg21457909	0.56755499	0.619076923	0.492011019	AGTR1	3	148416019
cg00711088	0.590404289	0.532	0.590082645		4	1401409	cg10812016	0.56754397	0.593538462	0.520681157		7	155385272
cg24741158	0.590406442	0.439384615	0.632506887		1	2844168	cg03196627	0.56752278	0.523384615	0.54600551	RASGRF1	15	79298606
cg04837959	0.590315745	0.524307692	0.601101928		1	29742186	cg20599295	0.56751854	0.540307692	0.558677686	RILP1	12	123957410
cg23786592	0.59028523	0.522461538	0.576859504	SFXN4	10	120925342	cg27636310	0.5675143	0.525846154	0.60716253		1	46632696
cg02586634	0.590169951	0.506769231	0.581267218	PPP2R2C	4	8449000	cg16126665	0.56751346	0.549538462	0.510743802	BC11B	14	99711752
cg8467168	0.590077559	0.468615385	0.57630854	HOSFC1	3	13114803	cg15444387	0.56750668	0.587384615	0.543801633	GLTSCR2	19	48248531
cg25143235	0.590016529	0.594307692	0.529476584	CHD5	1	6187886	cg22702618	0.56750074	0.500615385	0.577410468	CRF1	19	18705112
cg17179053	0.589931765	0.617230769	0.517906336	TYW3	1	75198211	cg20523839	0.56749735	0.524923077	0.558126722	CU68P2	10	11069315
cg08111036	0.589927527	0.579076923	0.573002755		19	2950164	cg233331981	0.56748972	0.523076923	0.543801633		10	7574780
cg01591431	0.58979021	0.561846154	0.558126722	C11orf63	11	122753443	cg23114616	0.56747192	0.461846154	0.624242424	TNFRSF25	1	6526398

ILMINID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg1176689	0.589773257	0.548	0.581267218	KCNF1	9	138594307	cg15225267	0.56746514	0.542461538	0.579063361		6	31276187
cg185332727	0.589719856	0.401230769	0.591184573	C17orf51	17	21454896	cg134032233	0.56746175	0.523384615	0.565289256	KIAA0913	10	75547933
cg11436025	0.58970036	0.446153846	0.585674931	DSCAM1	11	117376238	cg10142500	0.56745412	0.516307692	0.579063361		2	29174548
cg06040980	0.589663912	0.501230769	0.611019284	FLJ9582	22	21356317	cg04391972	0.56745158	0.537307692	0.560881543		4	182569038
cg20852302	0.589604577	0.537307692	0.540495868	UPF2	20	44784898	cg14094375	0.567427	0.626461538	0.466666667		9	133306791
cg09812413	0.589532528	0.601538462	0.550964187		10	12085555	cg15234263	0.56741852	0.509240769	0.553571901	8MS1P5	10	467627703
cg01661130	0.58952829	0.587384615	0.603305785		16	51110257	cg16287766	0.56741174	0.546769231	0.560330579	PACSIN1	16	34495124
cg11557030	0.589478279	0.493230769	0.558126722	KCNK10	14	88793311	cg07788140	0.56740665	0.516307692	0.590082645		12	113921202
cg05054998	0.58919093	0.554769231	0.588429752	C21orf70	21	46378625	cg06768599	0.56740242	0.626461538	0.503030303	LTBR	14	24785488
cg06548416	0.589186692	0.552066116	0.552066116	MYOM3	1	24438703	cg22582981	0.56740157	0.531692308	0.578512397	H53ST3A1	17	13504472
cg1842308	0.589151939	0.586153846	0.563636364	RNF5P1	6	32145870	cg00326464	0.56739479	0.423384615	0.558677686	TFAP2A	6	10404084
cg17826980	0.589121424	0.524	0.568044077	PAPPA2	1	176771674	cg06840109	0.56738462	0.564	0.523966942		7	63498367
cg1480234	0.589101081	0.602153846	0.545454545		5	87986067	cg10781852	0.56736766	0.514769231	0.570247934	FLJ10661	8	8065532
cg18770678	0.589062937	0.587076923	0.562534435	QRICH1	3	49069678	cg13244559	0.5673541	0.463692308	0.601652893	ADRA2C	4	3769810
cg12840312	0.588992583	0.548615385	0.59338843	FAM508	6	3849381	cg20816934	0.56734308	0.555384615	0.565289256		7	75799644
cg15445000	0.588970545	0.563076923	0.596143251	MED1	17	37608096	cg18512336	0.56732952	0.555692308	0.546000551	PRDM14	8	70981068
cg04187088	0.58889252	0.570153846	0.568044077	VGLL2	6	117586225	cg14042780	0.56732698	0.514461538	0.573553719		7	53106426
cg22741264	0.5888635728	0.610769231	0.543801653		12	115105519	cg14195925	0.56731596	0.520923077	0.587878788	HRNBP3	17	77224403
cg19588770	0.588578936	0.633846154	0.45785124	CENPO	2	25016434	cg22504285	0.5673007	0.546153846	0.544903581	ATP2A2	12	110718598
cg115050510	0.588511125	0.545846154	0.503030303	TEX101	19	43912492	cg11123440	0.56729646	0.578153846	0.497520661		8	11619852
cg09508613	0.588498411	0.602461538	0.561432507	SPRE2	16	89926162	cg13395698	0.56727357	0.54	0.573002755		11	74146974
cg24631784	0.588490782	0.579692308	0.4815427		8	122368294	cg13989864	0.56725747	0.494461538	0.63036303		12	126374295
cg13794009	0.588372113	0.539076923	0.602203857	DDAH2	6	31697530	cg15701622	0.56725662	0.510769231	0.564187328	MCRS1	12	49962202
cg13174084	0.588358551	0.596307692	0.557575758	LOC388965	2	84517877	cg02610227	0.56725408	0.580307692	0.553169044	YIF1B	19	38801653
cg19658046	0.58833736	0.508307692	0.621487603	FAM193B	5	17681988	cg26386839	0.56722441	0.538769231	0.548208366	THBS4	5	79378955
cg26069539	0.588214452	0.540307692	0.63415978		17	49021653	cg26054842	0.56721848	0.439076923	0.662809917		7	155915712
cg20580578	0.588198347	0.553846154	0.559779614	HIP1	7	75189143	cg08467108	0.56720576	0.461846154	0.55261708	EGFR	19	41304898
cg15013768	0.588114431	0.585230769	0.547658402	COMMD1	2	62132624	cg21808635	0.56720153	0.524307692	0.558677686	EGLN2	7	55225133
cg08019449	0.588074592	0.428923077	0.613774105	H1A-DQB1	6	32632800	cg07546293	0.56719559	0.556307692	0.561983471	GNGL3	16	851255
cg13888983	0.588044925	0.554153846	0.575206612	FREQ	9	132981932	cg15833099	0.56719135	0.553538462	0.546000551	GPR39	2	133407999
cg18686498	0.588015257	0.484615385	0.591184573	NALCN	13	102069757	cg23370710	0.56719051	0.546769231	0.557024793	TMS3	7	47354947
cg1496419	0.587990676	0.501538462	0.595041322	LYPD5	19	44306685	cg01332259	0.5671761	0.555692308	0.550964187	CREB5	7	28450797
cg02679779	0.587973723	0.633230769	0.537190083	FAM114A2	5	153417925	cg27093000	0.5671761	0.522769231	0.584022039	TRAIP1	12	112563698
cg14097619	0.587965247	0.524	0.572451791		3	183945054	cg06877486	0.5671744	0.477538462	0.595041322	GTFF2	13	45694166
cg03184452	0.587945751	0.561846154	0.581818182	DLC1	8	13105155	cg23344136	0.56715067	0.508307692	0.573002755	ODZ3	4	183686196
cg15952586	0.587940665	0.572	0.586225895		7	18459971	cg19876775	0.56714982	0.538769231	0.590633609	LRR32	11	76381937
cg27428708	0.587932189	0.571076923	0.557024793		16	54851977	cg15557309	0.56713711	0.578461538	0.537741047	MIR1182	1	231155948
cg06107159	0.587886417	0.431076923	0.690991736	LIMS2	2	128432243	cg26888284	0.56713202	0.541230769	0.55927865		15	20487955
cg16212881	0.587830473	0.516307692	0.596694215	DHX16	6	30640519	cg18100830	0.56712609	0.486153846	0.586225895		7	123903317
cg1715676	0.587824625	0.524615385	0.606060606	VWA38	2	98928898	cg14511521	0.5671193	0.54	0.56584022		2	71113568
cg16592586	0.587816063	0.586769231	0.530578512	GIR3A3	4	175751001	cg26988138	0.56711676	0.524	0.555922865	GNF7	19	2639474
cg22812331	0.5877847	0.594461538	0.566942149		6	33869169	cg27070162	0.56711676	0.548923077	0.575757576	RASGEF1C	5	179597526
cg18759081	0.587748252	0.517230769	0.596143251	COL9A3	20	61467939	cg17730530	0.56711422	0.532307692	0.560881543	CASC3	17	38296876
cg26023303	0.587732994	0.618461538	0.549862259		14	103541561	cg20955032	0.56710913	0.460307692	0.631404959	MIR549F2	2	213297541

ILMNID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg16357719	0.587640602	0.587076923	0.549862259	RNF51	16	2317538	cg16705744	0.56709727	0.507076923	0.584573003		3	27756039
cg04516952	0.587589744	0.602769231	0.560330579	KLK6	19	51472042	cg25285743	0.56707777	0.555076923	0.551515152	LMO3	12	16701533
cg01130611	0.587556686	0.556615385	0.584573003		3	46868978	cg05477834	0.567077438	0.491692308	0.617630854	PCDH17	5	140536175
cg12626372	0.587500742	0.4701155846	0.510192837	TMT1C3	12	38536556	cg02227919	0.56707099	0.523692308	0.564738292	LOC404925	2	171569132
cg16308530	0.587367663	0.571076923	0.575757576	AXIN2	17	63558518	cg22969763	0.56705573	0.539384615	0.563636364	KLR1	12	10607538
cg03147210	0.587254079	0.516923077	0.557024793	THEM4	1	151882284	cg05290300	0.56705149	0.547076923	0.543250689		16	3066763
cg00687447	0.58722865	0.556	0.572451791		16	1158657	cg19139727	0.56703963	0.477846154	0.504132231		10	35230860
cg19172278	0.58721085	0.579384615	0.562534435		7	152623304	cg09336982	0.56702776	0.491692308	0.553719008	CALHM2	10	105212481
cg0054641	0.587209154	0.563076923	0.553168044	TRIM31	6	30071503	cg26860182	0.56702183	0.532307692	0.575757576	FBRSL1	12	133066054
cg2029682	0.587159144	0.530461538	0.561983471	SLC30A3	2	27486061	cg00393673	0.56701674	0.516307692	0.563085399	LOC400794	1	165494483
cg27536187	0.587113371	0.584923077	0.54600551	SKI	1	2159365	cg05678498	0.56700911	0.532615385	0.537190083		12	94031876
cg01847344	0.587082009	0.548	0.558677686	C21orf29	21	43929576	cg02166394	0.56699725	0.563076923	0.505234416		5	2741746
cg16139559	0.58705997	0.470153846	0.63415978	RXFP3	14	37069791	cg19109570	0.5669786	0.478153846	0.604958678	SLC6A19	5	1201479
cg13249651	0.587054037	0.563384615	0.57630854		5	33926307	cg11123744	0.56695232	0.547692308	0.533333333		18	77376639
cg17875102	0.587011655	0.547384615	0.596143251	ALX4	6	33926307	cg08605326	0.56694978	0.469538462	0.590633609	CA10	17	50237401
cg13058901	0.586993007	0.533230769	0.583471074		11	44324759	cg26036375	0.56694978	0.586769231	0.497520661		2	122660039
cg20383948	0.586926891	0.593230769	0.568044077	COL18A1	21	4688137	cg25786388	0.56694469	0.595076923	0.521763085		13	100578831
cg14744537	0.586876881	0.600615385	0.553719008	RAB27B	18	52495740	cg09508668	0.56694045	0.493230769	0.554269972		13	28531046
cg27160879	0.586846366	0.577230769	0.553719008	LOC440356	16	29874506	cg15193614	0.56693537	0.524307692	0.583471074		5	2239477
cg06035456	0.586804832	0.500307692	0.607713499	COL13A1	10	71560212	cg15787146	0.56690401	0.532923077	0.554920937		17	70107185
cg13753428	0.586747192	0.539076923	0.534435262	RALGDS	9	136001623	cg09510604	0.56689553	0.533230769	0.571900826	NTRK1	1	156851440
cg19980648	0.586690401	0.631076923	0.522865014		2	219219072	cg17034508	0.566889468	0.533538462	0.550864187	TLX1B	10	102877719
cg17074338	0.5866726	0.624923077	0.509641873	GALNT4	11	11642133	cg10236261	0.56689044	0.561220769	0.553719008	#T81	12	110573183
cg25625308	0.586664124	0.484	0.611019284	CDC20B	5	54469057	cg00774005	0.56688922	0.468923077	0.644077135	DLX6AS	7	96433726
cg21096915	0.586655647	0.509538462	0.575206612	LOC84856	10	42971209	cg02098378	0.56688197	0.546769231	0.555571901		2	124584426
cg01017839	0.58656435	0.579384615	0.501928375		1	181448627	cg22753661	0.56687264	0.531692308	0.543250689	ADAMTS7	15	79092743
cg12265810	0.586559017	0.532923077	0.609917355	FOSB	19	45975694	cg09876447	0.56686925	0.617538462	0.528374656	MUC21	6	30950282
cg04976075	0.586490358	0.541230769	0.584573003	ADARB2	10	1230850	cg14157549	0.56686247	0.476615385	0.613774105		6	168393963
cg13753637	0.586476796	0.547692308	0.551515152	CSDA	12	10876057	cg14432665	0.56685145	0.501230769	0.575619835		9	88377422
cg02190253	0.586394575	0.556923077	0.564187328	C10orf119	10	121633065	cg13717425	0.56684891	0.413230769	0.654545455	AKR1E2	10	4868892
cg08317263	0.58637847	0.572	0.553168044	CCDC69	5	150603169	cg01109065	0.56683196	0.594153846	0.540495868	WNT5A	3	55508379
cg09462240	0.586370841	0.430769231	0.636363636		4	1409497	cg08495033	0.56682517	0.547384615	0.58292011		8	143680998
cg08310581	0.586336936	0.477846154	0.638016529	ARHGTF4	2	131797438	cg13158571	0.56682178	0.589230769	0.505785124	CYP4X1	1	47489195
cg19028462	0.586335241	0.586461538	0.560330579	MCF2L	13	113677628	cg21393051	0.56681924	0.510153846	0.560881543		1	208585310
cg08689344	0.58615978	0.548307692	0.604407713	GRM4	6	34031597	cg18661413	0.566815	0.597230769	0.463918846	SCHIP1	3	159478785
cg09910392	0.586058063	0.420923077	0.6		16	34809677	cg13668207	0.56681076	0.500307692	0.563636364		12	65904452
cg15938047	0.585968214	0.539384615	0.512396694	IPH3	16	87637283	cg25649283	0.56679296	0.508307692	0.556673829	EHMT1	9	140714075
cg02272280	0.585964823	0.641846154	0.495316804		18	12076435	cg21830368	0.56678025	0.578153846	0.531129477	9-Sep	17	75373327
cg11065385	0.585963975	0.4446769231	0.55261708	B4GALNT3	12	568798	cg03188200	0.56677771	0.547384615	0.558677686	SNORD115-9	15	25436244
cg04493054	0.58594448	0.462461538	0.639118457		4	84972436	cg04575609	0.56677432	0.521846154	0.56584022	BANP	16	38061548
cg20556675	0.585905488	0.649538462	0.495316804	OTUB2	14	94493284	cg19478951	0.56677008	0.562153846	0.562534435		10	133879084
cg21604741	0.585904641	0.600923077	0.573002755	IFT172	2	27713092	cg14981312	0.56676923	0.469538462	0.609366391	C1orf159	1	1050737
cg25831111	0.585903793	0.525230769	0.61322314	COASY	17	40713740	cg22708339	0.5667616	0.525846154	0.570747934		1	73500111

ILMINID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg09511963	0.585900403	0.500307692	0.612672176	LOC494141	11	18230625	cg10411339	0.56676075	0.572923077	0.533333333	TATDN1	8	125550661
cg11174640	0.585860907	0.617230769	0.539944904		6	27259237	cg04128348	0.56675567	0.586153846	0.53553719	KCNK2	1	215257222
cg11197905	0.585872431	0.493846154	0.582369146	ZNF518B	4	10459435	cg11433866	0.56674889	0.579076923	0.54214876		12	115126908
cg13083277	0.585659674	0.609846154	0.521763085	COL29A1	3	130064935	cg21097030	0.56674719	0.556	0.573002755	ASPG	14	104570926
cg00760290	0.585630954	0.531384615	0.585674931		16	33374131	cg20033981	0.5667438	0.531384615	0.530027548	ELF1	13	41590153
cg15380508	0.585611358	0.541230769	0.58625895	ODZ2	5	166874870	cg04298253	0.56674126	0.532	0.532469146	PRDM7	16	30143815
cg05820241	0.585597796	0.596307692	0.554820937	PTPRN	2	220174629	cg23329272	0.56673533	0.5	0.595041322		5	180085924
cg06251537	0.585595538	0.656615385	0.523966942		22	49798750	cg07211768	0.56672176	0.508615385	0.568044077		12	115102290
cg23684088	0.585569824	0.460615385	0.520110193		6	28510928	cg18058532	0.56671074	0.562461538	0.557024793	PAX6	11	31832710
cg13630250	0.585508794	0.516615385	0.581818182	COX16	14	70827336	cg08714996	0.56669549	0.4923307692	0.592837466	PCDHGB1	5	140729928
cg17953201	0.585474041	0.493846154	0.564738292	KIF26B	1	245851802	cg14462307	0.56669464	0.485538462	0.54606551	ALS2	2	202644572
cg18761461	0.585382496	0.569846154	0.557575758	HEBP2	6	138775231	cg27664418	0.56667599	0.471692308	0.608815427	MIR517A	19	54214531
cg07790170	0.585264675	0.641230769	0.524517906		4	111537570	cg26879386	0.56665141	0.531384615	0.556473829	PCDHGA4	5	140781179
cg14414944	0.58523416	0.5	0.584022039		2	583253	cg27619291	0.56664571	0.531384615	0.556473829	TRPA1	8	72988380
cg16590635	0.585213816	0.609230769	0.564738292	IPH3	16	87641681	cg25433760	0.56664039	0.603384615	0.534433262		19	264871
cg19740491	0.585160415	0.524923077	0.579614325	BAT1	6	31510208	cg04938780	0.56661242	0.524615385	0.564738292		2	130960303
cg22975331	0.585093452	0.577230769	0.568595041		17	22017237	cg13831106	0.56660564	0.508923077	0.549311295	MYT1L	2	2119853
cg01710432	0.585085823	0.571384615	0.564187328	HCM4	15	73635811	cg03566814	0.56658021	0.492	0.575757576	SMOC2	6	169019307
cg27367871	0.585046832	0.601846154	0.544352617	CXCR7	2	237478030	cg23232339	0.56655986	0.555384615	0.571349862	PRKZ	1	2003605
cg07836478	0.584983259	0.524615385	0.569146006		18	15330632	cg06881310	0.56655732	0.516923077	0.538292011	NHS1	6	138893718
cg27207756	0.584982412	0.438153846	0.652892562		4	1607291	cg23041020	0.56654122	0.517538462	0.562534435	TOILIP	11	1298152
cg15165230	0.584976478	0.525846154	0.596694215		4	3630746	cg17034331	0.56653189	0.452923077	0.641322314		5	77146318
cg03066754	0.584960373	0.547384615	0.601101928	DNAH17	17	76506831	cg07098168	0.56653104	0.471076923	0.592786501	INA	10	105049522
cg24298548	0.584945963	0.468923077	0.631955923		8	33792277	cg01387102	0.56652765	0.492615385	0.603305785	MLPH	2	238395125
cg06848958	0.584834923	0.501230769	0.521763085	BALAP3	16	1384655	cg06381133	0.56652087	0.554461538	0.461707989	FRS3	6	41747367
cg26889677	0.58480017	0.532	0.567493113		16	88349724	cg14874390	0.56651324	0.429846154	0.543250689	SGCE	7	94286304
cg10477905	0.584787455	0.587692308	0.52231405		2	177039295	cg14579997	0.56649968	0.492307692	0.592286501	PNRC1	6	89792529
cg16389456	0.584672176	0.562769231	0.551515152		12	123636548	cg06093811	0.5664946	0.524307692	0.584573003	MTFSD	16	26588436
cg23691246	0.584638271	0.585846154	0.568044077	CALHM2	10	105207034	cg25238274	0.56648866	0.445846154	0.598898072	ADAM12	10	127914098
cg24176383	0.584616232	0.554461538	0.585123967	ORZ21	19	8941059	cg23825213	0.56648527	0.523692308	0.544352617	KCNK2	18	77623475
cg18127012	0.584615385	0.569846154	0.54214876		4	160514445	cg24627419	0.56648273	0.562461538	0.553168044	C1orf61	1	156387415
cg17870484	0.584591651	0.556307692	0.53553719		5	2515708	cg22082564	0.56647171	0.500307692	0.599449036	UMPS	3	124449312
cg00738849	0.584534859	0.524	0.59338843	TUBB4Q	4	190906046	cg23332027	0.56646662	0.531384615	0.586225895		3	13681764
cg02270332	0.584493325	0.484615385	0.608264463		6	106475062	cg10043155	0.56646662	0.493846154	0.55979614		6	25218855
cg06127263	0.584487391	0.546769231	0.571900826	DTHD1	4	36324706	cg24413339	0.56646578	0.639692308	0.506336088	SPRN	10	13523754
cg15327547	0.584478067	0.507692308	0.630303030	ZNHIT1	7	100862300	cg23873703	0.56644982	0.424	0.644628099	KCNAB1	3	155838625
cg24852442	0.584464658	0.554769231	0.584022039	MYO7A	11	76903134	cg17148395	0.56644459	0.523076923	0.562534435	IGDCC3	15	55654217
cg13234848	0.584461115	0.539384615	0.562534435	AUIS2	7	69063971	cg08261138	0.56643102	0.595692308	0.556473829	KCTD13	16	29937626
cg24537912	0.584320407	0.446461538	0.720661157		7	46276807	cg22720431	0.56640729	0.501538462	0.571349862		13	107344392
cg07666891	0.584250901	0.549230769	0.552066116	MYRIP	3	39850280	cg18045108	0.56639712	0.532	0.534435262	TREH	11	118530628
cg14150199	0.584181394	0.556615385	0.559779614	LOC1003026	2	53994733	cg19586676	0.56639712	0.488615385	0.628650138	H1A-A	6	29913343
cg06067394	0.584087307	0.576615385	0.547107438	IGSF9B	11	133789110	cg16092346	0.56638779	0.578153846	0.466666667	KIAA1530	4	1363885

ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg18720628	0.58406103	0.479076923	0.633608815	NEDD4L	18	55862058	cg21718380	0.56638101	0.517230769	0.608915427		4	3827930
cg08515845	0.584037296	0.554461538	0.564738292	CDK5	7	150755180	cg14896608	0.56638999	0.538153846	0.515702479	RPA1	17	1799628
cg09807215	0.583968637	0.547384615	0.577961433		8	144510884	cg09414673	0.56636576	0.516651385	0.588980716	CPV2W1	7	1022797
cg14794228	0.583940665	0.556118044	0.553168044	ZNF577	19	52391166	cg27572240	0.56634287	0.548	0.557024793	RANGAP1	22	41682267
cg19251564	0.583938122	0.492923077	0.646280992		6	166418922	cg15405572	0.5663327	0.540923077	0.578512397	NAV3	12	78224552
cg27577149	0.583910998	0.521846154	0.614876033	PCDHGA1	5	140712441	cg16612072	0.56630473	0.617846154	0.536639118	DCPIA	3	53380856
cg05280206	0.583857597	0.501946154	0.539008264	PCDH8B1	5	140596187	cg26317899	0.56630049	0.547384615	0.470523416	ACACA	17	35527041
cg07276679	0.583771138	0.447384615	0.617630854	HCCA2	11	1575607	cg10154804	0.56628777	0.586153846	0.558126722	DMXL2	15	53914952
cg19376449	0.58369146	0.570769231	0.581267218	PCDH46	5	140246626	cg043206342	0.56627421	0.524307692	0.563836364		10	134310836
cg16466917	0.583663488	0.564923077	0.575206612	LOC285696	5	17216259	cg04651623	0.5662598	0.516651385	0.587878788		6	164614956
cg15048078	0.583638059	0.546769231	0.574104683	KIF3	4	38665599	cg14898623	0.56623895	0.478461538	0.588429752	NCOR2	12	124941871
cg06060608	0.583539733	0.547692308	0.563085399	P2RX5	17	3600383	cg23158877	0.56620216	0.523384615	0.581818182	C11orf73	11	86012876
cg14707006	0.583503285	0.617846154	0.468870523		5	87982684	cg10248981	0.56619862	0.523384615	0.566942149	LHPP	10	126157948
cg00625161	0.583498199	0.539384615	0.586225895	SNORD115	5-15	25429408	cg07490279	0.56618182	0.45692308	0.527823691	ATAD38	1	1407066
cg18253764	0.583474465	0.569230769	0.524517906	MAGE12	15	23883040	cg26870745	0.56617758	0.548	0.538842975		19	3133503
cg12291627	0.58347277	0.549846154	0.502479339				cg13524588	0.56617504	0.575538462	0.516804408	AES	19	3059753
cg14062420	0.583442255	0.72	0.403856749		5	178960377	cg01526870	0.56617504	0.562769231	0.54600551	LOH12CH1	12	12510070
cg234445461	0.583326128	0.508923077	0.6	PCDHGA4	5	140864733	cg15346242	0.56616402	0.479076923	0.545454545		3	196325552
cg06726860	0.583240517	0.640923077	0.509641873	FGF8	10	103530255	cg25643199	0.56616147	0.488923077	0.627548209		5	94953181
cg25588792	0.583232041	0.640923077	0.502479339	LAIH2	19	55013946	cg07097026	0.56615893	0.515076923	0.600550964	TCTE1	6	44257942
cg04752329	0.583230345	0.539076923	0.590082645	TCERG1L	10	122912961	cg14608581	0.56614537	0.578153846	0.556473829	SNRNP27	2	70122240
cg16447652	0.583226107	0.556923077	0.579046361		17	54674338	cg05482066	0.56614113	0.554461538	0.549862259		8	2106183
cg10620681	0.583224412	0.533230769	0.571349862	PPIRNP2	7	158280480	cg18465049	0.56613096	0.538153846	0.546556474		3	125678342
cg04027757	0.583165077	0.531384615	0.598347107	POM121L1P	22	22987124	cg02717117	0.56612164	0.547692308	0.577410468	C2orf63	2	55458694
cg11086883	0.583161687	0.478461538	0.633812672	HIA-A	6	29908891	cg27094323	0.56612079	0.539384615	0.550964187		7	1216898
cg06714180	0.583093028	0.571076923	0.515151515	HOXC4	12	54409525	cg06314643	0.56611401	0.517538462	0.534986226	SPDY1	7	44039612
cg01530981	0.583082009	0.578769231	0.566391185	CAMK1G	1	209756892	cg18109320	0.56606908	0.539692308	0.569696997		5	87985809
cg02205181	0.583038779	0.568846154	0.554269972	NFKB2	10	104154811	cg17133750	0.56606908	0.578153846	0.552066116		6	168402302
cg18107355	0.583033694	0.515692308	0.604958678		10	37940899	cg04940251	0.56603687	0.561846154	0.52231405		16	88935765
cg09067818	0.583011655	0.530769231	0.562534435	PLEKH2	2	48864246	cg16026212	0.56602585	0.499076923	0.55922885		5	65218564
cg01594981	0.582970968	0.554769231	0.581818182		13	32272949	cg26421646	0.56601229	0.548923077	0.543801653	SERINC2	1	31895741
cg23270757	0.582967578	0.493230769	0.601652893	OR2H1	6	29427699	cg24351819	0.56598771	0.5730769	0.575757576	PRMT6	1	107598744
cg23921487	0.582920958	0.516923077	0.601652893	SNRNP35	12	123950649	cg07302051	0.56595889	0.560923077	0.532231405		16	51671357
cg13847928	0.582903157	0.579384615	0.49862259		5	180587498	cg18675735	0.56594702	0.563076923	0.508339945		10	13425015
cg17533458	0.582815851	0.492307692	0.602203857	FAM172A	5	92956826	cg07142285	0.56594618	0.518153846	0.556473829	APCDD1L	20	57042703
cg05956608	0.582803136	0.571692308	0.579614325		1	207999258	cg00693293	0.56594448	0.626461538	0.447933884	IRP1	12	57559808
cg05222986	0.582770078	0.539384615	0.570798898	RPS12	6	133136266	cg10585648	0.56594024	0.524615385	0.552066116	CTC3	1	156307318
cg14670938	0.582734478	0.492923077	0.620385675		5	1927225	cg10121518	0.56593855	0.440615385	0.560881543	ARRGEP4	2	131794483
cg24748548	0.582670057	0.585538462	0.539393939		10	135153961	cg13506281	0.565936	0.491384615	0.570247934	MTUS2	13	29914200
cg20786524	0.5826548	0.569538462	0.602203857		1	18954525	cg08913010	0.56592498	0.506461538	0.596143251	SYT9	11	7272653
cg01925738	0.582647171	0.523692308	0.599449036	PCDH83	5	140480770	cg23359714	0.56590973	0.548307692	0.496969697	SFRP1	8	41168336

ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg10191684	0.58253865	0.505838462	0.573002755	SORBS3	8	22423014	cg20211711	0.56590379	0.585538462	0.544352617	C17orf61	17	7307357
cg27381584	0.582581903	0.584923077	0.568595041	C17orf62	17	80407749	cg02049733	0.56589362	0.539076923	0.590082645	DGKG	3	185912253
cg04342623	0.582572579	0.508615385	0.598347107	SFMBT2	10	7376174	cg03078973	0.56588175	0.556769231	0.552066116	RIPK4	21	43183466
cg08817693	0.582541216	0.577230789	0.58953168	C12orf23	12	107348944	cg00844166	0.56588006	0.564923077	0.506336088	SLOC6A1	5	101834675
cg07090424	0.582529349	0.578769231	0.54600551		1	22535491	cg15546071	0.56587328	0.523076923	0.557024793	CCNT2	2	135691756
cg13042926	0.582521721	0.617538462	0.474931129	GUCY1A2	11	106890247	cg24215443	0.56585209	0.540307692	0.56584022	TGF21	6	134210598
cg23304249	0.582441195	0.578461538	0.559779614	CCDC86	11	60608951	cg21008709	0.56584107	0.523384615	0.55922865	SEC22C	3	42622950
cg22066321	0.582403899	0.524923077	0.599449036	DIDO1	20	61542198	cg15746870	0.56583937	0.523384615	0.549311295	DPP9	19	4689573
cg03152121	0.582395423	0.438153846	0.618732782		3	26146913	cg03377603	0.56583853	0.578153846	0.544903581	TOP2B	3	25639601
cg19999682	0.582386346	0.548	0.49862259		4	14854863	cg07114310	0.56580377	0.500923077	0.613222314		7	51538921
cg06552810	0.582380165	0.526769231	0.56584022		11	13128660	cg02886205	0.56578536	0.536307692	0.5644187328		4	64215583
cg00804237	0.582297097	0.68	0.41046832	SLC79A4	7	5337292	cg21082879	0.56574952	0.516923077	0.559779614	MARK2	11	63607728
cg05952543	0.58225641	0.585730769	0.538297011	MKRN3	15	23810378	cg03225546	0.5657385	0.491692308	0.616528926		1	148244719
cg10067510	0.582252172	0.539384615	0.560881543	FAM134C	17	40761703	cg01389917	0.56573511	0.508923077	0.601652893		17	1956668
cg18907202	0.582236915	0.546153846	0.563636364	GTF2A1L	2	48844971	cg24406898	0.56573427	0.508615385	0.61046832	COL1A2	7	94023308
cg00713813	0.582213181	0.496384615	0.568942149	CCDC41	12	94853658	cg04151012	0.56572494	0.524307692	0.49476584	ZNF512	2	27806529
cg18494043	0.582203857	0.603384615	0.555371901	PCDH9	9	65602795	cg09243445	0.56570684	0.516307692	0.561983471	ASB18	2	237123016
cg04842962	0.582200466	0.678461538	0.435812672	MIR518A	13	67558489	cg20338397	0.5657046	0.548	0.584022039		7	55916316
cg24330386	0.582162323	0.555076923	0.566391185	CHPF	6	43655489	cg00695090	0.56565798	0.492307692	0.572451791	MAG11	3	65781822
cg04831505	0.58214876	0.555384615	0.543250689	TBC1D15	2	220409543	cg13415566	0.5656495	0.463076923	0.566942149		10	129423240
cg02740950	0.582107226	0.532615385	0.568044077	TNKB	12	722323240	cg01354088	0.56564696	0.586461538	0.519008264	KIF11	2	10183140
cg07058971	0.582103836	0.508	0.579614325	FAM110C	6	32055534	cg02844341	0.56564357	0.553538462	0.533994904	STQ14	7	116593761
cg00616554	0.582101293	0.563076923	0.544903581	C22orf9	22	45608492	cg12730088	0.56564272	0.492615385	0.582309146	H1A-B	6	31324962
cg19113954	0.581970757	0.563692308	0.485399449	SERP5C5	4	25162716	cg23918676	0.56559356	0.55076923	0.577961433	C1D	2	58290497
cg03101664	0.581952108	0.578769231	0.584022039	SNX15	11	64794610	cg02563389	0.56557321	0.516615385	0.54600551	MED13L	12	116714239
cg18101029	0.581940242	0.516307692	0.575757576		6	101841615	cg20956131	0.56557152	0.555384615	0.544903581	C13orf27	13	103426330
cg06880511	0.581922441	0.556923077	0.534986226		11	15672258	cg00554878	0.56557152	0.516615385	0.577961433	OSR1	2	19553950
cg24421094	0.581915965	0.507692308	0.614325069	NTSR2	2	11810597	cg10552385	0.56556389	0.44	0.6	RB1	13	48878037
cg02430385	0.581906336	0.562153846	0.567493113	DUOX1	15	45421951	cg19706449	0.56555287	0.514153846	0.569146006	ATPAF1	1	47133793
cg09072408	0.581869888	0.532615385	0.54214876		11	118663568	cg05363479	0.56555118	0.562153846	0.541597796		7	47092529
cg00561194	0.58185463	0.571384615	0.584022039		10	38069720	cg24033103	0.56554355	0.594769231	0.485399449		19	37797500
cg07270285	0.581842763	0.531076923	0.58980716	PAP2B	2	71455274	cg06667753	0.56553846	0.53207692	0.570798898	LOC441601	11	50239550
cg21570988	0.581834287	0.626461538	0.543250689	PRTT1	6	32116841	cg07264062	0.56553338	0.555076923	0.564187328	ARHGEF10	8	1785958
cg02826525	0.581830896	0.560615385	0.582369146	LOC375196	2	39168885	cg07367434	0.56549354	0.485230769	0.559779614	COL9A2	1	40781135
cg14939821	0.581829201	0.625538462	0.470523416	DOCK1	10	128974730	cg15125036	0.56549184	0.531692308	0.564187328		3	23691152
cg24907270	0.581817334	0.563076923	0.571900826	MTX3	5	79297410	cg00036263	0.56549015	0.516307692	0.552066116	KIF26B	1	245857353
cg1151927	0.581807163	0.617230769	0.563085399	DIP2C	10	557886	cg06381135	0.56548676	0.508615385	0.554820937	INPP5A	10	134505548
cg07040013	0.581773343	0.431692308	0.638016529		10	132099553	cg27639408	0.56548591	0.492307692	0.600550964		4	155662375
cg00590012	0.581725769	0.55076923	0.544903581	RARG	12	74960699	cg08692111	0.56548337	0.538769231	0.495867769		19	173123
cg07241573	0.581708837	0.524	0.58677686	GAI1	18	74960699	cg25186680	0.56543675	0.476	0.497520661	IQGAP1	15	90931462
cg26969197	0.581699513	0.602461538	0.574655647		8	65485770	cg00557175	0.5654181	0.518769231	0.569146006	KATNAL2	18	44626898
cg08470863	0.581683408	0.516615385	0.635812672	ROBO2	3	77524776	cg07453718	0.56540793	0.585538462	0.553719008	ARSA	22	51066776
cg07160118	0.581649502	0.540923077	0.557024753	FAPBP	6	33282418	cg02665399	0.56539775	0.530153846	0.561983471	PRKAR1B	7	611060

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg18636319	0.581620682	0.548	0.555922865	CKAR	4	26492206	cg09254101	0.56539691	0.461538462	0.607162534	OTX2OS1	14	57361117
cg14077219	0.5815605	0.532	0.56584022	RP2	6	111303792	cg05244468	0.56536809	0.498769231	0.608264463		16	33959595
cg00446707	0.581541852	0.523384615	0.546556474	DIRA3	1	68512928	cg25058261	0.56536247	0.595692308	0.4925561983	TRX3	12	115109452
cg14455319	0.581530833	0.595076923	0.551515152	ANKK1	11	113258908	cg10631074	0.565363	0.539692308	0.528374656	CDC144B	17	18529187
cg24612420	0.58146506	0.538461538	0.564187328	ACLY	17	40075182	cg07125991	0.56535368	0.549230769	0.556473829	PLBD2	12	113796636
cg19550904	0.581476584	0.632923077	0.536639118	C14orf73	14	103569127	cg09035995	0.56531079	0.452307692	0.539118457	MCPH1	8	6323131
cg06137273	0.581417249	0.601538462	0.519592229	GRM7	3	6905031	cg19103770	0.56531214	0.576923077	0.554820937	TFPI2	7	93520566
cg00421990	0.581380901	0.519076923	0.588429752	WDR60	7	158656429	cg14766700	0.56530451	0.548	0.546556474	FAM111A	11	38911067
cg26671477	0.581363001	0.508923077	0.580716253		1	121258490	cg03454333	0.56527485	0.529846154	0.554769972	ZFP37	9	115819364
cg03425348	0.58131239	0.534461538	0.5184573		2	20441979	cg24328279	0.56526976	0.492846154	0.574855647		15	98632054
cg17611686	0.581309599	0.524923077	0.51683471	C20orf56	20	22547532	cg14092997	0.56526128	0.508923077	0.582320211	KHSRP	19	6425737
cg06616806	0.581281627	0.586153846	0.563636364		1	148556049	cg23462514	0.56524687	0.525846154	0.584373003	RNF212	4	1107585
cg17581697	0.581760437	0.463076923	0.614325069		17	21827358	cg22826239	0.56522992	0.540923077	0.564187328	IGF2BP3	7	23510123
cg13058786	0.581258741	0.523384615	0.594490358	HRNRP3	17	77097868	cg04919081	0.56521466	0.561846154	0.553168044	NIPA2	15	23034300
cg17041723	0.581140072	0.523076923	0.611019284		6	169271958	cg21211367	0.56521297	0.56230769	0.466666667		2	16209418
cg22417032	0.581128205	0.538769231	0.491460055	PTPRN2	7	157350867	cg07450552	0.56519763	0.494769231	0.638181818	DNAIC18	5	138775419
cg17504591	0.581118033	0.579076923	0.553168044	JAKMIP3	10	133945918	cg25348323	0.565185	0.524	0.584573003		7	45131862
cg20242136	0.581092604	0.554769231	0.537190083		19	22715697	cg15610027	0.56516974	0.485538462	0.580716253	DTNA	18	32173227
cg07090980	0.581084128	0.444923077	0.660055096	NYX	17	873255	cg07619542	0.56516974	0.509846154	0.574104683	IP6K1	3	49823894
cg00147172	0.581082433	0.493538462	0.630303030		22	46423604	cg09199225	0.56516211	0.459692308	0.549311295	AGER	16	32149260
cg1956686	0.581062433	0.502461538	0.613774105		2	241085024	cg06014332	0.56515702	0.486769231	0.61322314	BBS1	11	56278041
cg02510164	0.581055308	0.521846154	0.604407713	MCF21	3	183146421	cg17217059	0.5651477	0.609538462	0.49862259	FAM160A1	4	152329364
cg13908310	0.581048527	0.429538462	0.597796143	SLC35F3	1	224404073	cg10264188	0.56513922	0.54	0.569146006	KCNK4	11	64060409
cg14994247	0.581029879	0.508615385	0.590082645	SLC6A3	5	1407005	cg15244965	0.56513922	0.454461538	0.599449036	C5MD1	8	3429480
cg03513088	0.581012079	0.562769231	0.584022039		7	57472367	cg05194618	0.56512905	0.500923077	0.616526926	KLHL28	14	45432309
cg07646362	0.580971392	0.547384615	0.603856749		16	34208469	cg24244500	0.56512821	0.539384615	0.533884298		13	112815961
cg13335974	0.580968849	0.444615385	0.67107438	GRIN2C	17	72839489	cg13578160	0.56512312	0.516933077	0.558677686	RGS12	4	3409017
cg04435719	0.580962916	0.556	0.521763085		6	28912166	cg24599270	0.56510956	0.563384615	0.544903581		6	3105492
cg10445421	0.58092562	0.491692308	0.601101928		1	39284267	cg17077639	0.56510278	0.486153846	0.606060606		4	165898825
cg19907812	0.580837466	0.539076923	0.598347107		13	112757082	cg05767411	0.56510023	0.609846154	0.511294766	TRIM61	4	38965386
cg0375765	0.580827294	0.548	0.569146006		14	103551944	cg03780701	0.56509091	0.531384615	0.559779614	ADAM32	8	38965386
cg11366662	0.580799322	0.524307692	0.57630854	TYK2	19	10465257	cg25400864	0.56508582	0.538153846	0.553168044	INTS2	17	60005610
cg23964682	0.580767959	0.531692308	0.590082645	ANKMY1	2	241497415	cg19098796	0.56505192	0.577538462	0.541597796	ECE2	3	183993726
cg22381454	0.580730663	0.571384615	0.560330579	LDLR	19	11199903	cg10756475	0.56504005	0.492	0.583471074		4	1757242
cg26089705	0.580674719	0.580923077	0.53553719		20	43371550	cg22067472	0.56503242	0.509230769	0.567493113	CVB3B	16	69458746
cg16451431	0.580595041	0.556615385	0.562534435	NXPH4	12	57616936	cg20026651	0.5650231	0.524615385	0.580165289	CDC25B	20	3776748
cg15903185	0.580581479	0.588	0.539944904		4	111530471	cg00486708	0.5650231	0.555384615	0.560881543	ZNF32	10	44145111
cg26180501	0.580573003	0.584923077	0.540495868	PTPRN2	7	138363265	cg00909094	0.56500615	0.523384615	0.593939394		5	2334514
cg10024527	0.580522145	0.547692308	0.549311295		20	44882188	cg05930586	0.56497309	0.492	0.588429752		5	114971559
cg08883213	0.580508582	0.540923077	0.575206612		19	55728836	cg18374181	0.56497224	0.618461538	0.45785124		21	34401798
cg02274103	0.580505192	0.548	0.561432507	GAA	17	78078650	cg05842420	0.56496885	0.502153846	0.543801653		2	237073843
cg07807817	0.580499258	0.530769231	0.596694215	COG6	14	744746425	cg05173749	0.56495783	0.476923077	0.61570248	MAP1B	5	71452580
cg14252488	0.580497563	0.547384615	0.572451791	KIF26B	1	245361403	cg23779887	0.56493559	0.570153846	0.543301653	TMEM161A	19	19249481

ILMNID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_au	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg19746940	0.580478915	0.663384615	0.482093664	TBX5	12	114832429	cg18312782	0.56494088	0.554769231	0.533333333	TEX2	17	62341028
cg10227187	0.580422971	0.546769231	0.545454545	NOXO1	16	2030509	cg21308379	0.56491206	0.516615385	0.597245179		17	44918854
cg10570241	0.58041958	0.540307692	0.583471074		7	158785291	cg27447296	0.56491206	0.564615385	0.502479339	MRPL33	2	27994265
cg03491478	0.58033398	0.557538462	0.523415978	MAPK8IP1	11	45906914	cg14735364	0.56489426	0.548307692	0.512396694	NRF1	7	129273471
cg01249054	0.580363636	0.540307692	0.564738292		6	140269571	cg21782511	0.56488409	0.523076923	0.52231405	FGFR3	4	1808376
cg05917963	0.580356855	0.555076923	0.574855647	CHUK	10	101988339	cg27493812	0.56487307	0.538461538	0.547107438		2	241562757
cg13459393	0.580352617	0.570153846	0.513498623	ZNF28	19	53326091	cg19414122	0.56486035	0.564	0.57355372		8	18896631
cg04215511	0.580346684	0.531384615	0.587327824	CNTNAP4	16	76310816	cg05063274	0.56485103	0.564	0.52231405	PTPNI4	1	214638188
cg03700679	0.58030854	0.600923077	0.531680441	TTC308	2	178418002	cg01445793	0.56485018	0.522769231	0.571349662	CAMKK1	17	3771578
cg05245226	0.580292435	0.634461538	0.503581267	LMO1	11	8285086	cg27373657	0.5648434	0.515692308	0.557575758	AP1G1	16	71843039
cg17286695	0.580208519	0.572	0.570247934	C6orf168	6	99791063	cg26220594	0.56483153	0.550769231	0.527823691		1	19110978
cg11347033	0.580171223	0.407384615	0.561432507	PCDH85	5	140515629	cg00630724	0.56483068	0.58461538	0.527823691	PTX3	10	103991299
cg15340795	0.580144946	0.532307692	0.58953168	ZNF432	19	52551618	cg11528835	0.56482136	0.538461538	0.568595041	CNTNAP3	9	39132860
cg16203863	0.580119517	0.463384615	0.563636364	WDR73	15	85197975	cg03814321	0.56480855	0.500923077	0.553168044	TTC39A	1	51810337
cg27396141	0.580077983	0.525230769	0.563085399	NSD1	5	176693630	cg11271345	0.56478815	0.554153846	0.572451791	ACTBL2	5	56779826
cg00538248	0.58007544	0.586769231	0.526721763	PTPR	20	41817011	cg22717608	0.56477913	0.500307692	0.556473829	PDHX	11	34937665
cg13045351	0.580073744	0.617846154	0.548760331	ZSCAN16	6	28092048	cg19518539	0.56476372	0.413230769	0.650688705		7	21301723
cg11012543	0.580038991	0.548307692	0.598347107	ALAS1	3	52231691	cg233316161	0.56476118	0.412615385	0.629752066	ZNF605	12	133532874
cg16160105	0.579997457	0.437538462	0.607162534	KRT83	12	52715030	cg11528328	0.56475864	0.525538462	0.542699725	CHC6	21	36041683
cg14587335	0.579996609	0.532615385	0.551515152	HRNP3	17	77478600	cg05620415	0.56472982	0.524307692	0.55922865	HS3T6	16	1961808
cg27649194	0.579970333	0.555692308	0.590082645	CACNA2D4	12	1974168	cg20645941	0.56471795	0.547384615	0.502479339	FBXQ15	18	71815161
cg033961510	0.579949142	0.633846154	0.480991736		10	2878438	cg06426293	0.5647171	0.64	0.458953168	SYTL3	6	159151854
cg24928391	0.579931341	0.531076923	0.605509642	SOX17	8	55370336	cg21270847	0.56471541	0.514153846	0.540495868	PBX1	1	164604806
cg07076730	0.579916084	0.583846154	0.562534435		15	25018449	cg26188685	0.56469945	0.657230769	0.412121212	HDC	15	50540269
cg26892635	0.579852511	0.533846154	0.579063361	ORF1	16	3253043	cg05450236	0.56469591	0.532923077	0.539944904	GSTO2	10	106036853
cg01182309	0.579809282	0.588307692	0.530578512	CYP27B1	12	58159692	cg08530838	0.5646832	0.516307692	0.558126722		5	114193858
cg27656638	0.579754185	0.555076923	0.585123967	HOXB3	17	46628224	cg02116933	0.56466624	0.539692308	0.516804408		20	20346166
cg07060913	0.579753338	0.579076923	0.533884298		16	86537142	cg00406010	0.56465522	0.540307692	0.58292011		8	2178551
cg15406755	0.579747404	0.562769231	0.543801653	THP11	22	26908877	cg16607898	0.56464336	0.531307692	0.612121212	AFAP1	4	7770089
cg22777979	0.579737232	0.614692308	0.503581267	B3GAT1	11	134283252	cg25224222	0.56463403	0.619384615	0.494214876	FOSB	19	45971050
cg05888252	0.579729604	0.594769231	0.579063361		1	175857819	cg08368520	0.56462556	0.540307692	0.544352617	FOXK1	7	4747026
cg25554926	0.579716889	0.570461538	0.549862259	CPLX1	4	812611	cg11320749	0.56462301	0.485846154	0.618181818		10	130069573
cg19378133	0.579716042	0.579384615	0.539393939	A2BP1	16	6068835	cg22204479	0.56462132	0.524923077	0.539393939	NRX2-4	20	21378227
cg09350548	0.579713499	0.618153846	0.519008264		16	77270312	cg18323236	0.56461962	0.508615385	0.577961433	NIPAL3	1	24743029
cg05312104	0.579686374	0.495692308	0.615977961	NAV2	11	19402086	cg04443186	0.56461623	0.532307692	0.561432507		13	113614132
cg12235622	0.579671117	0.474461538	0.639118457	C1orf84	1	43836713	cg07091678	0.56461708	0.532	0.575757576	CPTAC1	10	99734120
cg13012310	0.579631278	0.579076923	0.531129477		14	102050627	cg08720806	0.56461369	0.501230769	0.603856749	PNODX2	11	125147671
cg20584555	0.579587201	0.577846154	0.54214876	BEST1	11	61717780	cg18498115	0.56461369	0.524923077	0.592386501		4	85411336
cg14422341	0.579556686	0.562769231	0.571046683	CD34	19	531225	cg16097124	0.56459674	0.571384615	0.550964187	TMED10	14	75642688
cg07573872	0.579542276	0.539384615	0.597245179	SPNO2	19	1126342	cg08065574	0.56458233	0.547692308	0.574655647	NR2F6	19	17346507
cg02403933	0.579535495	0.534230769	0.611570248	MTNR1A	4	187475669	cg22713360	0.56458148	0.514769231	0.591735537	MLEC	12	121125837
cg21062780	0.579526171	0.564	0.556473829		6	889772	cg01210372	0.56457046	0.515692308	0.590082645	BCDKK	16	31119640
cg10144336	0.579520237	0.563076923	0.540495868		10	3068123	cg252021359	0.56456876	0.649230769	0.47585124	ESYT2	7	158622466
cg18000391	0.579496503	0.609230769	0.489807163	PLEC1	8	14500375	cg14709479	0.56455266	0.587692308	0.497520661	TLL10	1	1108924

ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ auc	ILMNID	MAPINFO	CHR	GENE	mean_cv_ spe	mean_cv_ sen	mean_cv_ auc	GENE	CHR	MAPINFO
cg15717448	0.57949057	0.548615385	0.559922865	cg16362133	171234429	5		0.491692308	0.56454842	0.56454842	SLC11A2	12	51420474
cg09611598	0.57947875	0.423692308	0.63415978	cg05302877	20382487	11		0.461846154	0.56452723	0.56452723	SP6	17	45930106
cg002332679	0.579453274	0.597615385	0.556473829	cg13164087	1400588	5	SLC6A3	0.506461538	0.566391185	0.566391185		6	97832666
cg16799785	0.579422759	0.541538462	0.577410468	cg18343290	82381568	15		0.577846154	0.561983471	0.561983471		7	140218068
cg02692938	0.579421911	0.493538462	0.586225895	cg27368891	76032509	8		0.571384615	0.536088154	0.536088154	TNEM150C	4	83482706
cg2098780	0.579401568	0.480615385	0.541322314	cg13636926	7613764	17	EFNB3	0.562769231	0.562769231	0.562769231	ARID3C	9	34627940
cg01699727	0.579393092	0.500923077	0.584573003	cg16159904	11001757	16	CHTA	0.546447034	0.536088154	0.536088154	LRR27	10	134146220
cg18710250	0.579381225	0.564615385	0.570798898	cg15731920	113818925	4	ANK2	0.671384615	0.484848485	0.484848485	IASP1	17	37036919
cg20117368	0.579378682	0.602769231	0.533884298	cg00876704	49061546	19	SULT2B1	0.56447034	0.548760331	0.548760331	PRN1	19	14544684
cg20763134	0.579373596	0.615692308	0.543250689	cg25526717	33291578	6	DAXX	0.562461538	0.567942149	0.567942149	PFKL	21	45719687
cg00248633	0.579363967	0.563692308	0.562088399	cg14976971	46094335	19	GPR4	0.531384615	0.567493113	0.567493113	Clorf96	1	229478633
cg26533311	0.579363424	0.539076923	0.554269972	cg13843611	27206911	6		0.495076923	0.594490358	0.594490358	RNF186	1	20143089
cg07509323	0.579332062	0.577846154	0.563085399	cg03352287	72703562	14	RG56	0.564447034	0.555371901	0.555371901	FZD2	17	42635778
cg15585987	0.57928968	0.595384615	0.566942149	cg17049621	50824749	8	SNHG1	0.546447034	0.515692308	0.515692308	CHTA	16	11001939
cg21728101	0.579279508	0.631076923	0.476033038	cg06585907	1979360	7	MAO11	0.492615385	0.604407713	0.604407713		16	11001939
cg27236770	0.579265099	0.468615385	0.571900826	cg08415582	18870922	19	CRIC1	0.569538462	0.543250689	0.543250689		8	57030523
cg01614102	0.57924306	0.516923077	0.609366391	cg24667575	115134918	12		0.56440771	0.56969697	0.56969697	NEURL18	8	172097064
cg25388061	0.579203221	0.562461538	0.549862259	cg11126351	176215887	5		0.562461538	0.564187328	0.564187328	KIAA0391	14	35594356
cg02330371	0.579162534	0.539692308	0.571349862	cg14677084	21380496	15		0.540615385	0.569146006	0.569146006	PCDHGA4	5	140857941
cg12292216	0.5791032	0.632307692	0.505785124	cg08396096	37388168	17		0.493846154	0.596143251	0.596143251	VPS37C	11	60828446
cg045660357	0.579057427	0.555384615	0.555922865	cg10960055	37498025	1	GRK3	0.56434753	0.562534435	0.562534435	DCAKD	17	43112088
cg14553794	0.579033694	0.585538462	0.521212121	cg09874796	53895291	12	TARBP2	0.547692308	0.562534435	0.562534435		6	28664226
cg24107021	0.578990464	0.589230769	0.504132231	cg09650880	96297058	13	DZP1	0.517846154	0.582369146	0.582369146	DIP2C	10	661730
cg1592613	0.57897775	0.524307692	0.590082645	cg19335742	78200484	15		0.56433128	0.569146006	0.569146006	PAK6	15	40566880
cg10506102	0.578975207	0.500307692	0.579063361	cg04741860	40142930	4	N4BP2	0.56433128	0.597796143	0.597796143		2	558749
cg16692735	0.578969273	0.540615385	0.571349862	cg19306761	135259660	10		0.532	0.550964187	0.550964187		7	149732847
cg13793678	0.578961544	0.532307692	0.559779614	cg03873322	48832808	4	OCIAD1	0.54214876	0.54214876	0.54214876	NMBR	6	142409202
cg20672044	0.578945539	0.585230769	0.404407713	cg21414905	72096498	16	HPR	0.586153846	0.497520661	0.497520661		6	11901382
cg10172584	0.578925196	0.579076923	0.548209366	cg23465978	19639581	19	YIEFN3	0.548615385	0.563085399	0.563085399	MGMT	10	131330293
cg10443987	0.578913329	0.492	0.55922665	cg21931784	64023712	7	ZNF680	0.594353846	0.578512397	0.578512397		11	28030401
cg17405812	0.578897224	0.500615385	0.604407713	cg23632840	17085500	10	ADAR2	0.477538462	0.604407713	0.604407713	C2orf94	20	10414722
cg03022891	0.578886205	0.453846154	0.656198347	cg15713378	1947791	11	TNNI3	0.524307692	0.590082645	0.590082645	DIP2C	10	709839
cg18582073	0.578877728	0.540307692	0.602754821	cg05720027	85202042	15	NMB	0.56426362	0.523986942	0.523986942	FND1	6	159654415
cg16511494	0.578876881	0.429538462	0.622038567	cg17919132	209788337	1	LAMB3	0.56425175	0.548760331	0.548760331		13	112858755
cg0636244	0.578837042	0.476615385	0.594490358	cg16985667	3572706	10		0.56424836	0.523692308	0.523692308	KRTAP4-5	17	39306789
cg07883772	0.578777707	0.506307692	0.643801653	cg07211140	93064234	11	CCDC67	0.56424836	0.532	0.532	WT1	11	32455025
cg15143338	0.578773469	0.406769231	0.696969697	cg11208348	101039724	14		0.5642445	0.517230769	0.517230769	ZNF277	7	111849025
cg19500098	0.578707553	0.655384615	0.528374656	cg03278924	10300506	13		0.500307692	0.552066116	0.552066116	ELP2	18	33709663
cg06110575	0.578617504	0.565230769	0.537190083	cg21297282	45229649	2		0.56419326	0.595076923	0.595076923	STGAINAC5	1	77525228
cg15603354	0.578607332	0.478461538	0.598347107	cg14538441	107578208	11	SLN	0.56418987	0.525230769	0.525230769	PIGY	4	89444921
cg10632728	0.578547064	0.485230769	0.6	cg08685733	18875315	1		0.56416698	0.500307692	0.500307692		17	42215609

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg01215957	0.578525111	0.556	0.571900826	FAM168A	11	73309852	cg15352671	0.5641602	0.532307692	0.561983471	LOC255512	11	1331497
cg18289746	0.578462386	0.592923077	0.568595041	TUBE1	6	112408679	cg079245535	0.56415342	0.523692308	0.573553719	HBC1D1	4	37893355
cg17366353	0.578374232	0.563076923	0.578512397		19	5477638	cg197263355	0.56415088	0.517538462	0.561983471	BTBD11	12	108027812
cg14810735	0.578331002	0.625846154	0.455198347	ANGPT1	8	108441547	cg066395538	0.56414495	0.486461538	0.597796143	OPCML	11	133373892
cg17030482	0.578308116	0.507384615	0.594490358		8	11539330	cg15647005	0.5641424	0.538153846	0.57630854		12	3579236
cg15397231	0.578294554	0.530153846	0.570247934	HLA-C	6	31239074	cg101006333	0.56413562	0.524923077	0.562334435	ZNF443	19	12552133
cg14678162	0.57825302	0.585538462	0.538292011		2	104640450	cg078727333	0.56411782	0.430769231	0.655096418	MTFMT	15	65322308
cg16832666	0.578237762	0.555384615	0.563085399	ORT1L1	1	248005356	cg02282041	0.56411358	0.430769231	0.655096418	CDH13	16	82908556
cg23194912	0.578230134	0.414461538	0.63415978		2	3034048	cg19556079	0.564110426	0.561846154	0.548709366	BAT4	16	31632638
cg14658900	0.578229286	0.562769231	0.571349862	ZNF174	16	3450923	cg13402656	0.56410426	0.40923077	0.707438017	DLGAP2	8	1511478
cg14239592	0.578211465	0.539076923	0.552066116	ZNF274	14	10635451	cg14392925	0.56408137	0.478153846	0.574104683	NRXN2	11	64490736
cg17113548	0.578180971	0.406769231	0.607713499		14	10635451	cg06370862	0.56407459	0.462153846	0.622038567	TRAPPC10	21	45431089
cg09411798	0.578176732	0.585538462	0.544903581	ZNF639	3	179041374	cg24702522	0.56408985	0.523692308	0.570661157	TFAP2D	6	50680748
cg15803018	0.578170799	0.572	0.560330579		11	45744972	cg00040027	0.56408561	0.530769231	0.54600551	SLC25A12	2	172747482
cg25369980	0.578124179	0.594461538	0.558126722	ENC1	5	73937328	cg21751231	0.56404323	0.508	0.56584022	TCEB1	8	74884456
cg05540369	0.578114007	0.468307692	0.64077135	ZNF75A	16	3355553	cg09958470	0.56404153	0.486153846	0.572151791	SLC16A12	10	91294839
cg00813752	0.578110617	0.545538462	0.583471074	CHGA	14	93389146	cg2651171	0.56403645	0.547384615	0.5211763085	EPHA10	1	38218194
cg05169951	0.578096207	0.50846154	0.603856749	KREMEN2	16	3017955	cg06778672	0.56404747	0.603076923	0.526178799	GRHL3	1	24645184
cg20560392	0.578080949	0.508923077	0.554820937	ANKK1	1	65617331	cg25739366	0.56404323	0.548307692	0.580716253	APLNK	11	57004007
cg00841679	0.578054844	0.531384615	0.568044077		8	94230311	cg17186373	0.56404323	0.508	0.56584022	TCEB1	8	74884456
cg0584664	0.57802331	0.548923077	0.596143251	INS-IGF2	11	2154934	cg03271761	0.56404238	0.477538462	0.56804077	KCNIP4	4	21699209
cg06184252	0.577989405	0.554769231	0.568044077	ATP6V0A1	17	40610858	cg00708678	0.56404153	0.486153846	0.572151791	SLC16A12	10	91294839
cg25660566	0.577952956	0.577538462	0.530027548		7	155262852	cg1717442	0.56403645	0.547384615	0.5211763085	EPHA10	1	38218194
cg14382976	0.577932613	0.556615385	0.533333333	ASZ1	7	117067624	cg09322349	0.56403475	0.526153846	0.571349862	GRHL3	1	24645184
cg15890623	0.57786904	0.478153846	0.63415978		5	5883984	cg22496723	0.56401611	0.537846154	0.574104683	CACNG2	22	36960779
cg05270572	0.577832592	0.555076923	0.555371901	STEAP3	2	119981334	cg21838488	0.56398729	0.437846154	0.528374656	RNF219	13	79234435
cg27331871	0.577820725	0.532	0.577961433	INS-IGF2	11	2171478	cg02143895	0.56398305	0.54769231	0.547107438	HRNB3	17	77394015
cg01093138	0.577813944	0.541538462	0.577961433	CDH13	16	83374884	cg21252727	0.56397033	0.541230769	0.530578512		12	28149882
cg14227837	0.577786819	0.546153846	0.579514325	SHISA2	13	26620952	cg10180297	0.56395423	0.537615385	0.574655647	H3S3T2	16	22827393
cg26685515	0.5777772409	0.524923077	0.583471074		1	181394054	cg06773116	0.56395338	0.522769231	0.508539945	KCNIP4	4	20884431
cg23181900	0.577760542	0.571076923	0.554269972	COL6A6	3	130368798	cg10414985	0.56395168	0.447076923	0.56969697		10	102978033
cg1149163	0.577728332	0.562461538	0.580716253		5	2387826	cg13814950	0.56394959	0.563769231	0.559779614	DAPK1	9	90112515
cg07916377	0.577709684	0.568538462	0.542699725	FECB	18	55254112	cg07572984	0.56394151	0.601846154	0.529476584		14	30858152
cg05741219	0.577680865	0.517846154	0.602203857	SF3B14	2	24295212	cg18677906	0.56393049	0.5	0.570798898	GALNT9	12	132851967
cg03558837	0.577673236	0.570461538	0.553719008	ESPNL	2	239029375	cg17213843	0.56391693	0.538769231	0.520110193	HERC3	4	389513558
cg01042202	0.577663912	0.508923077	0.566391185	PCDH6	5	140242159	cg19052128	0.56390591	0.487076923	0.628650138		13	76444293
cg23331340	0.577655435	0.524	0.577961433	KCNK3	2	18058900	cg20873718	0.56389828	0.577230769	0.424242424	IFNG	7	2559473
cg05983061	0.577625768	0.595384615	0.52892562	MOBP	3	39544192	cg05221664	0.56389659	0.563076923	0.571349862		10	133484920
cg13571283	0.577607968	0.548923077	0.575757576	MRPL23	11	1975380	cg22584065	0.5638915	0.507692308	0.597245179	FGF14	13	103052828
cg27809937	0.577585082	0.46846154	0.631404959	BCK	2	24249273	cg10008501	0.56386946	0.477846154	0.6		8	9763558
cg06709178	0.577580843	0.448	0.607713499	INPP5B	14	56991775	cg16968865	0.56385251	0.548615385	0.549311295		18	45534737
cg2128892	0.577551176	0.544923077	0.555371901	PTPRG	3	38328759	cg00414835	0.56382708	0.595692308	0.526721763	GDF10	10	48439452
cg27040468	0.577503404	0.540307692	0.56384022	BICC1	10	60456715	cg25444002	0.56382539	0.454153846	0.56969697	FGF14	13	102520143
							cg24703717	0.56382369	0.563076923	0.556473829	HLA-F	6	29691168

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg22166489	0.577480822	0.564	0.530072548	WDR66	12	122356033	cg23636941	0.56376551	0.508615385	0.583471074	AHCY	20	32899639
cg203339553	0.577477432	0.424923077	0.596143251	C2orf3	2	75937869	cg05766510	0.56376012	0.587076923	0.536088154	PTPRN2	7	157454062
cg25055120	0.577468955	0.578769231	0.499724518		2	13448495	cg18595005	0.56375534	0.580923077	0.500275482	NXN	17	783036
cg19276110	0.577447764	0.532307692	0.578859504	PIA2G5	1	20396593	cg26873935	0.56373554	0.584661538	0.474380165		16	72459516
cg04852280	0.577440136	0.564307692	0.553168044	ZNF593	1	26466234	cg14688662	0.56372197	0.546769231	0.549662259		7	1688174
cg23247171	0.577433355	0.562461538	0.552066116		14	105649747	cg07674095	0.5637118	0.539076923	0.557575758		4	3702475
cg10202558	0.577424031	0.556307692	0.561432507	TMEFM192	4	166034152	cg19117501	0.56370502	0.569846154	0.555922865	GLI1L2	11	134206074
cg23415538	0.577411316	0.523384615	0.601652893	SLC25A39	17	424021174	cg10070101	0.56369739	0.580615385	0.507988981		17	78963290
cg01223757	0.577396906	0.546769231	0.530578512		13	21520203	cg15541040	0.56368892	0.492923077	0.61322314		2	3486749
cg07461072	0.577389277	0.560923077	0.562534435	GSGL	16	28073940	cg27631817	0.56367875	0.549538462	0.501928375		16	10060938
cg19847637	0.577358762	0.547076923	0.55922865	RASGRP3	2	33738782	cg11267630	0.56366857	0.524307692	0.57079898		17	46811527
cg15627457	0.577347743	0.532307692	0.574655647		8	141535348	cg10464773	0.56366773	0.540923077	0.551315152		16	164171506
cg00901051	0.5773452	0.563076923	0.526721763		1	208132490	cg26889647	0.56366773	0.463076923	0.545454545	CLK1	2	201729339
cg26704289	0.57734181	0.58	0.507438017		15	73611289	cg13617192	0.56366095	0.532307692	0.5184573		12	34534365
cg08389686	0.577332486	0.524307692	0.620936639	GPH2	5	179763678	cg02323875	0.56365247	0.515692308	0.557024793	TXLNA	1	32645332
cg23080818	0.577317228	0.586153846	0.511294766		2	161504905	cg02338553	0.56364823	0.593846154	0.573415978	LOC146481	16	34712713
cg10935612	0.577313838	0.573615385	0.548556474	UBE2OL1	5	6450951	cg03633114	0.5636423	0.569846154	0.49201019	CYR1	21	27945841
cg03997764	0.577300275	0.499692308	0.575206612	FAM968	16	66988417	cg18826958	0.56362819	0.455384615	0.585674931		22	46275235
cg19757176	0.577273999	0.562769231	0.527272727	TPN3	1	154128041	cg10001282	0.56361856	0.563076923	0.558126722	NEUROG3	10	7133478
cg16472210	0.5772757046	0.511692308	0.561432507	GBE1	3	91810671	cg21128600	0.5635889	0.617846154	0.50523416	ZNF705D	8	11972998
cg26122626	0.577252808	0.571384615	0.544352617	PRDM13	6	100058086	cg11282156	0.56358466	0.547692308	0.541597796	SYNPR	3	63428840
cg03578473	0.577244331	0.515384615	0.581818182	NEUROD1	2	182546504	cg19378036	0.56357618	0.516307692	0.570798898	HNF1B	17	36105364
cg07154956	0.577240941	0.577846154	0.550964187	MA2K3	17	21187834	cg13507084	0.5635711	0.538769231	0.578512397	CNTFR	9	34555042
cg07424214	0.577237355	0.634769231	0.451239669	LOC255167	5	6585532	cg25973354	0.56355669	0.515384615	0.566942149		19	1395646
cg02941741	0.57723416	0.577230769	0.543801653	NGF	2	233793921	cg22880770	0.56355584	0.509528462	0.573553719	NEURL3	2	97166341
cg03564727	0.577232465	0.539692308	0.56584022	KCNAT	19	49577706	cg23991721	0.56351939	0.563384615	0.578512397	C19orf25	19	1237877
cg13495235	0.57720534	0.593230769	0.524517906	C9orf24	9	34379714	cg06821774	0.56351685	0.532307692	0.564738292	LMNL	3	197686168
cg14032218	0.577177368	0.547692308	0.564738292	METTL13	1	171755066	cg01959541	0.56348125	0.632615385	0.450137741		6	159291228
cg15353061	0.577172282	0.444	0.647933884	CYP2E1	10	135342413	cg03721622	0.5634787	0.556307692	0.560881543		8	78478065
cg26546664	0.577170587	0.593538462	0.514600551		16	34407868	cg20685672	0.56347362	0.508615385	0.574104683		2	181987552
cg06374079	0.577162958	0.523384615	0.590082645		6	27187908	cg18135206	0.56345751	0.530076923	0.57630854	TECP2	14	102964638
cg08932236	0.577144431	0.5853846154	0.561983471		12	116008258	cg16521978	0.56344904	0.515384615	0.564738292		6	33129673
cg1229184	0.577142615	0.594769231	0.509641873		6	28710807	cg11533311	0.56344649	0.484923077	0.626997245		5	66541588
cg00737264	0.577089214	0.462769231	0.585123967	SMOC2	6	169049498	cg20993503	0.5634431	0.541230769	0.551515152	LOC387763	11	43964098
cg10000841	0.577088366	0.421846154	0.63030303	CSTII	20	23419162	cg06775570	0.56343547	0.415076923	0.619283747		11	119201635
cg14270494	0.577074804	0.46	0.523966942		18	5605013	cg06945523	0.56342615	0.617230769	0.479338843	SHISA4	1	201857621
cg01835665	0.577063785	0.58461538	0.568595041	HEXIM1	17	43224550	cg09432202	0.56341344	0.55076923	0.561432507		16	163768551
cg15325872	0.577044289	0.586461538	0.562088399	RNASIH1	2	3606125	cg26504248	0.5634082	0.523076923	0.575757576	SNORA76	17	62223554
cg13143830	0.577037508	0.436615385	0.656198347	C2orf69	2	200778665	cg18492757	0.56338716	0.557230769	0.561983471		8	2548901
cg13540960	0.577005298	0.539692308	0.574655647	WTL	11	32450244	cg23406436	0.56334478	0.438769231	0.563085399	ZNF253	19	19975451
cg16824113	0.57700106	0.54	0.598347107		12	132166485	cg03614649	0.5633315	0.55076923	0.532782369	AFG3L1	16	90037935
cg22789915	0.576982412	0.570769231	0.579063361	COL18A1	21	46850449	cg20463151	0.56332443	0.463384615	0.619834711		8	11546573
cg26925981	0.57697563	0.539384615	0.591735537		8	73913918	cg25508322	0.56332359	0.541230769	0.573002755		13	86268291

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg02380278	0.57697563	0.539384615	0.591735537	ZNF366	5	71756236	cg23046586	0.56332104	0.487384615	0.599449036	ZFP14	19	36833695
cg02392634	0.576952744	0.524615385	0.602754821	ANK1	8	41733831	cg14990366	0.56331002	0.509846154	0.531129477	CACNB4	2	152831129
cg04442406	0.576944268	0.586769231	0.52892562	PLSCR2	3	14637398	cg13071333	0.56328036	0.5234315978	0.5234315978	LCE3A	1	152595832
cg27135645	0.576901886	0.493846154	0.543230689		3	164056753	cg19300401	0.56327697	0.594153846	0.484848485		6	16962712
cg05301470	0.576890867	0.523384615	0.568044077		6	28558006	cg01511740	0.56326764	0.550153846	0.540495888		8	65291990
cg19148440	0.576884933	0.648307692	0.472176309	HLI7RE	3	9957031	cg03921886	0.56326171	0.563384615	0.47107438	PCM1	8	17781617
cg01818121	0.576883238	0.578769231	0.550413223	GRAMD2	15	72476917	cg11008866	0.56326086	0.523384615	0.55922865	CASR	3	121902442
cg03456771	0.576857809	0.579076923	0.538842975		7	56242072	cg09860783	0.56322611	0.515384615	0.566942149	ZC3H12D	6	149777984
cg27406623	0.576838313	0.601230769	0.514600551		2	236164597	cg05601199	0.56322356	0.524307692	0.579063361	AHRR	5	389490
cg27302994	0.576834075	0.516923077	0.608264463		1	30240265	cg14092571	0.56321085	0.524	0.55922865		14	90743983
cg05626549	0.576773045	0.539076923	0.558126722	PRKCDP	11	6341895	cg17881353	0.56320068	0.578769231	0.544903581	WNT9B	17	44928210
cg1367404	0.576714558	0.603076923	0.477134986		3	27674614	cg14646974	0.56318457	0.547692308	0.551515152	HSD17B7P2	10	38645378
cg0356090	0.576705539	0.609846154	0.509090909		16	1328562	cg26730347	0.56318457	0.516923077	0.638016529	SLCSA1	22	32438947
cg15246706	0.576695063	0.516	0.58677686		16	1060558	cg05977230	0.56316338	0.483692308	0.579063361	LRPS	11	68115417
cg16464486	0.576693367	0.526153846	0.514049587	JAKMIP3	10	133929519	cg22937354	0.56315999	0.548923077	0.539944904	OR5D16	11	55606216
cg11860629	0.576645052	0.445230769	0.663707989	LOR	1	153234268	cg23231779	0.56315914	0.431076923	0.654545455	THEM4	1	151887143
cg26658272	0.576639118	0.485846154	0.6	SLCSA18	5	1241765	cg05274242	0.5631405	0.585538462	0.561432507		16	31540559
cg01159980	0.576624709	0.595692308	0.544903581	FAM1348	5	16599017	cg14442405	0.56313965	0.470153846	0.627548209		2	3045331
cg07786675	0.576622166	0.563076923	0.548209366	SFN	1	27189805	cg03372209	0.56312778	0.547692308	0.550413223	ZEAT	8	135725058
cg15308271	0.576610299	0.524615385	0.585674931	GCKR	2	27720440	cg25191241	0.56311337	0.524615385	0.590082645	UMPS	3	124448994
cg03387145	0.576534011	0.601230769	0.560881543		15	100465995	cg21390082	0.56311083	0.486153846	0.612121212	SIN12L	17	33842255
cg24768630	0.576518754	0.439076923	0.649035813	HHLA2	3	108041508	cg10625367	0.56310066	0.452923077	0.614876033		12	6241975
cg20443138	0.576500954	0.547384615	0.509641873	DUSP19	2	183943496	cg10915263	0.56309981	0.501538462	0.583471074	CD163L1	12	7592647
cg01312445	0.576467048	0.524615385	0.551515152		16	86536684	cg17806623	0.56309218	0.55538462	0.536639118	KL	13	33590002
cg14821900	0.576452638	0.555384615	0.550964187	KCNK7	11	65363274	cg08483376	0.56308286	0.548307692	0.549862259	EVX1	7	27285136
cg20365687	0.57644501	0.454153846	0.566391185		13	60834809	cg09328979	0.56308286	0.548	0.547658402		16	88700622
cg23616527	0.57643399	0.571076923	0.581267218		10	50961649	cg06900355	0.56307947	0.569846154	0.531680441		5	150554476
cg24625984	0.57641958	0.445230769	0.593939394	LOR	17	130279707	cg25454379	0.56306336	0.585538462	0.531129477	DPFA5	18	76069154
cg16044948	0.576403475	0.587692308	0.511294766		1	197879380	cg18057665	0.56305319	0.540307692	0.582369146		6	74064140
cg16040504	0.576394999	0.524615385	0.584022039	HPCAL4	1	40157202	cg02979703	0.56305149	0.479692308	0.573002755		8	37558858
cg08983961	0.576296673	0.408923077	0.653443526	NPB	17	79859947	cg23117796	0.56305065	0.578769231	0.519592239		15	101389700
cg14449259	0.576266158	0.514153846	0.62369146	HRNBP3	17	77174290	cg01416043	0.56304217	0.540615385	0.568595041	KCNF1	2	11053694
cg08766812	0.576258529	0.624615385	0.467768595		2	57987283	cg13703866	0.56302776	0.491384615	0.615977961	TPO	2	1479523
cg12297125	0.57621869	0.593538462	0.499173554	PRDM16	1	2998522	cg09809504	0.56301505	0.529846154	0.521763085	CACNA1H	16	1227402
cg06525969	0.576183937	0.515076923	0.585674931		16	88453808	cg08397273	0.56301166	0.460307692	0.619283747	ZNF544	19	58740455
cg23690528	0.576171223	0.509846154	0.611570248	SGCE	7	9428570	cg03495660	0.56300826	0.562769231	0.52112121		16	1074500
cg05131382	0.576170375	0.563692308	0.573553719		3	138661281	cg24208325	0.56299725	0.555692308	0.531680441	ODZ4	11	78780863
cg10796815	0.576156813	0.484307692	0.563636364	GPR88	1	101004474	cg21219373	0.56298538	0.571692308	0.515151515	PPP1R13L	19	45885940
cg8840250	0.576149184	0.424307692	0.56236915		5	1855934	cg14891195	0.56297605	0.571692308	0.565289256	KHLH14	18	30933000
cg14225039	0.576143184	0.554769231	0.564187328		7	125916407	cg02144238	0.56297351	0.497307692	0.570798898	ACCN4	2	220591374
cg00406909	0.576139013	0.532615385	0.566942149		2	175194079	cg25753693	0.56296419	0.617270769	0.517355372	KIRS	19	51504734
cg17229388	0.576118669	0.594769231	0.576859504	C13orf16	13	111973143	cg02599361	0.56296334	0.467153846	0.626465281	ADAMT52	5	176548373
cg17050807	0.576112736	0.562461538	0.504132231		22	42548356	cg24793722	0.56296249	0.517538462	0.531129477		4	824416
cg24864707	0.576067811	0.565538462	0.555922865		11	121916987	cg07588819	0.56295571	0.570461538	0.577772727		2	132110512

ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg14033514	0.576060182	0.462461538	0.608815427		17	19483721	cg011175812	0.56294554	0.464307692	0.512947658	KCNA6	12	4919001
cg105382687	0.576057539	0.501846154	0.581818182		8	48091424	cg027432699	0.56293537	0.484307692	0.597796143	GPN1	2	27873401
cg176002719	0.576053401	0.445230769	0.474380165	VWDE	7	12444013	cg027723433	0.56293113	0.430461538	0.602203857	GPR83	11	94133505
cg18921954	0.576049163	0.603076923	0.520110193		5	33938909	cg10854819	0.56292859	0.56307692	0.536088154		15	31515852
cg133280155	0.576036448	0.445230769	0.675482094		14	72219812	cg04687616	0.5629252	0.540307692	0.56493113	11-Sep	4	77929804
cg03723481	0.576021191	0.444923077	0.625344353	MAD1L1	7	20717123	cg02250046	0.56290061	0.547692308	0.5494903581	CPN1	10	101825185
cg08402652	0.575969485	0.523384615	0.592286501	AMOTL1	11	94501461	cg021590985	0.56290061	0.408615385	0.66446281		10	123852191
cg17083209	0.57595338	0.556615385	0.575206612	LPHN2	1	82274574	cg14159747	0.56289129	0.532923077	0.525068871		11	113255604
cg04107099	0.575945751	0.468538462	0.612121212	ELOVL4	6	80657826	cg06357691	0.56288875	0.460307692	0.581818182	ESRRA	11	64072954
cg2739258	0.575900826	0.546153846	0.564738292		5	180526995	cg05513226	0.56288281	0.540923077	0.527823691		12	117147009
cg26367197	0.575894893	0.563076923	0.547107438		18	76307273	cg24153199	0.56287179	0.492307692	0.6	SO52	14	50698328
cg05300717	0.575759235	0.570153846	0.560330579	DKFZp761E1	11	65546210	cg05839532	0.5628701	0.533230769	0.557024793	BRUNOL4	18	35143863
cg01180479	0.575869464	0.586769231	0.542699725	MAP3K11	11	65374833	cg24512544	0.56285908	0.562769231	0.531129477	EIF2C2	8	141588943
cg18988170	0.575850816	0.454769231	0.627548209	CABYR	18	21719263	cg14808329	0.5628523	0.648	0.400550964	PHF15	5	133863570
cg05969021	0.57581691	0.571384615	0.537741047	ID3	1	23888005	cg03847932	0.56284976	0.554153846	0.528374656	PPP4R2	3	73045556
cg18821144	0.5757847	0.617538462	0.520661157		12	68881398	cg10836036	0.56284552	0.493230769	0.55261708	FDPS	1	155278714
cg24616795	0.575780462	0.58	0.563636364	SFMBT1	3	53032891	cg14168530	0.56284298	0.515692308	0.571349862		2	45155991
cg27333018	0.575730451	0.618461538	0.45785124		19	2897514	cg11491998	0.56282772	0.476923077	0.614325069		6	30419493
cg26209816	0.575724518	0.492923077	0.560881543	ADAP1	7	978305	cg00063649	0.56280329	0.507692308	0.575757576	CI5orf61	15	67813698
cg25481705	0.57570926	0.516307692	0.582369146	KLC3	19	45850053	cg27452939	0.56280144	0.533538462	0.549311295		2	156698957
cg07947490	0.575679593	0.556	0.514049587	ACVR1C	2	158484692	cg01278873	0.56277601	0.571384615	0.537782369	ZNF233	19	44764101
cg14447047	0.575646535	0.538461538	0.563085399	RUNX2	6	43389860	cg26380578	0.56277432	0.540615385	0.577961433	ARFGAP1	20	61916036
cg17611045	0.575632125	0.592615385	0.49476594	FGF12	3	152789245	cg19814830	0.56276838	0.484	0.639669421	PCDHGA2	5	140735545
cg10196128	0.575614325	0.548307692	0.572451791	ZNF284	19	44576295	cg18810983	0.56276669	0.524	0.58044077	C3orf59	3	192632111
cg08688169	0.575597372	0.561846154	0.546556474	P2RX7	12	121570081	cg09480837	0.5627616	0.554769231	0.548209366	PLCE1	10	95789501
cg01352656	0.575592287	0.531384615	0.575757576		2	118561080	cg14791747	0.56275736	0.649230769	0.445730028		5	124309096
cg07006673	0.575531257	0.565538462	0.537741047	HNF4A	20	43029732	cg06755871	0.56275397	0.525846154	0.55922865		2	356507
cg27393285	0.575521933	0.546461538	0.554820937		19	57277288	cg10804974	0.56275143	0.564	0.540495868	PCDH814	5	140603270
cg02131512	0.575478703	0.524923077	0.572451791	SEZ6L2	16	29882803	cg17718276	0.56274465	0.460307692	0.613272314	TIME132D	12	129648291
cg10717312	0.575406554	0.571076923	0.534986226	SNED1	2	241976244	cg19989581	0.56273617	0.516307692	0.566391185		15	27924256
cg06769094	0.575406654	0.538769231	0.554820937	SF-RP1	8	41151382	cg26429499	0.56271159	0.476923077	0.590082645	SHANK2	11	70563792
cg04026675	0.575363424	0.562153846	0.548556474	CDKN2A	9	21974514	cg1154964	0.56270566	0.432923077	0.53553719	HFCW1	7	43484715
cg20602954	0.575332062	0.601538462	0.514049587	NKAIN3	8	63776762	cg08950364	0.56270057	0.554769231	0.539944904		1	108039344
cg2224628	0.575321043	0.486153846	0.607162534	CCDC1028	18	66381506	cg17378535	0.56269718	0.476	0.55979614		17	41406661
cg06740275	0.575317652	0.546769231	0.571900826	TSR1	17	2239739	cg03837036	0.56267938	0.431692308	0.618181818		12	63015414
cg0026576	0.575297309	0.462769231	0.612121212		6	4481538	cg08975528	0.56267769	0.571076923	0.503581267	ZBTB12	6	31867700
cg18723873	0.575293071	0.523384615	0.571900826	CYP1A1	15	75018222	cg16680645	0.5626743	0.554153846	0.541046832		1	32254925
cg17842426	0.57528968	0.539076923	0.564187328	PSK	10	124739963	cg11339341	0.5626709	0.515692308	0.583471074	LOC645323	5	87969130
cg22423398	0.575283747	0.602153846	0.548209366	NCRNA0016	2	132919865	cg26807620	0.56266921	0.571076923	0.5505964187	DIGAP2	8	1497078
cg04723073	0.575276118	0.524307692	0.563085399		2	183473088	cg16115627	0.56266667	0.495692308	0.546556474	PPP6	7	154684433
cg06211550	0.575271032	0.492923077	0.577410468		16	3230726	cg19270089	0.56265565	0.493538462	0.566391185	CADPS	3	62440705
cg01145232	0.575269337	0.556	0.571349862	RAET1G	6	150245071	cg25558663	0.56264632	0.579692308	0.431955923		20	44882514

ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_auc	ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg20749955	0.575254079	0.556923077	0.554820937	cg23036947	0.56764124	0.616307692	0.507988981	MXRA8	1	1294018
cg08359167	0.575227803	0.541846154	0.570798898	cg034446636	0.56263785	0.477538462	0.595592287	ENAH	1	63784592
cg24195486	0.575213393	0.539384615	0.563085399	cg02105641	0.562637	0.516	0.608815427	SULF1	8	70378740
cg15816012	0.575197288	0.548	0.537190083	cg02283643	0.56263022	0.649846154	0.516753444	MTRF1L	6	153324006
cg26596350	0.575175249	0.577538462	0.546556474	cg13538257	0.56261327	0.535076923	0.573002755	DHX16	16	30640512
cg19471466	0.575129477	0.664923077	0.420936639	cg05125483	0.56263089	0.515692308	0.573002755	ACCN1	17	31662900
cg03418847	0.575115067	0.556615385	0.566942149	cg03745383	0.56259868	0.470769231	0.531129477	ZNF345	19	37342585
cg06894708	0.575105743	0.555384615	0.587878788	cg16484811	0.56259868	0.587384615	0.531129477	LOC1001330	7	76178571
cg05663480	0.575089638	0.554769231	0.57630854	cg06911873	0.56258868	0.524615385	0.566391185	91		
cg06264150	0.575072685	0.578461538	0.554820937	cg02067211	0.56258699	0.561538462	0.548760331	TMEM30C	3	99903438
cg11429271	0.575068447	0.539692308	0.563085399	cg20843202	0.56257343	0.531692308	0.538292011	PAX4	7	127255970
cg1925132	0.575043865	0.557846154	0.511294766	cg18170071	0.56257173	0.407384615	0.608264463	PXMP4	20	32308480
cg03957109	0.575037932	0.537846154	0.450137741	cg04730850	0.5625658	0.548307692	0.553719008	1		
cg24214699	0.575032846	0.508615385	0.554269972	cg23842690	0.56255986	0.547076923	0.579063361	1		
cg16148593	0.575015046	0.548	0.577961433	cg17174814	0.56255478	0.508307692	0.548209366	15		
cg07838427	0.574984531	0.545230769	0.587327824	cg09299053	0.56254461	0.485230769	0.602703857	ZNF715	7	57509762
cg17250863	0.57496334	0.586769231	0.521212121	cg022447539	0.56253444	0.541846154	0.52892562	FSCN1	7	5633086
cg22503388	0.574949777	0.533538462	0.511294766	cg06142341	0.56250477	0.501230769	0.58953168	2		
cg14856738	0.574945539	0.507384615	0.57353719	cg03862437	0.56249705	0.531076923	0.581818182	TMEM44	3	194353432
cg04006010	0.574942149	0.554769231	0.563636364	cg27441048	0.56246662	0.501230769	0.602203857	BC17B	7	72972316
cg2063520	0.574900615	0.539384615	0.584022039	cg27303855	0.56246323	0.546615385	0.548709366	ZSCAN20	11	33937682
cg13980233	0.574892138	0.438769231	0.624242424	cg16827215	0.56245391	0.530461538	0.510192837	GALNT4	11	11292635
cg04146151	0.574777707	0.432307692	0.601101928	cg15099161	0.56244628	0.508	0.55922865	2		
cg10880514	0.574772621	0.640307692	0.423444353	cg18606843	0.56244543	0.508615385	0.574104663	7		
cg08173535	0.574767535	0.533846154	0.570247934	cg18649451	0.5624412	0.555384615	0.544903581	14		
cg25940354	0.574734478	0.601538462	0.505785124	cg15744869	0.56239712	0.516307692	0.56969697	ADAMTS13	15	94323963
cg01923692	0.574731935	0.564	0.525619835	cg13049458	0.56239288	0.515384615	0.58953168	TBC1D12	10	96219442
cg1526941	0.574728544	0.524	0.555371901	cg15975960	0.56238779	0.417230769	0.555371901	LTID1	1	62660624
cg19544445	0.574689553	0.489230769	0.566391185	cg13350140	0.56238017	0.516307692	0.569146006	CNN2	19	1035450
cg08373250	0.574689553	0.509230769	0.608815427	cg17311921	0.56236321	0.555692308	0.560881543	RET	10	43599566
cg00773093	0.574688705	0.500923077	0.608815427	cg11595964	0.56237084	0.524	0.57630854	CNGA3	2	98962564
cg22040889	0.574648019	0.587692308	0.506887052	cg17311921	0.56236321	0.555692308	0.560881543	UMHC1	5	176429459
cg15401952	0.57462259	0.571076923	0.514049587	cg16178091	0.56234965	0.493538462	0.578512397	1		
cg05384646	0.574621742	0.562153846	0.562088399	cg16439365	0.56233778	0.540615385	0.553168044	15		
cg03031672	0.574618351	0.468615385	0.635812672	cg02104456	0.56233355	0.516	0.542699725	BTBD1	16	96598888
cg11488033	0.574604789	0.54	0.575757576	cg23468699	0.56232507	0.492615385	0.572551791	MIR496	14	101526893
cg04714132	0.574582751	0.572923077	0.510743802	cg22021832	0.56232507	0.546769231	0.536088154	CYC117	19	42944717
cg27539482	0.574568341	0.509230769	0.6	cg06362358	0.56231998	0.446153846	0.637465565	1		
cg19402285	0.574523416	0.523384615	0.58292011	cg14266287	0.56231744	0.547692308	0.539393939	ASB1	2	239334506

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg14204124	0.574514092	0.609846154	0.547658402	FRMPD1	9	37651660	cg236550904	0.56230642	0.515076923	0.557024793	CNTD2	19	40730628
cg2376120	0.574502225	0.548923077	0.543250689	MEGF11	15	66331911	cg08578305	0.56229964	0.564	0.526170799	KRAS	12	25403937
cg18526696	0.574480186	0.501538462	0.595041322		12	64975934	cg23549869	0.56229625	0.547692308	0.547658402	HSD17B1	17	40706776
cg05207973	0.574450519	0.527169231	0.58980716	SLC34A1	5	176823642	cg13011928	0.56228862	0.517538462	0.578912397	IFAP2A	6	10599281
cg20673840	0.574447976	0.556307692	0.594490358	ESM1	5	54281362	cg05445108	0.56228269	0.523384615	0.584022039	RPH3A1	17	114319
cg10090591	0.574442043	0.585846154	0.541046832	CCDC84	11	118669922	cg13755538	0.56226055	0.501230769	0.578512397	CENPI	13	25496950
cg09953499	0.574388642	0.595692308	0.530027548	WBSR27	7	73264916	cg23501896	0.56225726	0.408307692	0.562344435		1	201976446
cg05065686	0.574338631	0.561846154	0.557575758	RAS10B	17	34068474	cg07193301	0.56224709	0.587076923	0.546556474	SNCAIP	5	121648993
cg27837940	0.574326764	0.540615385	0.570798898	FAM159B	5	63985374	cg08148553	0.562242	0.586461538	0.55927865	AP1M2	19	10698020
cg36363971	0.574298792	0.5416615385	0.560606061		4	145430689	cg035708536	0.56223946	0.468615385	0.602754821		1	113285118
cg08724891	0.57428523	0.532	0.519539229	CDK5R2	2	219826145	cg02182594	0.56223268	0.485338462	0.598347107	NTM	11	7810876
cg01537571	0.574277601	0.478769231	0.621487603	TM2D3	15	102193344	cg17519938	0.56223183	0.547076923	0.554269972	LRRC47	1	3713580
cg05616792	0.574266582	0.595076923	0.54600551	M8P	18	74844009	cg15934191	0.5622172	0.680307692	0.547451791		6	98298775
cg13110283	0.574255641	0.548615385	0.592837466	KCNK1	1	233751191	cg00324979	0.56221572	0.569358462	0.54214876		16	89152443
cg07275305	0.574255563	0.531384615	0.573002735	TRIM72	16	31227884	cg07856667	0.56220725	0.569358462	0.54214876		10	94455710
cg10715884	0.574252172	0.576384615	0.533884298	TMEM161B	5	87564247	cg23283423	0.56219677	0.547692308	0.550413223	RYN	17	821546
cg066663310	0.574211485	0.563076923	0.56804077		5	176153056	cg01865066	0.56217758	0.555692308	0.550964187		16	1178445
cg02800216	0.574211485	0.539692308	0.585674931		2	237542624	cg08077337	0.56217419	0.500923077	0.559779614	F2D7	2	20290063
cg18219563	0.574178428	0.523076923	0.564187328	SLC30A3	2	27487458	cg24777388	0.56216826	0.571076923	0.546556474	GYPC	2	127414883
cg22874255	0.57417758	0.461538462	0.621487603	ONCUT3	19	1768630	cg23213397	0.56216741	0.493307692	0.567493113	BDKRB2	14	96670738
cg16793394	0.574166561	0.430769231	0.631404959	OTX2	14	57278084	cg25883868	0.56215554	0.422769231	0.636363636	APOBEC3F	22	39436644
cg08950509	0.574158932	0.548615385	0.532782369	COL6A6	3	130368139	cg19305468	0.56214452	0.556307692	0.537474047	APC	19	1468359
cg175127779	0.57414876	0.556307692	0.563085399	CNO17	8	17104072	cg18176922	0.56213435	0.517230769	0.567493113	SIMO1	18	12420412
cg25440278	0.57414876	0.578769231	0.563085399	SNED1	2	241857607	cg21445204	0.56212672	0.60958462	0.474931129		2	220140750
cg24620761	0.574143675	0.498769231	0.58677686	ZIC4	3	147123199	cg06928484	0.56212418	0.449538462	0.58677686	VANGI2	1	160390288
cg18097224	0.574141979	0.555076923	0.56069697	TCF7	5	133451237	cg03354009	0.5621174	0.540307692	0.510743802	NUP107	12	59080458
cg05972503	0.574121636	0.52769231	0.583471074		15	89901456	cg06843899	0.56210468	0.531384615	0.526721763		5	42949542
cg06835814	0.574095359	0.493846154	0.595041322	KIAA1751	1	1905080	cg24410690	0.56209621	0.522769231	0.554269972	SLC15A4	12	129299481
cg00888521	0.574092816	0.625538462	0.469421488	RASGRF2	5	80256971	cg01513909	0.56209282	0.562769231	0.531680441	FAU	11	64889881
cg04640920	0.574088578	0.539692308	0.58677686		15	37180889	cg21964928	0.56208773	0.570461538	0.526721763		5	153872958
cg20442322	0.574081797	0.585846154	0.527823691	DNAH6	2	84743743	cg15375488	0.56208091	0.485846154	0.57630654		2	4050845
cg03606628	0.574068235	0.532923077	0.591184573	IL4I1	19	50393234	cg20544552	0.56205876	0.563692308	0.550964187		11	85593693
cg02978505	0.574020767	0.454153846	0.642424242		16	85292322	cg04366249	0.56205637	0.561846154	0.509641873	SGCE	7	94285501
cg01421830	0.573975842	0.539384615	0.586255895	KCNT1	9	138626955	cg24899205	0.56204959	0.523692308	0.53677686	TYNBD2	22	19929184
cg1437063	0.57396228	0.508	0.596694215	ADAR2	10	1715848	cg02458198	0.56201144	0.56307692	0.563636364	SPATA22	17	3375211
cg14802355	0.573947023	0.524	0.598347107	MANBA	4	103683140	cg27277583	0.56199873	0.501538462	0.586225895	CL50F27	15	76484625
cg03602360	0.573943632	0.531692308	0.580716253	KLHDC2	2	79664699	cg24335302	0.56199619	0.518153846	0.493112948	TRIM45	1	117664588
cg236635292	0.573940242	0.557538462	0.572066612		14	50234668	cg11182018	0.56198771	0.500307692	0.556473829	PCDH82	5	140474062
cg01504485	0.573930918	0.526153846	0.566947149		4	47843915	cg25694480	0.56198771	0.603846154	0.475482094		20	62032681
cg23205874	0.573927527	0.50384615	0.532231405	RGS19	20	62711523	cg10134527	0.56197923	0.525230769	0.576858504	ZBTB22	6	33283015
cg13749113	0.573922441	0.562769231	0.562534435		15	24506388	cg13655169	0.56197659	0.639692308	0.475482094		14	73929570
cg23036405	0.573911422	0.57230769	0.545454545		5	170763642	cg14179288	0.56197499	0.678461538	0.403856749	TLE4	9	82286351
cg2155471	0.573848697	0.547076923	0.552066116	DNAJC17	15	4106201	cg22082267	0.56197076	0.516923077	0.544903581	OR8K1	11	56112153

ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg26089165	0.573847001	0.634153846	0.460606061		2	23702420	cg19748485	0.56195635	0.483384615	0.620385675	PCDHGA4	5	140743927
cg11248869	0.573819877	0.493230769	0.602203857	FOXJ3	1	42801011	cg22933549	0.56195126	0.507076923	0.594490358		5	2372100
cg18566132	0.573811401	0.539384615	0.557024793		8	103613344	cg23507211	0.56192922	0.504538462	0.504683196	SEMA6A	5	115860190
cg16448525	0.573787667	0.625846154	0.51184573	FLJ42709	5	92918072	cg03526756	0.56192329	0.508615385	0.566942149		11	8698743
cg14191616	0.573786819	0.501538462	0.636914601	SH3RF3	2	109788051	cg10816323	0.56191651	0.531076923	0.55979614	PPP2R5C	14	102263580
cg07200616	0.573763933	0.454461538	0.655647383	BSGN1L1	17	80964885	cg137286491	0.56191481	0.477846154	0.580716253		10	134605694
cg06653293	0.573741894	0.504307692	0.563636364	CLT0rf91	17	1619494	cg15160514	0.56190803	0.515394615	0.562334435		12	32552765
cg05953287	0.573722151	0.545338462	0.561432507	TMEM48	1	54303980	cg03832227	0.56189277	0.516615385	0.550413223	ASTL	2	96798320
cg03616722	0.573706294	0.424615385	0.618732782		13	112602415	cg23397023	0.56189193	0.499384615	0.553719008		7	149112887
cg15253398	0.573697817	0.516	0.563636364	CACHD1	1	84936613	cg20089264	0.56188515	0.523384615	0.558677686	ASCC2	22	30234304
cg06581093	0.57368595	0.563384615	0.546556474		2	132181575	cg14571813	0.56187921	0.570461538	0.560330579		3	129062831
cg04728296	0.573643264	0.485846154	0.577410468	CPE	4	166414645	cg16991886	0.56187836	0.524307692	0.574855647		6	11832079
cg17631269	0.573638483	0.501230769	0.587878788	CHD5	1	6209018	cg24546155	0.56187582	0.578769231	0.501928375		18	44786426
cg13688770	0.573615597	0.530461538	0.575206612	ZFYVE28	4	2367137	cg16504661	0.56186648	0.562461538	0.55922865	CAPN14	2	31441554
cg04802236	0.573615597	0.573230769	0.557024793	RPL23	17	37010032	cg06048354	0.56186056	0.428307692	0.648484848	PCDH84	5	140501812
cg27359374	0.573606273	0.611076923	0.466115702		6	158375774	cg11339066	0.5618497	0.516	0.560881543	MIR596	8	1764618
cg14597321	0.573578605	0.501846154	0.611570248		6	13443660	cg02270518	0.56182561	0.484923077	0.58677686		6	29404728
cg04936446	0.573535071	0.423692308	0.571349862	RASSF8	12	26111348	cg05264472	0.56181903	0.517538462	0.54600551	SDHA	5	246469
cg11019923	0.573529965	0.539076923	0.574104683	DNAH4	15	78557532	cg02825985	0.56180208	0.547076923	0.561432507		22	49579410
cg10750464	0.573520661	0.508	0.592286503	CDC151	19	11534062	cg27096292	0.56179699	0.555076923	0.547107438	ANKRD11	16	89422212
cg13837721	0.573468451	0.515384615	0.643526171		11	116147580	cg24133569	0.56179445	0.58	0.526170799	BPGM	7	134330825
cg09361691	0.573478279	0.594769231	0.529476584	HTF2A	13	47471169	cg17531142	0.56176648	0.410153846	0.552066116	PVT1	8	128940078
cg21766898	0.573447764	0.579076923	0.548760351	TPA2B	6	50814089	cg16786640	0.56175122	0.507384615	0.581267218	DOK7	4	3485263
cg13499022	0.573408773	0.579692308	0.546556474	WDR85	9	140473606	cg29988655	0.56174613	0.433538462	0.38677686	HLA-H	16	29856070
cg0856204	0.573407078	0.492615385	0.561983471		17	35242555	cg23525438	0.56174529	0.468307692	0.60050964		6	17016326
cg01597066	0.573367239	0.539384615	0.573553719	SPHK2	19	49127506	cg25541928	0.56174444	0.515692308	0.573002755	SCG3	15	51973199
cg08227269	0.573342657	0.510153846	0.580716253	NEUROD1	2	182545555	cg09433929	0.56174105	0.501538462	0.598347107	CA3	8	36350913
cg17342807	0.573333079	0.483692308	0.603509642		4	379766	cg13223838	0.56173427	0.507692308	0.558677686	TBC1D22A	22	47370226
cg15148028	0.573318076	0.571076923	0.53353719	KIF76B	1	245755568	cg15738694	0.56173172	0.463692308	0.584573003	LOC144742	12	119740840
cg17191933	0.573308752	0.571692308	0.557575758	IPO5	13	98628505	cg19895882	0.56173172	0.478461538	0.555922865	KIAA0564	13	42534974
cg05282880	0.57329519	0.585230769	0.540495868	DIP2C	10	464225	cg00870753	0.56172579	0.554769231	0.549862259	UBE2MP1	16	34404772
cg19492498	0.573184997	0.583076923	0.471625344	MBL2	10	54531460	cg14646039	0.56170884	0.476923077	0.570798898		5	152726842
cg07365502	0.573147701	0.556615385	0.526170799		5	71852464	cg27119358	0.56168765	0.530461538	0.586225895	PRKD2	19	47217695
cg20291846	0.573133291	0.564615385	0.549311295	ADAM17	2	9696659	cg21341586	0.5616868	0.462153846	0.612121212	EIF4E	4	99851281
cg08289782	0.573095995	0.431692308	0.674380165	POLR3A	10	79789985	cg23453683	0.56167239	0.448307692	0.61046832	SECI	19	49168736
cg00563300	0.573095147	0.469846154	0.577410468	KCTD5	16	2747028	cg25506248	0.56167069	0.568923077	0.542699725		12	130502267
cg1346321	0.573041746	0.540307692	0.562534435	CUX2	12	111652316	cg25101405	0.56166645	0.501846154	0.564187328	ATAD3C	1	1398028
cg09169779	0.57302225	0.570153846	0.567493113		13	20751710	cg09699751	0.56165798	0.571076923	0.544903581	NCL	2	232320598
cg21018997	0.573012926	0.508923077	0.609366391		17	59572908	cg1975890	0.5616512	0.523076923	0.585874931	CTNND2	5	11903145
cg05563903	0.573007841	0.484307692	0.588429752	HQCA1	2	237415999	cg19786602	0.56164018	0.515692308	0.558677686		17	7966376
cg06673652	0.573006145	0.540615385	0.545454545		5	5672077	cg15819717	0.56163848	0.564307692	0.52892562	HQXC9	12	54393100
cg06617886	0.572999364	0.632	0.438567493	ZNF441	19	11877017	cg21176048	0.56161305	0.524	0.573553719	PEX5L	3	179755235
cg03366574	0.572994278	0.524307692	0.588429752		7	2764599	cg00566431	0.56159271	0.548615385	0.5184573	UBE2L6	11	57335606
cg04833819	0.572578173	0.430461538	0.592837466	ABR	17	927551	cg08111922	0.56159186	0.578153846	0.494714876	MGC23270	14	105287408

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ auc	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg15106537	0.572971392	0.46769231	0.619283747	cg20562678	0.56158847	0.563076923	0.558126722		13	28369080
cg04996251	0.572966306	0.547384615	0.558677686	cg09784259	0.56158508	0.547076923	0.566942149	SELE	1	169703422
cg17059564	0.572949354	0.524307692	0.555371901	cg11855834	0.56158169	0.50120769	0.553168044	IGSF3	1	111717224
cg12857867	0.572948506	0.547692308	0.579063361	cg15536663	0.56156389	0.570769231	0.541597796	EPH41.4A	5	111665548
cg15464382	0.572944268	0.54	0.575553719	cg26538035	0.56156135	0.518461538	0.553719008		1	231178722
cg22737347	0.572921382	0.536692308	0.578512397	cg22248789	0.56155965	0.516923077	0.600550964		2	63284574
cg06631222	0.572901038	0.492	0.580165289	cg01267532	0.56153931	0.515076923	0.568044077	MFSD6L	17	8702816
cg16161715	0.572858566	0.572307692	0.528374656	cg06557606	0.56153592	0.58520769	0.533884298	ATP6V1H	8	54756224
cg24833731	0.572801017	0.447076923	0.544903581	cg20971527	0.56153253	0.548307692	0.546556474	ENO1	1	8921294
cg11964358	0.572748464	0.487692308	0.601101928	cg23941942	0.56152575	0.547384615	0.576859504	NUP35	2	18398968
cg14360022	0.572734901	0.588923077	0.496969697	cg08017787	0.56153897	0.483846154	0.604958678	PIPRN2	7	158182836
cg03401726	0.572728968	0.572	0.538297011	cg05406923	0.56149354	0.476923077	0.585123967		6	99090481
cg21950155	0.572727273	0.467384615	0.669421488	cg18587364	0.56148893	0.517538462	0.591184573	ZNF562	19	9785779
cg09004398	0.572713711	0.584923077	0.557024793	cg118345456	0.56148506	0.542153846	0.515702479	PROX1	1	214175695
cg2146346	0.572709472	0.454461538	0.598347107	cg13354250	0.56148337	0.5	0.561983471		6	169558175
cg08335929	0.572689977	0.516615385	0.581818182	cg2138998	0.56148082	0.601230769	0.476033028	SFRS9	12	120903935
cg00436603	0.572684891	0.509230769	0.588429752	cg11341199	0.56147828	0.470153846	0.556473829		5	2643534
cg26749518	0.572682348	0.500307692	0.587327824	cg07782112	0.56145878	0.452923077	0.636363636		2	53107842
cg26184765	0.572677262	0.438153846	0.648484848	cg04727522	0.56145539	0.586769231	0.497520661	C18orf22	18	77793182
cg24133536	0.572652681	0.516923077	0.541597796	cg10474368	0.56144776	0.549230769	0.550964187	XRRAL	11	74660190
cg00074360	0.572639966	0.548538462	0.561983471	cg20295992	0.56142573	0.524307692	0.548708366		6	108440280
cg25809290	0.572635728	0.554769231	0.536693118	cg226441902	0.56141979	0.462153846	0.597245179	MYEOV2	2	241076601
cg11077770	0.572622166	0.576923077	0.557024793	cg02920148	0.5614181	0.500923077	0.54600351	YIF1A	11	66056751
cg09633595	0.572584022	0.531384615	0.564738292	cg02154765	0.56140538	0.586461538	0.48815427		8	1110111
cg19986012	0.572560288	0.555076923	0.560330579	cg23281527	0.56138504	0.609538462	0.498071625	KLHDC7A	1	18809531
cg26962439	0.572546726	0.406461538	0.665911846	cg05711145	0.56137148	0.484307692	0.615977961	ADAD2	16	84224633
cg16708264	0.572536554	0.547692308	0.55922865	cg04807864	0.56136893	0.531076923	0.529476584		2	130750050
cg19172487	0.572515363	0.563692308	0.555371901	cg07791427	0.56136554	0.533230769	0.569146006	HOXC8	12	54402704
cg07825681	0.572515363	0.500923077	0.519008264	cg26865747	0.56136046	0.539076923	0.571900826		6	28601377
cg17005446	0.572452638	0.531384615	0.563636364	cg00163372	0.56135368	0.562461538	0.558677686	MYC	8	128752988
cg19211853	0.572440771	0.548307692	0.582369146	cg20641660	0.56134266	0.461538462	0.367493113	C11orf30	11	76155487
cg04740229	0.572439076	0.591615385	0.556473829	cg08099141	0.56132401	0.548	0.529476584	C12orf57	12	7053170
cg08162948	0.572437381	0.546769231	0.519559229	cg22233844	0.56128756	0.564307692	0.539593939	ITGAE	17	3658293
cg10614445	0.572427209	0.585846154	0.554269972	cg00428487	0.56127654	0.542461538	0.548760331	SLC29A4	7	5323393
cg08922032	0.57240178	0.506153846	0.598898072	cg15482971	0.56126807	0.48520769	0.56969697	GABRB3	15	27019448
cg02333352	0.572396694	0.492307692	0.561983471	cg26086649	0.56123755	0.586461538	0.516253444		4	147556288
cg16801954	0.572395847	0.485538462	0.590082645	cg13331266	0.56123246	0.446461538	0.609366391	FLJ33360	5	6327070
cg21113773	0.572379741	0.593230769	0.548760331	cg00418613	0.561223162	0.516	0.552066116		6	20093995
cg26476755	0.572372113	0.555692308	0.562534435	cg26082202	0.5612189	0.548307692	0.575553719	C14orf118	14	76617583
cg14917512	0.572338207	0.623846154	0.474931129	cg21380560	0.56121212	0.516	0.555371901		6	41434320
cg20423484	0.572294978	0.563384615	0.567493113	cg23706819	0.56121127	0.484615385	0.570798898		4	89152212
cg14825735	0.572281416	0.555384615	0.555922865	cg24069376	0.56120788	0.555384615	0.576859504	EXOG	3	38537580
cg04704219	0.572254291	0.500615385	0.589531168	cg08075908	0.56120278	0.501230769	0.566391185		2	176997277

ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO	ILMINID	mean_cv_ auc	mean_cv_ sen	mean_cv_ spe	GENE	CHR	MAPINFO
cg13076847	0.572239881	0.492	0.508264463		4	3664401	cg13676220	0.56119347	0.557220769	0.539944904	C9orf171	9	135333078
cg06613286	0.572223471	0.587076923	0.521763085	TMEHM132C	12	128850903	cg19743077	0.56119263	0.515692308	0.568595041		1	244081054
cg23357643	0.5712213605	0.513846154	0.58292011	EP5813	1	110305591	cg10906729	0.56119178	0.546769231	0.500826446	LOC404266	17	46682390
cg0895932	0.572208519	0.433307692	0.611019284	LCF1C	1	152778380	cg19437757	0.56119178	0.469230769	0.583471074	PTRN2	7	157697057
cg02246609	0.572204281	0.456	0.605917355		6	28584003	cg17462074	0.56119008	0.416	0.61046832	GRIN28	12	14133168
cg19611416	0.572203433	0.533307692	0.549311295	STK38L	12	27397833	cg20681948	0.56118839	0.523076923	0.591735537		12	103342297
cg1959457	0.57220089	0.563692308	0.585123967	NUF2	1	163290999	cg08198377	0.56118839	0.563394615	0.57024793	TMX1	14	51707975
cg00837838	0.572182242	0.523692308	0.554820937	B2M	15	45003774	cg13173903	0.56118754	0.532	0.57630854	RPN2	20	35807784
cg230333870	0.57215427	0.547692308	0.560330579	TMPRSS5	11	11357528	cg17434690	0.56118754	0.516307692	0.558126722	MYO15B	17	73613031
cg08952028	0.572132231	0.500615385	0.571349862	CHRD12	11	74442900	cg2226227	0.561185	0.536307692	0.563085399		4	3726292
cg08102516	0.572072049	0.538461538	0.553371901	SOK1	7	3411356	cg24754692	0.56117906	0.554769231	0.553922865	PPP3CA	4	102246975
cg13838861	0.572066963	0.608923077	0.507988981		22	20343300	cg20744163	0.56117822	0.522461538	0.571900826	PCNXL3	11	65401703
cg16017089	0.572061878	0.532	0.550413223	ARHGEF17	11	73053830	cg20744163	0.56117652	0.444615385	0.596143251	ZMIZ1	10	80999841
cg10678427	0.572040687	0.484307692	0.599449036		3	127176163	cg22496683	0.56116635	0.561230769	0.550413223	RBM46	4	155702610
cg25114504	0.571996609	0.539692308	0.566391185	RSPO2	8	109092720	cg25132161	0.56116635	0.50769231	0.520661157	USP13	3	179370779
cg16785447	0.571956771	0.617846154	0.4815427	DAD1	14	23058779	cg02743029	0.56115957	0.563692308	0.534435262	SERTAD4	1	210405212
cg02829845	0.571948294	0.484307692	0.555922865		4	188963260	cg20797905	0.56115787	0.556	0.574104683	RPTOR	17	78747227
cg06809298	0.571936427	0.516923077	0.557024793	COX7A2	6	75953833	cg02508421	0.56115787	0.570153846	0.555922865		3	49145923
cg01302668	0.571909303	0.524615385	0.554820937	FAM162B	6	117087238	cg14953804	0.56114601	0.437230769	0.543250689	DIP2B	12	51021072
cg19770292	0.571908455	0.578769231	0.561432507		5	1868593	cg13147090	0.56111634	0.415076923	0.620855675	HQSEC3	12	247963
cg06900062	0.571896588	0.548	0.580165289		6	27236425	cg24530489	0.56111295	0.585846154	0.567493113		15	56299380
cg10024000	0.571872854	0.501230769	0.584573003	NF1	17	29422363	cg13608545	0.5610926	0.539076923	0.572451791		22	43166139
cg00532020	0.571855054	0.471384615	0.634710744		5	180511822	cg08316083	0.56108413	0.493615385	0.557575738	ATP8B3	19	1796207
cg01065905	0.571850816	0.532	0.54600551	SIC30A7	1	101416548	cg18619880	0.5610746	0.546153846	0.558677686		8	92828658
cg04771433	0.571849121	0.423384615	0.649586777	PRKCZ	1	2064716	cg01022370	0.56107311	0.568	0.536639118	MFHAS1	8	8743074
cg24541425	0.571844882	0.501230769	0.538292011		12	130529058	cg04017458	0.56107226	0.524923077	0.521765085		17	41438330
cg25749040	0.571836406	0.588307692	0.53553719	KLK1	19	51323301	cg19741148	0.56107141	0.595692308	0.51184573		5	553895
cg09171874	0.571819453	0.523692308	0.566942149	S100A3	1	153520801	cg08535018	0.56105785	0.540615385	0.56969697	NCN18	8	12272982
cg00981737	0.571807586	0.416	0.660045096		14	105103747	cg04027548	0.56105192	0.594461538	0.478236915	NCN12	10	126138800
cg14007688	0.571773767	0.594461538	0.497520661	DBH	9	136529227	cg02698338	0.56105022	0.500307692	0.534986226	NCN12	10	126138800
cg16184827	0.571773681	0.470461538	0.61267176	CNGA3	2	98961824	cg27034150	0.56104937	0.564	0.528374656	SUR1A1	16	28635053
cg76066440	0.571771986	0.593538462	0.543801653	PLIN3	19	4868007	cg01194279	0.56103751	0.479384615	0.555922865		12	114073142
cg13590414	0.57177029	0.446153846	0.658402204	RPUS03	3	9885815	cg18982286	0.56102564	0.540615385	0.561432507	SHANK3	22	51336325
cg10659624	0.571758423	0.554461538	0.566942149		9	128173245	cg00653081	0.56099936	0.540307692	0.553719008	TAOK1	17	27717927
cg03832220	0.57175249	0.531384615	0.56969697	LOC284100	17	36244714	cg03658275	0.56097648	0.54	0.579063361	LAMA2	16	129285274
cg04857710	0.571737232	0.632615385	0.525619835		4	186401692	cg08841613	0.56097224	0.507692308	0.544903581		2	220553978
cg27505178	0.571730451	0.554153846	0.565289256	LOC779375	3	129830468	cg06627617	0.56096631	0.586769231	0.5184573	ASAP2	2	9471179
cg20784768	0.571653316	0.523384615	0.562085399	SIC35F3	1	234041456	cg224929017	0.5609519	0.665230769	0.429752066	C2orf42	2	233734820
cg14434870	0.571641449	0.522153846	0.593939394		6	99279054	cg22694560	0.56094681	0.493230769	0.566942149	ADCV3	2	25135724
cg0473621	0.571639754	0.539076923	0.552066116	HSD17B4	5	118788720	cg01833485	0.56092732	0.531692308	0.55922865	ESRRG	1	216860692
cg2734137	0.571631278	0.548307692	0.563085399	NR2F2	15	96858114	cg04936149	0.56091375	0.517230769	0.548209366	HBA2	16	221811
cg02239021	0.571588986	0.533846154	0.56804077	CYBSRL	1	54665433	cg03693386	0.56091375	0.554769231	0.527823691	CNR1	6	98875236
cg07601542	0.571563162	0.548615385	0.548760331	OLIG2	21	34396986	cg02558625	0.56088154	0.515384615	0.543801653	COL4A1	13	110951418

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	GENE	CHR	MAPINFO
cg21808131	0.571562619	0.547384615	0.566942149	ARHGAP5	14	32597140	cg07370905	0.5608773	0.607692308	0.465564738	WDR52	3	113015910
cg11493544	0.571561772	0.562461538	0.560881543	GRB2	17	734012155	cg24586196	0.56086459	0.571384615	0.520110193	MAEL	1	166958406
cg10703784	0.571557533	0.588	0.526721763		12	28127353	cg06683241	0.560862341	0.564615385	0.52892562	HNPPSA	10	134351765
cg14242557	0.571556686	0.460307692	0.564187328	ZNF483	9	114288341	cg17469361	0.56085781	0.501846154	0.503030303	CHAT	4	6536474
cg05077167	0.571555838	0.618461538	0.547658402	ZFP3612	2	43453540	cg25609507	0.56085781	0.535384615	0.513498623	C2orf54	10	50821597
cg00627998	0.571545666	0.571692308	0.571900825	CRTAC1	10	98686032	cg01538581	0.56082556	0.640307692	0.4445179063	C2orf54	2	241832900
cg04733156	0.571541428	0.578153846	0.522865014	LOC654342	2	91847990	cg27165836	0.56081204	0.593538462	0.52221405		1	4193882
cg06766189	0.571488027	0.531692308	0.585123987	NGF	2	233790293	cg01274907	0.56081119	0.514769231	0.560330379	PCDH9	5	140568075
cg18488996	0.571478703	0.523692308	0.56584022	PAR5	15	25230295	cg09461388	0.56080695	0.478461538	0.59949036	CELSR1	22	46763229
cg07009284	0.571467684	0.531692308	0.568044077		8	76318515	cg17061192	0.56080102	0.501230769	0.55757578		4	185926620
cg01115070	0.571449036	0.586307692	0.459504132		11	43969000	cg01156070	0.56079339	0.539076923	0.574104683		6	100064885
cg10797813	0.571445645	0.554461538	0.570247934	PHF1	6	33377848	cg02805106	0.5607883	0.564	0.548760331	RHCG	15	90039409
cg03547722	0.571431235	0.555384615	0.573002755	NURP1	16	10837705	cg25587159	0.56078745	0.507384615	0.591735537		3	197836971
cg15825321	0.571428693	0.54	0.571900826	ARHGAP23	17	36666642	cg13848773	0.56078067	0.522769231	0.5592865	HERC4	10	69835125
cg10144483	0.571422759	0.587692308	0.499173554		4	180981242	cg19595234	0.56077722	0.522769231	0.514049587	TOX2	20	42543778
cg16222698	0.571419369	0.516923077	0.566391185	REXO2	11	114310578	cg01907820	0.56076626	0.462769231	0.602703857	SCNN1A	12	6473007
cg18222083	0.571413435	0.523076923	0.566942149	TNFRM106A	17	41364007	cg22908423	0.56074423	0.532307692	0.560881543	C11orf9	11	101947241
cg24509168	0.571410045	0.538153846	0.546556474	KCNJ9	1	160051356	cg01060910	0.56074423	0.486153846	0.573553719	LANCE1	2	211341456
cg19096571	0.571410045	0.556615385	0.550413223	GSTO2	10	106035131	cg07614486	0.56073914	0.525230769	0.55757578	BARAA	1	229406711
cg10074506	0.571404959	0.593538462	0.558677686	SEZ6	17	27333617	cg16908962	0.56073236	0.555692308	0.558126722	ZAK	2	174132279
cg18048405	0.571396482	0.439692308	0.626446281	SA536	1	100598378	cg10149837	0.56073151	0.531384615	0.585123967	LOC1001296	16	87738623
cg23106854	0.571392244	0.586153846	0.557024793	WDR88	19	33623072	cg02402630	0.56073066	0.515384615	0.552066116		7	155996361
cg18867912	0.571382072	0.432923077	0.639118457		3	193692556	cg09461420	0.56072643	0.515076923	0.562534435	CBFA2T3	16	88994295
cg22150680	0.571381225	0.448	0.574104683	SORCS2	4	21934487	cg03704006	0.56070778	0.410769231	0.659504132	SDCCAG1	14	50320632
cg01166759	0.571354348	0.515384615	0.547658402	KRTAP3-1	17	39165697	cg19509763	0.56069591	0.462153846	0.607713499		1	4087676
cg26038749	0.571349015	0.532923077	0.575757576	SVIL	10	29759454	cg14277848	0.56068489	0.561230769	0.546556474	GPATCH4	1	156570825
cg14957731	0.571319347	0.485846154	0.568595041		1	170899285	cg20458353	0.56067302	0.484615385	0.55979614		7	21260793
cg14726783	0.571317552	0.547384615	0.564738292	DCC	18	128080305	cg18179997	0.56067133	0.532	0.573553719	SKIV2L	6	31930875
cg24368876	0.571315109	0.510769231	0.582369146	FAM150A	8	53478706	cg14312578	0.56066879	0.541230769	0.538292011	HAS1	19	52217388
cg06429194	0.571312566	0.579692308	0.528374656	KLF14	7	130418324	cg17320856	0.56066709	0.508615385	0.554268972	GAINT9	12	132687617
cg16964748	0.571304937	0.563692308	0.527272727	C16orf62	16	19566559	cg05524450	0.56066455	0.525846154	0.536088154	C19orf43	19	12845012
cg16021627	0.571304937	0.564307692	0.522865014	NEU1	6	31829976	cg07587706	0.56065353	0.53230769	0.590633609	MCCC1	3	182816873
cg20426994	0.571277813	0.523384615	0.553719008	IGM2	6	10881782	cg07634101	0.56063759	0.548	0.546556474		11	366375
cg07210205	0.571266794	0.530461538	0.56969697	CCDC108	2	2129351	cg16736080	0.56063149	0.524615385	0.532314405	FAM181A	14	94393497
cg18187244	0.571259165	0.539692308	0.574104683	SNRG2	2	1229351	cg25511237	0.56061454	0.500307692	0.561432507	ADH5	4	100010080
cg21084508	0.571243908	0.532923077	0.56969697	BLCAP	20	36156231	cg25073725	0.56061284	0.500307692	0.564758292		20	57978889
cg14160807	0.571242212	0.563076923	0.57658402	SPAG1	8	101177733	cg20347090	0.56060382	0.524615385	0.551515152	PRMT8	12	3607093
cg00191853	0.571210002	0.609538462	0.525619835	F5	1	169555944	cg07978099	0.56060013	0.58461538	0.557024793	OSGIN1	16	83986941
cg25256723	0.571202373	0.548307692	0.585123967		3	143993048	cg25766748	0.56059928	0.501846154	0.558677686	CACNA1A	19	13614882
cg20566934	0.571190506	0.569538462	0.55979614	GAINT9	12	132892417	cg06874323	0.5605908	0.524	0.549862259		10	102986525
cg26076948	0.571160839	0.477538462	0.627038567										

ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_auc	ILMINID	mean_cv_auc	mean_cv_sen	mean_cv_auc	CHR	MAPINFO	GENE	CHR	MAPINFO
cg04373285	0.571159992	0.561846154	0.575757576	cg00781405	0.560580663	0.483384615	0.559779614	19	34404619	ZNF611	19	53207899
cg22607959	0.57113541	0.584615385	0.498071625	cg19430677	0.56056707	0.423076923	0.551515152	10	54603444	PAOX	10	135192755
cg09210519	0.571131172	0.507076923	0.569146006	cg26789064	0.56055435	0.587692308	0.534435262	4	106335252		4	4866130
cg05723527	0.571118457	0.594461538	0.494214876	cg0703610	0.5603062	0.594153846	0.523415978	17	45405871	SMARCD2	17	61919098
cg16781264	0.571115914	0.438769231	0.576859504	cg10662479	0.56052254	0.465846154	0.591735537	20	101087575	WFD3	20	44420225
cg24141298	0.571096419	0.586153846	0.523966942	cg05956462	0.56052384	0.516923077	0.580165289	3	74778002		3	127175400
cg27371953	0.571087095	0.595384615	0.543801653	cg2088263	0.56052299	0.572615385	0.499724518	8	28897999	GRAM2	8	51746830
cg11293176	0.571066751	0.587076923	0.505785124	cg08242000	0.56051357	0.407076923	0.568044077	16	242714091	NRP2	16	144922888
cg17127769	0.571065056	0.549846154	0.553168044	cg27013478	0.56050265	0.524307692	0.55972865	5	70178002	FLYWCH1	5	2961939
cg05056748	0.571057427	0.584615385	0.499173554	cg03877420	0.5605018	0.477538462	0.574104683	7	169724590		7	1004367
cg14903689	0.571040475	0.539076923	0.525068871	cg05067136	0.56049826	0.523384615	0.539359393	13	235903542	C7orf30	13	23338602
cg12557350	0.571030303	0.524307692	0.567493113	cg01870826	0.56049826	0.616923077	0.512396694	7	46875218	LOC389458	7	5111875
cg15509177	0.571012503	0.562769231	0.553719008	cg04003871	0.56049841	0.5	0.570474934	2	126432983		2	241290895
				cg15575375	0.56048909	0.508923077	0.606060606	1	19919208	NFIA	1	61649973
cg22027267	0.570987074	0.618461538	0.528374656	cg17490089	0.56048315	0.516307692	0.56584022	19	51544518	KANK3	19	8398300
cg04804717	0.570985378	0.453846154	0.646780992	cg19815327	0.56047552	0.493538462	0.561983471	3	1300320	ZNF860	3	32023353
cg27268110	0.570977775	0.625846154	0.489807163	cg05848389	0.56046874	0.508307692	0.557206612	14	82127764	DYNCH1	14	102508999
cg02508881	0.570971816	0.563076923	0.526170799	cg01248385	0.56046281	0.555384615	0.531129477	13	142216119	MCF2L	13	113636953
cg26280827	0.570970121	0.4141153846	0.615977961	cg05313771	0.56044755	0.596	0.436914601	19	49000996	ONECUT3	19	1762435
cg05064106	0.570965095	0.525538462	0.571900826	cg02213971	0.56041958	0.493846154	0.602754821	11	50528213	ANO1	11	69994230
cg01560685	0.570904853	0.578769231	0.551515152	cg16978004	0.56038313	0.540307692	0.573553719	15	32322028	SNORD116-17	15	25330742
cg17436861	0.570897224	0.523384615	0.58677686	cg05460986	0.56038228	0.525230769	0.493112948	6	176900842	RNASET2	6	167370529
cg244449136	0.570876033	0.524923077	0.575553719	cg01977082	0.5603772	0.462461538	0.555371901	8	30128219	COX6C	8	100906160
cg15154047	0.570874338	0.569846154	0.590533609	cg19442545	0.56037635	0.547692308	0.562334435	10	143210309	FUT11	10	75533431
cg03634397	0.570871795	0.500923077	0.5553719008	cg14115163	0.56037211	0.540307692	0.529476584	10	91147145	HERC4	10	69835215
cg05668580	0.570862471	0.571384615	0.527272727	cg14470803	0.56037211	0.564615385	0.467217631	9	161494307	DBC1	9	122132379
cg24618492	0.570860776	0.493230769	0.602203857	cg01135950	0.56037127	0.548923077	0.530578512	5	19140922	SLC6A3	5	1443198
cg18736800	0.570850604	0.523384615	0.544903591	cg18864926	0.56036448	0.563076923	0.576859504	14	24914284	CCDC85C	14	99989889
cg22760475	0.570837042	0.563076923	0.561983471	cg2876561	0.56036279	0.515384615	0.555371901	7	149380129	PHF14	7	11013721
cg21548749	0.57082687	0.516	0.6	cg00214530	0.56035346	0.570769231	0.51184573	6	32096524		6	133890532
cg01230067	0.570793812	0.585230769	0.527823691	cg00352468	0.56034668	0.438461538	0.619834711	2	7493222	EEF1E1	2	10687659
cg13336665	0.570789726	0.547384615	0.551515152	cg21717369	0.56034329	0.523692308	0.566391185	6	134872610	ICDRLG	6	8103092
cg03427058	0.570787031	0.431384615	0.571349862	cg17249942	0.56033397	0.470153846	0.616753444	9	1565025	COG5	9	140043007
cg05337441	0.570771774	0.492615385	0.582369146	cg19565406	0.56033227	0.54	0.547933884	7	7493222	GRIN1	7	107205105
cg07750655	0.570769231	0.491692308	0.563085399	cg14896906	0.56033143	0.461846154	0.62369146	6	21266568		6	168167789
cg0699008	0.570753669	0.478461538	0.79614325	cg18054526	0.56032634	0.541230769	0.556473829	4	51213855	PH4	4	74848016
cg14242735	0.570753973	0.438153846	0.570798898	cg13126871	0.56031617	0.515692308	0.545931295	3	34258406		3	67019880
cg04793390	0.57073363	0.564	0.555922865	cg03655504	0.56031617	0.537	0.560881543	18	24288969		18	72837386
cg00105470	0.570722611	0.524615385	0.55261708	cg21412355	0.56031193	0.460615385	0.6	1	154332333	PM20D1	1	205819251
cg15838301	0.57071922	0.579076923	0.532231405	cg14893161	0.56031193	0.424615385	0.552066116	1	7719899		1	15930761

85

GENE	GENE	GENE	GENE
TXNIP	KIAA0391	C5orf45	PCDHB5
NUDT1	VPS37C	MIR517A	ZNF432
GALNT10	DCAKD	TRPA1	WDR73
BRD2	PAK6	NHSL1	NSD1
MAP2K1	NMBR	TOLLIP	PTPRT
LRRC30	C20orf94	INA	ZSCAN16
C6orf48	FNDC1	MLPH	ALAS1
ABCA13	KRTAP4-5	FRS3	KRT83
HEATR1	ZNF277	PNRC1	CACNA2D4
ZNF267	ELP2	MTHFSD	SOX17
TDRD12	ST6GALNAC5	KCNG2	OR1F1
PAM	PIGY	C1orf61	CYP27B1
IKZF3	LOC255512	UMPS	HOXB3
RNF145	TBC1D1	SPRN	TFIP11
TRIP10	BTBD11	KCNAB1	B3GAT1
ADAMTS2	OPCML	IGDCC3	CPLX1
ZNF844	ZNF443	KCTD13	A2BP1
C14orf80	MTFMT	TREH	NAV2
RPGRIP1L	BAT4	KIAA1530	C1orf84
HDAC4	DLGAP2	RPA1	BEST1
NCOR2	FLJ12825	CYP2W1	CDC34
C7orf50	SLC25A12	RANGAP1	SBNO2
LRRC10B	NRXN2	NAV3	MTNR1A
SLC7A11	TRAPPC10	DCP1A	PLEC1
ATXN1	TLX3	ACACA	EFNB3
KLF11	APLNR	DMXL2	CHTA
HLA-E	TCEB1	C11orf73	FLJ36000
HLA-A	SLC16A12	LHPP	SULT2B1
CNTNAP5	EPHA10	ATAD3B	DAXX
TXNDC6	RNF219	AES	GPR4
DNMT3A	HS3ST2	LOH12CR1	RGS6
LOC389033	DAPK1	TCTE1	SNTG1
MESDC1	HERC3	SNRNP27	CRTC1
TTC30A	LFNG	SPDYE1	GRIK3
EIF3H	FGF14	SERINC2	TARBP2
KIRREL	GDF10	PRMT6	COL5A1
SUGT1L1	HLA-F	APCDD1L	DZIP1
RPL31	AHCY	CCT3	N4BP2
CBR1	GLB1L2	MTUS2	OCIAD1
TACC1	CLK1	SYT9	HPR
FAM38A	TXLNA	C17orf61	YJEFN3
PAX7	LOC146481	DGKG	ZNF680
ATHL1	CYYR1	RIPK4	TNNT3

GENE	GENE	GENE	GENE
NF2	NEUROG3	SLCO6A1	NMB
ASCL2	ZNF705D	CCNT2	LAMB3
CYP2J2	SYNPR	SEC22C	CCDC67
TRIML1	HNF1B	DPP9	SLN
RGPD2	CNTFR	TOP2B	FAM168A
DACT2	NEURL3	MARK2	TUBE1
MIR596	C19orf26	COL1A2	ANGPT1
CHRNA2	LMLN	ZNF512	HLA-C
HRNBP3	TECPR2	ASB18	OR11L1
RAI1	LOC387763	MAGI1	ZNF174
EBF3	SHISA4	ST7OT4	ZNF274
GRAMD2	SFTA3	HLA-B	ZNF639
MAD1L1	SNORA76	AKR1CL1	ENC1
IPO9	ZNF253	C1D	ZNF75A
FAM101B	AFG3L1	MED13L	CHGA
CACNA1B	ZFP14	C13orf27	KREMEN2
KY	CACNB4	OSR1	AK3L1
ASTN1	LCE3A	RB1	INS-IGF2
RBM47	PCM1	ATPAF1	ATP6V0A1
SLC32A1	CASR	LOC441601	ASZ1
TLE3	ZC3H12D	COL9A2	STEAP3
CTBP2	AHRR	INPP5A	CDH13
FXYD5	WNT9B	IQGAP1	SHISA2
MUC2	HSD17B7P2	KATNAL2	COL6A6
OSBPL5	SLC5A1	ARSA	FECH
PTPRN2	LRP5	PRKAR1B	SF3B14
C8orf38	OR5D16	OTX2OS1	ESPNL
ARHGEF10	ZFAT	TBX3	KCNS3
ZNF516	SLFN12L	CCDC144B	MOBP
TNXB	CD163L1	PLBD2	MRPL23
VASN	KL	MCPH1	BOK
ZNRF1	EVX1	TFPI2	INPP5B
ZNF776	DPPA5	FAM111A	PTPRG
GNG7	KCNF1	ZFP37	BICC1
TBX4	TPO	KHSRP	WDR66
F2R	CACNA1H	RNF212	C2orf3
C10orf119	ZNF544	IGF2BP3	PLA2G5
SLIT2	PPP1R13L	NIPA2	ZNF593
TNFRSF8	KLHL14	DNAJC18	TMEM192
PHACTR2	ACCN4	DTNA	SLC25A39
CAMTA1	KLK8	IP6K1	GSG1L
NDRG4	KCNA6	AGER	RASGRP3
GRIN2D	GPN1	BBS1	GFPT2

GENE	GENE	GENE	GENE
RFC1	GPR83	FAM160A1	UBE2QL1
ODZ4	CPN1	KCNK4	FAM96B
SEMA3D	SOS2	CSMD1	TPM3
TTC12	BRUNOL4	KLHL28	GBE1
ADAM12	EIF2C2	RGS12	PRDM13
CNNM2	PHF15	TRIM61	NEUROD1
MAP3K14	PPP4R2	ADAM32	MAP2K3
KCNMA1	FDPS	INTS2	LOC255167
NLRP2	C15orf61	ECE2	NGEF
PPP2R2C	ZNF233	CYB5B	KCNA7
KIAA1310	ARFGAP1	CDC25B	C9orf24
PRKAG2	PCDHGA2	ZNF32	METTL13
PIP4K2A	C3orf59	MAP1B	CYP2E1
CST9	PLCE1	TMEM161A	SMOC2
KCNAB2	PCDHB14	TEX2	CSTL1
LILRA4	TMEM132D	MRPL33	HEXIM1
PLEKHA7	SHANK2	NRF1	RNASEH1
ASCL4	HECW1	FGFR3	C2orf69
ANK2	ZBTB12	PTPN14	WT1
RPL38	LOC645323	CAMKK1	ZNF366
MYT1L	DPP6	AP1G1	ANK1
C1orf96	MXRA8	PITX3	PLSCR2
LMO3	ENAH	CNTNAP3	IL17RE
THRA	SULF1	ACTBL2	PRKCD8P
ZNF598	MTRF1L	PDHX	LOR
SOX2OT	ACCN1	ZNF605	SLC6A18
TRIM4	ZNF345	CLIC6	FAM134B
SLC7A6	LOC100133091	HS3ST6	SFN
RHOA	TMEM30C	FBXO15	GCKR
KIF26A	PAX4	SYTL3	HHLA2
LUZP1	PXMP4	PBX1	DUSP19
PHF20	ZNF716	HDC	KCNK7
PRR16	FSCN1	AFAP1	HPCAL4
TXNL4A	TMEM44	FOXK1	NPB
CCDC85A	BCL7B	NKX2-4	PRDM16
C6orf89	ZSCAN20	NIPAL3	SGCE
CCDC64B	LOC100287834	PKNOX2	GPR88
MED18	ADAMTSL3	TMED10	C13orf16
ESRP1	TBC1D12	NR2F6	VWDE
CARS2	L1TD1	MLEC	AMOTL1
SHROOM3	CNN2	BCKDK	LPHN2
DNAJB5	RET	ESYT2	ELOVL4
GOLSYN	UIMC1	SLC11A2	DKFZp761E198

GENE		GENE		GENE		GENE
SPSB4		BTBD1		SP6		MAP3K11
C3orf21		MIR496		TMEM150C		CABYR
SIAE		CXCL17		ARID3C		ID3
ODF2L		ASB1		SGCD		SFMBT1
SFXN4		CNTD2		LASP1		ADAP1
IQSEC1		KRAS		PKN1		KLC3
CHD5		HSD17B1		PFKL		ACVR1C
TYW3		RPH3AL		RNF186		RUNX2
C11orf63		CENPJ		CRCT1		FGF12
KCNT1		SNCAIP		FZD2		ZNF284
C17orf51		AP1M2		NEURL1B		P2RX7
DSCAML1		NTM		SNTG2		HNF4A
FLJ39582		LRRC47		BLCAP		SEZ6L2
UPF2		FZD7		SPAG1		SNED1
KCNK10		GYPC		F5		SFRP1
C21orf70		BDKRB2		GALNT9		CDKN2A
MYOM3		APOBEC3F		UBE2MP1		NKAIN3
RNF5P1		APC2		DHX29		CCDC102B
PAPPA2		SLMO1		SH3PXD2A		TSR1
QRICH1		VANGL2		DUOX2		CYP1A1
FAM50B		NUP107		NMS		PSTK
MED1		SLC15A4		COL4A3BP		NCRNA00164
VGLL2		FAU		TRIM27		RAET1G
CENPO		TXNRD2		WNT5A		PPP1R3B
TEX101		SPATA22		LCP2		KCNQ1OT1
SPIRE2		C15orf27		SH3BP4		SLC24A4
DDAH2		TRIM45		FAM53B		LILRB3
LOC388965		PCDHB2		LOC100101938		FBXL2
FAM193B		ZBTB22		C10orf58		COG5
HIP1		TLE4		LMTK3		NFASC
COMMD1		OR8K1		MOV10L1		PCBD2
HLA-DQB1		SEMA6A		DBN1		C15orf48
FREQ		PPP2R5C		TRIM10		ANXA2
NALCN		ASTL		GABRG2		LRRC27
LYPD5		ASCC2		EPN2		PHYHIP
FAM114A2		CAPN14		HMGXB3		KRI1
DLC1		PCDHB4		ATF6B		GGT7
LIMS2		SDHA		NEUROG1		PDZRN4
DHX16		ANKRD11		MIB2		IQSEC3
VWA3B		BPGM		SOX15		SHPK
GLRA3		PVT1		APOB		GRIK2
COL9A3		DOK7		ALG1		EXOC2
RNPS1		HLA-H		C1orf35		OCA2

GENE	GENE	GENE	GENE
KLK6	SCG3	DKFZp686A1627	LOC286094
TMT3C3	CA3	LOXL2	PKD1
AXIN2	TBC1D22A	LOC399959	LGALS13
THEM4	LOC144742	IRF6	NR6A1
TRIM31	KIAA0564	NMUR2	TLR10
SLC30A3	PRKD2	BTN3A2	C1orf170
SKI	EIF4E	SIM2	ZC3H12A
C21orf29	SEC1	UGGT1	ESRRA
RFXP3	ATAD3C	APEX1	KCNIP4
ALX4	NCL	DEFB116	CDH4
COL18A1	CTNND2	PSMD14	FRMPD1
RAB27B	HOXC9	MEST	MEGF11
LOC440356	PEX5L	UQCRC1	SLC34A1
COL13A1	UBE2L6	MIR548F5	ESM1
RALGDS	MGC23270	AK5	CCDC84
GALNTL4	SELE	NCOA7	WBSCR27
CDC20B	IGSF3	CCDC152	RASL10B
LOC84856	MFSD6L	CASC3	FAM159B
FOSB	ATP6V1H	ITPR2	CDK5R2
ADARB2	ENO1	AGK	TM2D3
CSDA	NUP35	C12orf39	MBP
CCDC69	ZNF562	TTLL10	KCNK1
ARHGEF4	PROX1	FAM57A	TRIM72
MCF2L	SFRS9	VIPR2	TMEM161B
GRM4	C18orf22	NELL2	ONECUT3
JPH3	XRRA1	LOC150622	OTX2
B4GALNT3	MYEOV2	FKBP10	CNOT7
OTUB2	YIF1A	PRKAA2	ZIC4
IFT172	KLHDC7A	TMC5	TCF7
COASY	ADAD2	TCF21	KIAA1751
LOC494141	HOXC8	ACOXL	RASGRF2
ZNF518B	MYC	C19orf63	DNAH6
COL29A1	C11orf30	FLJ43860	IL4I1
ODZ2	C12orf57	ADRB3	MANBA
PTPRN	ITGAE	FAM19A3	KLHDC2
COX16	GABRB3	SLC25A22	RG519
KIF26B	FLJ33360	GRASP	DNAJC17
HEBP2	C14orf118	EEFSEC	FOXJ3
BAT1	EXOG	TFAP2D	FLJ42709
HCN4	C9orf171	NR5A2	SH3RF3
CXCR7	LOC404266	PCOLCE2	B3GNTL1
DNAH17	GRIN2B	TACC2	C17orf91
BAIAP3	TMX1	NOS1	TMEM48

GENE	GENE	GENE	GENE
CALHM2	RPN2	FGFR2	CACHD1
OR2Z1	MYO15B	NFAT5	CPE
TUBB4Q	GPR133	PEBP4	ZFYVE28
DTHD1	PPP3CA	FKBP4	RPL23
ZNHIT1	PCNXL3	HTR6	RASSF8
MYO7A	ZMIZ1	DSCR3	DNAJA4
AUTS2	RBM46	SYCP1	CCDC151
MYRIP	SERTAD4	HOXA6	HTR2A
LOC100302652	RPTOR	HCN2	TFAP2B
IGSF9B	DIP2B	C2orf89	WDR85
NEDD4L	ATP8B3	TRPM1	SPHK2
CDK5	MFHAS1	RB1CC1	IPO5
ZNF577	KCNJ8	UNC5A	MBL2
PCDHGA1	NKX1-2	ANKRD43	ADAM17
PCDHB13	SULT1A1	SAR1B	POLR3A
HCCA2	SHANK3	LOC440173	KCTD5
CUL3	TAOK1	EBF2	CUX2
PCDHA6	LAMA2	PRICKLE4	IQCA1
LOC285696	ASAP2	ZFP41	ZNF441
KLF3	C2orf82	PPP2R5E	ABR
P2RX5	ADCY3	PES1	NDUFS4
SNORD115-5	ESRRG	SLC39A14	RASD2
MAGEL2	HBA2	SNORD115-48	TMCO6
PCDHGA4	CNR1	GNB4	AGAP1
FGF8	COL4A1	KRT14	GEMIN7
LAIR2	WDR52	KBTBD7	YBX1
TCERG1L	MAEL	PPP2R2D	PCDHB9
POM121L1P	CHAT	PTH2	PRHOXNB
HOXC4	C2orf54	SEN7	NINL
CAMK1G	CELSR1	PCDHA2	RBM24
NFKB2	RHCG	LOC441666	GABRG3
PLEKHH2	HERC4	ITPRIP	KLK9
OR2H1	TOX2	TMEM121	SLC39A8
SNRNP35	SCNN1A	KSR2	PAX3
FAM172A	C11orf9	PLEKHM2	C4orf51
RPS12	LANCL1	SPATA18	PDE6B
PCDHB3	ATP6V0E1	NGB	PARVB
SORBS3	RAB4A	SIX6	RBM26
C17orf62	ZAK	SLC34A2	ZNF20
SFMBT2	LOC100129637	C2orf63	BTNL2
C12orf23	CBFA2T3	AGBL5	C15orf2
GUCY1A2	SDCCAG1	COL6A3	SLC12A7
CCDC86	GPATCH4	DNAJC1	C11orf86

GENE	GENE	GENE	GENE
DIDO1	SKIV2L	DLL1	GNA11
SLC29A4	HAS1	SLC1A2	NTRK1
MKRN3	C19orf43	ALKBH4	TMEM132C
FAM134C	MCCC1	BCAN	EPS8L3
GTF2A1L	FAM181A	RASA3	LCE1C
CCDC41	ADH5	LOC728989	STK38L
PCDH9	PRMT8	MORN5	NUF2
MRPS18A	OSGIN1	LONRF2	B2M
CHPF	CACNA1A	ANKH	TMPRSS5
TBC1D15	ZNF611	TRIM2	CHRD12
FAM110C	PAOX	SLC37A2	RSPO2
C22orf9	SMARCD2	ATPGD1	SDK1
SEPSECS	WFDC3	ELOVL7	ARHGEF17
SNX15	GRM2	GFRA4	DAD1
NTSR2	NRBP2	ZNF830	COX7A2
DUOX1	FLYWCH1	ACAT2	FAM162B
PAIP2B	C7orf30	IFRD1	NF1
PRRT1	LOC389458	WWP1	SLC30A7
LOC375196	NFIA	SRPR	PRKCZ
DOCK1	KANK3	TTC39A	KLK1
MTX3	ZNF860	CADPS	S100A3
DIP2C	DYNC1H1	KIAA1324	DBH
RARG	ANO1	MGMT	CNGA3
GALR1	SNORD116-17	POP5	PLIN3
ROBO2	RNASET2	MIR548H4	RPUSD3
TAPBP	COX6C	MYH14	LOC284100
CCKAR	FUT11	psITPTE22	LOC729375
RPF2	DBC1	BCL6	HSD17B4
DIRAS3	CCDC85C	JAG2	NR2F2
ANKK1	PHF14	CCDC124	CYB5RL
ACLY	EEF1E1	ERICH1	OLIG2
C14orf73	ICOSLG	ASPSR1	ARHGAP5
GRM7	GRIN1	KCNG1	GRB2
WDR60	PF4	ARHGDIG	ZNF483
C20orf56	PM20D1	CSNK1A1	ZFP36L2
JAKMIP3	COX8A	GJD3	CRTAC1
NXN	ETFA	LOC144571	LOC654342
MCF2L2	DAB1	EPB41L4A	PAR5
SLC35F3	C17orf102	NKX2-1	PHF1
SLC6A3	MIR663	MAEA	NUBP1
GRIN2C	SLC12A2	GRHL3	ARHGAP23
TYK2	CTTN	CRIP3	REXO2
ANKMY1	MAPK9	C2orf77	TMEM106A

GENE		GENE		GENE		GENE
LDLR		SHISA9		USP13		KCNJ9
EMILIN2		FBXL7		MACC1		GSTO2
NXPH4		BCL2L14		DLG5		SEZ6
GAA		TAAR8		OC90		SASS6
COQ6		EIF4E3		OLFM1		WDR88
TBX5		LMBRD1		BRIP1		SORCS2
NOXO1		CENPB		SPG7		KRTAP3-1
MAPK8IP1		FAM54A		LRP1		SVIL
CHUK		PRPF8		FAM150B		DCC
ZNF28		TMEM102		ARL4A		FAM150A
CNTNAP4		SFRS8		CACNG2		KLF14
TTC30B		PDZD8		GNASAS		C16orf62
LMO1		AXIN1		STC1		NEU1
C6orf168		PACSIN1		C16orf11		GCM2
STK10		LTB4R		PLEKHA5		CCDC108
NOL10		HS3ST3A1		C1orf159		MIR1182
KEAP1		TFAP2A		TATDN1		RASGEF1C
KTN1		FLJ10661		KCNK2		MIR548F2
BANP		ADRA2C		ASPG		PCDHB17
RGL3		PRDM14		ELF1		LOC440925
FJX1		ATP2A2		PRDM7		KLRC1
IFT140		MCRS1		PAX6		FBRSL1
H2AFY2		YIF1B		PCDHGB1		LOC400794
SDHAF2		THBS4		ALS2		SLC6A19
MYOM2		EGLN2		TNFRSF25		CA10
AGTR1		EGFR		KIAA0913		TLX1NB
RASGRF1		GNG13		BMS1P5		IFT81
RILPL1		GPR39		GTF2F2		DLX6AS
BCL11B		TNS3		ODZ3		ADAMTS7
GLTSCR2		CREB5		LRRC32		MUC21
CRLF1		TRAFD1		EHMT1		AKR1E2
CUGBP2				SNORD115-9		CYP4X1
						SCHIP1

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs2647533	0.600166137	0.726769231	0.438567493	rs9866986	0.548781522	0.578461538	0.520110193
rs9657672	0.597081161	0.562461538	0.620385675	rs10867619	0.548779826	0.566769231	0.490909091
rs17615862	0.593204069	0.687384615	0.4877603306	rs2303879	0.548749735	0.48	0.595041322
rs4625729	0.592214452	0.710153846	0.453443526	rs7813392	0.548746345	0.531384615	0.496418733
rs12129855	0.590158508	0.504307692	0.605509642	rs6080124	0.548738292	0.493538462	0.6
rs10492998	0.585599915	0.749230769	0.414325069	rs7283316	0.548738292	0.648923077	0.41707989
rs7714670	0.585252808	0.686461538	0.465564738	rs4778426	0.548738292	0.587076923	0.512396694
rs3910138	0.585233312	0.556307692	0.590082645	rs17349719	0.548698877	0.476923077	0.620385675
rs2925034	0.585103624	0.414769231	0.734986226	rs9421389	0.548687434	0.634153846	0.418732782
rs9379179	0.584609875	0.507692308	0.629201102	rs1105572	0.5486726	0.680307692	0.407713499
rs8067791	0.584551388	0.508	0.672176309	rs15997	0.548672176	0.572923077	0.517355372
rs7191801	0.583680441	0.602461538	0.547107438	rs480414	0.548651833	0.539076923	0.537190083
rs702279	0.583624921	0.632307692	0.532321405	rs319999	0.548650985	0.662769231	0.428099174
rs10954965	0.583118033	0.547692308	0.598347107	rs12694971	0.54864039	0.514153846	0.569146006
rs1034383	0.582500106	0.461230769	0.672727273	rs8048114	0.548636576	0.507076923	0.575206612
rs1515983	0.582111464	0.711076923	0.427548209	rs4790290	0.548628947	0.569846154	0.530027548
rs10848835	0.581927951	0.702461538	0.444077135	rs11653310	0.548624709	0.663384615	0.411570248
rs1372876	0.581858868	0.461846154	0.676584022	rs10744957	0.548621742	0.46	0.640220386
rs3923367	0.581850392	0.703692308	0.416578926	rs9870523	0.548607756	0.596923077	0.484848485
rs677242	0.581724942	0.478461538	0.643526171	rs534870	0.548603094	0.530769231	0.543250689
rs2289631	0.581533376	0.428615385	0.691460055	rs697686	0.548591227	0.539076923	0.543250689
rs2251788	0.58133079	0.711384615	0.438016529	rs1303577	0.548575122	0.656307692	0.431404959
rs7110986	0.580402204	0.406769231	0.738842975	rs6488494	0.548572579	0.462153846	0.588980716
rs4969277	0.580323374	0.509230769	0.651790634	rs461409	0.548570036	0.673230769	0.412121212
rs11581005	0.580141132	0.691076923	0.41046832	rs1990594	0.548547997	0.407692308	0.673829201
rs4607200	0.579439712	0.633846154	0.48815427	rs9358075	0.548526383	0.417230769	0.663911846
rs2496622	0.578689977	0.632615385	0.533884298	rs2270637	0.548517906	0.421846154	0.676584022
rs11098735	0.578522568	0.454769231	0.710192837	rs274034	0.548508582	0.609230769	0.506336088
rs1463366	0.578432719	0.408615385	0.705785124	rs9596775	0.54848951	0.545846154	0.544903581
rs4761128	0.578388218	0.743846154	0.408815427	rs9576140	0.548484425	0.688	0.406060606
rs11051794	0.578059335	0.508615385	0.65599449	rs6726230	0.548470862	0.640923077	0.452892562
rs7804911	0.577905065	0.475692308	0.667768595	rs6580870	0.548467472	0.461538462	0.635812672
rs16858984	0.577430388	0.680923077	0.41322314	rs6977125	0.5484662	0.615076923	0.47107438
rs10117055	0.577379106	0.445846154	0.700826446	rs10872009	0.548465777	0.633230769	0.472727273
rs41388348	0.577335029	0.507692308	0.647933884	rs11172333	0.548463234	0.507384615	0.495316804
rs2228225	0.577225683	0.662769231	0.437465565	rs609749	0.54846281	0.593846154	0.508539945
rs1751956	0.577203221	0.405230769	0.741046832	rs2414817	0.548450943	0.538153846	0.532782369
rs3808942	0.577012503	0.744615385	0.406060606	rs10984217	0.548446705	0.507384615	0.582369146
rs735942	0.576965035	0.568615385	0.572451791	rs2197465	0.54844289	0.458153846	0.587878788
rs5994376	0.576459419	0.491384615	0.658953168	rs2817781	0.548406866	0.672	0.402754821
rs9967367	0.576206823	0.578153846	0.548209366	rs11233339	0.548394151	0.413846154	0.678236915
rs12963675	0.576171223	0.428307692	0.703581267	rs9518832	0.548333121	0.671076923	0.408264463

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs4516035	0.576136893	0.477230769	0.646280992	rs4667005	0.54828036	0.554769231	0.507988981
rs761248	0.57593558	0.624923077	0.506336088	rs4035	0.548320407	0.650153846	0.427038567
rs1408102	0.575768595	0.546769231	0.561983471	rs11863694	0.548317016	0.681230769	0.41707989
rs4440831	0.575636787	0.516923077	0.593939394	rs13432421	0.548311507	0.546461538	0.514049587
rs1035775	0.575432507	0.554153846	0.586225895	rs14336089	0.548300911	0.572923077	0.495316804
rs2278507	0.574475524	0.499384615	0.619283747	rs1017104	0.548283111	0.421846154	0.651239669
rs1042078	0.574278449	0.547384615	0.578512397	rs7756967	0.548282687	0.430769231	0.657300275
rs11063610	0.574182666	0.639384615	0.516804408	rs2838820	0.548282263	0.404923077	0.673278237
rs12417345	0.574056368	0.618153846	0.507438017	rs4884684	0.548276754	0.470153846	0.604407713
rs9368920	0.574006781	0.601538462	0.534435262	rs13088237	0.54827633	0.556923077	0.49862259
rs9567688	0.573800805	0.561538462	0.580165289	rs10484867	0.548275482	0.464615385	0.55922865
rs6845132	0.573664336	0.712923077	0.419824711	rs240734	0.548259801	0.571384615	0.514600551
rs17801625	0.573318924	0.476923077	0.668319559	rs4771969	0.54824751	0.554769231	0.517906336
rs1029549	0.573285442	0.500307692	0.634710744	rs4785602	0.548244967	0.469846154	0.620385675
rs17041890	0.573279085	0.476307692	0.669421488	rs4721135	0.548239881	0.579384615	0.539393939
rs2201593	0.573223564	0.648923077	0.489256198	rs17105685	0.548222929	0.422461538	0.675482094
rs978935	0.573185421	0.639076923	0.480951736	rs11839514	0.548221233	0.500923077	0.584573003
rs360833	0.573053189	0.655692308	0.489807163	rs6473804	0.548201738	0.476615385	0.591735537
rs3846663	0.573049799	0.530461538	0.618732782	rs2910400	0.548200042	0.586461538	0.515151515
rs1483179	0.572929032	0.640923077	0.468870523	rs11122844	0.548199619	0.420923077	0.659504132
rs10491954	0.572914177	0.646153846	0.488705234	rs249276	0.548188175	0.594461538	0.473278237
rs2606635	0.57290231	0.570153846	0.558677686	rs2400963	0.548166561	0.602153846	0.494214876
rs875373	0.572875609	0.515384615	0.630853994	rs1487864	0.548163594	0.429846154	0.659504132
rs10948653	0.572775588	0.693846154	0.416528926	rs10495941	0.548151727	0.609846154	0.480440771
rs2237066	0.572736173	0.492307692	0.614876033	rs367164	0.548144946	0.468307692	0.619834711
rs10023064	0.572705658	0.650153846	0.484297521	rs4555439	0.548110617	0.608923077	0.4815427
rs2122481	0.572661157	0.452923077	0.662258953	rs989364	0.548107226	0.507692308	0.587327824
rs1749859	0.572627252	0.507692308	0.644077135	rs2824671	0.54809875	0.694461538	0.40661157
rs8004280	0.572589532	0.664615385	0.428650138	rs17649190	0.548096631	0.439076923	0.66446281
rs8126205	0.572574274	0.491076923	0.638016529	rs6596370	0.548094512	0.537846154	0.550413223
rs4532668	0.572495868	0.469230769	0.601652893	rs7070956	0.548078406	0.679076923	0.411019284
rs10914615	0.572202585	0.531076923	0.593939394	rs292213	0.54807544	0.609538462	0.496418733
rs6550981	0.571946599	0.421846154	0.714049587	rs4818873	0.548060606	0.469230769	0.631955923
rs6860985	0.571847001	0.517230769	0.653443526	rs10500030	0.548058063	0.491076923	0.60203857
rs4983125	0.571437593	0.562461538	0.565289256	rs17157663	0.548010595	0.482769231	0.602754821
rs12904303	0.57140623	0.467384615	0.652341598	rs11117291	0.547998305	0.634153846	0.450688705
rs6823628	0.571282475	0.523692308	0.612121212	rs4754095	0.547980081	0.637692308	0.412121212
rs3825932	0.571208731	0.577538462	0.544903581	rs5995155	0.547961433	0.554769231	0.53553719
rs1269287	0.571081585	0.734461538	0.412121212	rs6933583	0.547959737	0.608923077	0.483746556
rs6944935	0.571072685	0.577538462	0.537741047	rs930514	0.547919051	0.664923077	0.407162534
rs10926222	0.571021674	0.531692308	0.598898072	rs3006496	0.54791566	0.539692308	0.549862259
rs3135093	0.570770078	0.610153846	0.517906336	rs1811272	0.547905065	0.554461538	0.538842975

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs7684254	0.570458572	0.578769231	0.503030303	rs6140981	0.547893198	0.507384615	0.60407713
rs10914446	0.570342869	0.641230769	0.500275482	rs4858604	0.547893198	0.547076923	0.545454545
rs2800156	0.570323797	0.492	0.660055096	rs17828333	0.547885993	0.586461538	0.507438017
rs3996004	0.570189023	0.452615385	0.67107438	rs7742836	0.54788345	0.491692308	0.606060606
rs3116602	0.570181394	0.734461538	0.411019284	rs270215	0.54786904	0.6	0.449035813
rs5771040	0.570048315	0.508	0.603856749	rs2080652	0.547855902	0.430153846	0.680991736
rs3008527	0.570013986	0.569846154	0.55922865	rs17635849	0.547843187	0.453846154	0.642975207
rs12932881	0.569812672	0.511692308	0.564187328	rs9613438	0.547828353	0.672	0.420385675
rs719787	0.569774529	0.640307692	0.512396694	rs9648374	0.547815639	0.561846154	0.521212121
rs9298792	0.569717737	0.406769231	0.739393939	rs4405545	0.547813944	0.648923077	0.422038567
rs1032034	0.569708413	0.461230769	0.649586777	rs1506452	0.547810553	0.566153846	0.48815427
rs2970359	0.569681712	0.561846154	0.566942149	rs2442562	0.5477936	0.671384615	0.412121212
rs17422587	0.569660521	0.489846154	0.608264463	rs11234795	0.547770714	0.483384615	0.604958678
rs4791845	0.569579996	0.570153846	0.566391185	rs10062513	0.54777029	0.499076923	0.592837466
rs2052003	0.569478703	0.609538462	0.507988981	rs320947	0.547762662	0.687076923	0.405509642
rs2047382	0.569439288	0.695076923	0.421487603	rs1484444	0.547760542	0.553846154	0.520661157
rs2834506	0.569335961	0.633846154	0.50137741	rs10973014	0.54775588	0.57538462	0.507438017
rs3904775	0.569291375	0.679692308	0.446280992	rs2352452	0.547754185	0.562769231	0.54214876
rs2397146	0.569257046	0.602461538	0.537741047	rs748142	0.547736385	0.443692308	0.643526171
rs7044631	0.569195168	0.539076923	0.605509642	rs2024269	0.547731299	0.655692308	0.437465565
rs1202592	0.569148125	0.695076923	0.425895317	rs2186919	0.547724094	0.473846154	0.55922865
rs7560385	0.569118881	0.546461538	0.568044077	rs7669724	0.547716465	0.452615385	0.633057851
rs1045873	0.569082009	0.546769231	0.603856749	rs612040	0.547701208	0.642153846	0.425895317
rs4629410	0.568975207	0.453846154	0.67107438	rs10842230	0.547692732	0.422461538	0.662809917
rs1523320	0.568899343	0.527384615	0.48815427	rs9300935	0.547679593	0.448	0.643526171
rs2475802	0.568778555	0.571692308	0.574655647	rs10490845	0.547676626	0.557846154	0.51184573
rs2712735	0.568765416	0.545538462	0.574104683	rs4715624	0.547669845	0.624615385	0.469421488
rs6816344	0.568716253	0.572	0.558677686	rs6545185	0.547646959	0.510133846	0.579614325
rs10113903	0.568656071	0.678769231	0.445179063	rs1475111	0.547636364	0.602461538	0.463911846
rs11135724	0.568527654	0.632615385	0.487052342	rs10168628	0.547626616	0.531076923	0.551515152
rs304650	0.568510278	0.695076923	0.45599449	rs1152663	0.547606273	0.585846154	0.500275482
rs2000658	0.568490782	0.405230769	0.731129477	rs1323998	0.547597796	0.541538462	0.514049587
rs1629874	0.568447976	0.672923077	0.428650138	rs11678782	0.547582962	0.547692308	0.543801653
rs17554682	0.568422971	0.554769231	0.568044077	rs354731	0.5475783	0.576307692	0.469421488
rs2436134	0.568389913	0.487384615	0.614325069	rs25736	0.54757491	0.486461538	0.58677666
rs896560	0.568387794	0.648307692	0.487603306	rs2426002	0.547555838	0.648923077	0.443526171
rs967963	0.568379318	0.577538462	0.549311295	rs1514992	0.547534647	0.406769231	0.680991736
rs989661	0.568217843	0.483692308	0.655096419	rs6817250	0.547532528	0.685230769	0.428099174
rs9821733	0.568214452	0.674461538	0.405509642	rs12243919	0.547520237	0.438769231	0.672727273
rs850056	0.568138165	0.468	0.662258953	rs9465181	0.547502861	0.476923077	0.613774105
rs2917702	0.568109769	0.687692308	0.401101928	rs7812091	0.547464293	0.452307692	0.639669421
rs7651106	0.567990676	0.576923077	0.546556474	rs11003694	0.547457512	0.639076923	0.465013774

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs17302818	0.567987285	0.524307692	0.61046832	rs18109570	0.547454122	0.492615385	0.596143251
rs2812378	0.567913117	0.616615385	0.477134986	rs4878520	0.547443526	0.556923077	0.459504132
rs537124	0.567853783	0.594461538	0.52231405	rs2511733	0.54743505	0.430461538	0.64077135
rs11006606	0.567827506	0.483692308	0.631955923	rs1174246	0.547427845	0.536923077	0.44077135
rs11107153	0.567760542	0.672923077	0.446280992	rs7834780	0.547410892	0.546153846	0.55469972
rs10039846	0.567671964	0.468	0.636914601	rs13251640	0.547399449	0.405230769	0.680440771
rs10514824	0.567658826	0.702769231	0.42972066	rs114873	0.547396906	0.445538462	0.643526171
rs1095014	0.567597796	0.538461538	0.574104683	rs2280630	0.547390125	0.601230769	0.487052342
rs856168	0.5675605	0.505846154	0.608264463	rs17788859	0.547363848	0.625230769	0.483911846
rs4948800	0.567515575	0.446461538	0.661157025	rs4953715	0.547355796	0.516923077	0.555371901
rs1405609	0.567303818	0.408923077	0.690909091	rs10952132	0.547354524	0.437538462	0.657300275
rs4869954	0.567267641	0.554461538	0.570247934	rs385130	0.547340962	0.461230769	0.611570248
rs1335477	0.567235431	0.680307692	0.456138347	rs750160	0.547327824	0.509230769	0.58292011
rs1479835	0.567157449	0.445846154	0.66446281	rs373533	0.547326552	0.444307692	0.646831956
rs4840037	0.567151363	0.562769231	0.578512397	rs1798805	0.547320195	0.612923077	0.487603306
rs9408826	0.567112524	0.423384615	0.704683196	rs8083764	0.547297733	0.587076923	0.511294766
rs1873870	0.567010384	0.570461538	0.545454545	rs12578981	0.547290951	0.562153846	0.533884298
rs1898199	0.566827718	0.532923077	0.585123967	rs2190921	0.54728078	0.444	0.63415978
rs1076447	0.566825599	0.618461538	0.50523416	rs43111	0.547280356	0.438153846	0.649035813
rs2578414	0.566737021	0.5	0.626446281	rs1366810	0.54727527	0.563384615	0.52892562
rs3775214	0.566663276	0.693538462	0.404407713	rs453577	0.547273999	0.648	0.455096419
rs10517889	0.56660167	0.423384615	0.693663912	rs4239263	0.547257894	0.508923077	0.562534435
rs1777458	0.566578936	0.438769231	0.656138347	rs10768122	0.547231617	0.468	0.604958678
rs4751335	0.566399237	0.470769231	0.563085399	rs4705028	0.547217707	0.42	0.641322314
rs10799006	0.566232676	0.592923077	0.534435262	rs1688631	0.547203645	0.451692308	0.553168044
rs4640066	0.566167408	0.609538462	0.506336088	rs2219770	0.547203221	0.636	0.407713499
rs7004854	0.566138589	0.586153846	0.564738292	rs10736390	0.547200254	0.438461538	0.634710744
rs10860174	0.566118669	0.530461538	0.6	rs6000135	0.547187116	0.512923077	0.584022039
rs10979395	0.566044077	0.531384615	0.561983471	rs2393904	0.547168468	0.585846154	0.508539945
rs3743180	0.566015681	0.421846154	0.674931129	rs1875108	0.547166773	0.586461538	0.516804408
rs4859673	0.565978809	0.657538462	0.459504132	rs780807	0.54714982	0.437230769	0.656749311
rs17353923	0.565972452	0.492923077	0.620385675	rs10115383	0.547143463	0.478461538	0.532231405
rs17691755	0.565965671	0.531384615	0.61046832	rs2688515	0.547143039	0.416615385	0.651790634
rs8017825	0.565894045	0.468307692	0.650137741	rs10930659	0.547123119	0.487384615	0.461157025
rs6971101	0.565843611	0.453538462	0.683016529	rs7223424	0.547109981	0.546461538	0.538842975
rs10832565	0.565771138	0.459384615	0.552066116	rs2606167	0.547099809	0.617230769	0.460055096
rs1410113	0.565710108	0.424	0.685399449	rs12613274	0.547084552	0.516	0.55922865
rs12327456	0.565535071	0.687076923	0.436363636	rs2115063	0.54708328	0.522153846	0.554269972
rs2060715	0.565441831	0.429538462	0.685950413	rs29645	0.547079466	0.593538462	0.516804408
rs13119207	0.565416826	0.570153846	0.571349862	rs1992405	0.547066328	0.453230769	0.636914601
rs4706705	0.565369782	0.742769231	0.407713499	rs2919852	0.547059123	0.531076923	0.537741047
rs7709207	0.565309176	0.514769231	0.612672176	rs12510308	0.547057004	0.680307692	0.41046832

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs2282498	0.56329858	0.476307692	0.624793388	rs689338	0.547026912	0.680307692	0.411019284
rs2780626	0.565290104	0.680615385	0.456749311	rs7565113	0.547012926	0.655384615	0.451790634
rs34628921	0.565273575	0.560923077	0.546556474	rs17667613	0.547012079	0.484615385	0.62646281
rs217519	0.565260437	0.724923077	0.401101928	rs17176567	0.547010807	0.524615385	0.548209366
rs1740186	0.565204069	0.664307692	0.452892562	rs17320342	0.547004026	0.665846154	0.42369146
rs255155	0.565196864	0.663076923	0.430853994	rs6049473	0.546996398	0.444923077	0.631955923
rs2124141	0.56512651	0.508	0.616528926	rs748696	0.54698114	0.489823077	0.61707989
rs372883	0.56510659	0.433230769	0.568044077	rs41406547	0.546967578	0.405846154	0.654545455
rs655823	0.565019284	0.619076923	0.507438017	rs1442343	0.54696673	0.505846154	0.60661157
rs12893510	0.565018012	0.600615385	0.525619835	rs4582347	0.546962492	0.562153846	0.507988981
rs10833849	0.564867557	0.431076923	0.654545455	rs17685537	0.546951897	0.507384615	0.58953168
rs7617553	0.56482348	0.664307692	0.467768595	rs12198460	0.546950625	0.587076923	0.512396694
rs6463117	0.564786607	0.633538462	0.496418733	rs4704134	0.546943844	0.625846154	0.446280992
rs160134	0.564700148	0.523076923	0.598347107	rs357833	0.546934944	0.586153846	0.430853994
rs10937304	0.564650562	0.427692308	0.695316804	rs3813097	0.546927739	0.665538462	0.426997245
rs501082	0.564550964	0.556307692	0.477134986	rs2146959	0.54691672	0.422461538	0.591735537
rs6038855	0.564550117	0.475076923	0.636363636	rs6959554	0.546916296	0.492	0.584573003
rs2150367	0.564531469	0.407076923	0.523415978	rs538399	0.54686459	0.523076923	0.562534435
rs4904639	0.5644573	0.600923077	0.515151515	rs607230	0.546854418	0.585846154	0.492011019
rs2158231	0.564445857	0.48923077	0.605509642	rs6432412	0.546795084	0.547692308	0.543250689
rs733697	0.564367875	0.414769231	0.711294766	rs4671975	0.546784488	0.467692308	0.624242424
rs4981135	0.564333545	0.401538462	0.679898907	rs10842277	0.546781522	0.648	0.441873278
rs3740615	0.564222929	0.430769231	0.694214876	rs6461569	0.546750159	0.524	0.551515152
rs445593	0.564206823	0.460307692	0.667768595	rs3809618	0.546725154	0.578153846	0.522865014
rs17356059	0.564128417	0.678769231	0.43415978	rs4483218	0.546678957	0.523384615	0.557024793
rs954492	0.564088154	0.710461538	0.404958678	rs17236056	0.546664548	0.439076923	0.647933884
rs10774354	0.564018224	0.492923077	0.618181818	rs2147104	0.546660733	0.608615385	0.512396694
rs17681684	0.564008476	0.608923077	0.494214876	rs38826	0.546635304	0.672307692	0.401101928
rs1110116	0.564008053	0.678769231	0.41046832	rs470259	0.546633609	0.579076923	0.517355372
rs7316328	0.563977538	0.555384615	0.566391185	rs4129470	0.546631914	0.656307692	0.430853994
rs11170013	0.563966942	0.529846154	0.595041322	rs4851752	0.546628523	0.421538462	0.666115702
rs10846858	0.563921594	0.476923077	0.654545455	rs1388206	0.546623013	0.610153846	0.471625344
rs3476	0.563855054	0.461230769	0.663911846	rs2416313	0.546614961	0.492615385	0.602203857
rs11638818	0.563769019	0.648615385	0.4815427	rs909975	0.546611994	0.578461538	0.446831956
rs10128061	0.563630006	0.575076923	0.422038567	rs10121214	0.546604365	0.594461538	0.488705234
rs2033083	0.563626192	0.423384615	0.573553719	rs1908587	0.546590379	0.505538462	0.506336088
rs9296302	0.563567281	0.407076923	0.717906336	rs12450199	0.546589532	0.588	0.474931129
rs2951874	0.563490146	0.493230769	0.617630854	rs1958333	0.546580831	0.484615385	0.60661157
rs4948428	0.563429116	0.534769231	0.499173554	rs35713415	0.546574274	0.415076923	0.665564738
rs2259298	0.5633340114	0.688307692	0.411570248	rs2641698	0.54657046	0.657230769	0.444628099
rs682431	0.563336724	0.672923077	0.444628099	rs15153188	0.546556474	0.547384615	0.550413223
rs2717907	0.563320619	0.422153846	0.708539945	rs9832360	0.546555626	0.640923077	0.44077135

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs343970	0.563287137	0.452	0.673829201	rs4757484	0.546555202	0.649230769	0.424793388
rs22482302	0.563269337	0.672615385	0.443526171	rs2184401	0.546555202	0.600923077	0.480991736
rs926392	0.563240517	0.436923077	0.690358127	rs93934914	0.546554355	0.431076923	0.630853994
rs6005236	0.563226955	0.476	0.626446281	rs791075	0.546532316	0.688615385	0.407162534
rs1927103	0.563177368	0.500615385	0.61046832	rs9660184	0.546520449	0.648	0.436914601
rs10021880	0.563151939	0.424307692	0.680991736	rs10174389	0.546519178	0.586461538	0.492561983
rs3822365	0.563048951	0.678461538	0.442424242	rs950089	0.546513668	0.413230769	0.671625344
rs10323916	0.563009536	0.641230769	0.480991736	rs481855	0.546505616	0.553538462	0.543250689
rs989430	0.562959525	0.437846154	0.676033058	rs8826441	0.546499158	0.546789231	0.525068871
rs2143877	0.562891138	0.562769231	0.544903581	rs1029349	0.546496292	0.444923077	0.639118457
rs4699559	0.562887052	0.625230769	0.492011019	rs6785581	0.546492053	0.407076923	0.672176309
rs1425025	0.562868404	0.554769231	0.553719008	rs1422587	0.546483577	0.656	0.435812672
rs7099380	0.562863318	0.537846154	0.477134986	rs6532676	0.546477644	0.4735538462	0.591735537
rs7723716	0.562833651	0.579692308	0.492561983	rs9316663	0.546475948	0.617846154	0.436914601
rs7454457	0.562637847	0.507692308	0.599449036	rs12610130	0.54646832	0.506769231	0.570247934
rs329675	0.562582751	0.695692308	0.41322314	rs4720876	0.546465353	0.624923077	0.463911846
rs12670725	0.562530621	0.427384615	0.672176309	rs13072883	0.546463658	0.422789231	0.645730028
rs7037570	0.562434414	0.494153846	0.629752066	rs11207282	0.546452214	0.592615385	0.485950413
rs5003374	0.562402628	0.510153846	0.580716253	rs4304868	0.546436957	0.4366615385	0.660055096
rs4936957	0.562393304	0.594769231	0.516233444	rs7178655	0.5464217	0.538461538	0.527272727
rs4509530	0.562386523	0.600307692	0.519008264	rs12981961	0.54641619	0.523692308	0.551515152
rs2351379	0.562328459	0.507692308	0.612672176	rs157093	0.546391185	0.671384615	0.445179063
rs7961115	0.562302183	0.679076923	0.402203857	rs2246538	0.546388118	0.563076923	0.5184573
rs4731689	0.562286501	0.632923077	0.469972452	rs10817777	0.54638737	0.578153846	0.506336088
rs6533118	0.562273787	0.661538462	0.469421488	rs904648	0.546375079	0.416	0.674380165
rs2815378	0.562230981	0.553538462	0.568044077	rs1680726	0.546374232	0.498461538	0.587878788
rs1076344	0.562166137	0.507384615	0.616528926	rs17651892	0.546361517	0.452	0.666115702
rs6108927	0.562097054	0.561538462	0.559779614	rs586225	0.546356855	0.608923077	0.465013774
rs17042619	0.562095359	0.421538462	0.69862259	rs4130266	0.546339055	0.586769231	0.509090909
rs1947234	0.562050011	0.640923077	0.492011019	rs6979397	0.546327612	0.624923077	0.456198347
rs10743273	0.562019496	0.592615385	0.550964187	rs957979	0.546313702	0.663076923	0.417630854
rs2382859	0.562014834	0.406153846	0.678787879	rs710050	0.546302606	0.595076923	0.482644628
rs4754513	0.561966518	0.496307692	0.592837466	rs2666229	0.546301335	0.609846154	0.4815427
rs12474952	0.561965671	0.557230769	0.54600551	rs6482199	0.546301335	0.423076923	0.650688705
rs136944	0.561957618	0.404923077	0.716804408	rs2910810	0.546286925	0.489846154	0.58953168
rs1452494	0.561928375	0.431076923	0.666666667	rs9526823	0.546282687	0.562461538	0.526721763
rs9521520	0.561891502	0.568615385	0.547107438	rs2884208	0.54627082	0.539384615	0.548760331
rs2541953	0.561888536	0.476615385	0.646280992	rs2252783	0.546269549	0.654789231	0.441322314
rs6873118	0.56188684	0.570461538	0.528374656	rs2885	0.54626192	0.477230769	0.621487603
rs1997716	0.56186904	0.538153846	0.564187328	rs883410	0.546261072	0.405538462	0.687603306
rs560056	0.561837677	0.585846154	0.519592279	rs2071864	0.546261072	0.405538462	0.538842975
rs4933299	0.561835982	0.632615385	0.469421488	rs2866180	0.54625302	0.406461538	0.63030303

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs461863	0.561820301	0.672615385	0.419283747	rs1448872	0.546249205	0.484615385	0.59184573
rs1034593	0.561773257	0.445846154	0.640220386	rs8057341	0.546244967	0.570461538	0.50523416
rs6958639	0.561764781	0.641230769	0.495904132	rs2140330	0.546236915	0.444307692	0.631955923
rs10177101	0.561755033	0.562461538	0.550964187	rs6471128	0.5462331	0.562769231	0.519008264
rs10780305	0.561722823	0.507384615	0.61707989	rs11231741	0.546205552	0.546461538	0.54214876
rs1863037	0.561719008	0.595076923	0.504603196	rs291640	0.546188599	0.484615385	0.625895317
rs9379986	0.561719008	0.492307692	0.61045832	rs750753	0.546180971	0.444923077	0.620385675
rs4940339	0.561691884	0.492615385	0.626997245	rs2823253	0.546164018	0.584923077	0.473278237
rs2079644	0.561625344	0.484923077	0.609917355	rs8898585	0.546151727	0.627076923	0.412121212
rs11085020	0.561596949	0.641846154	0.469421488	rs9315638	0.546146641	0.641230769	0.441322314
rs7981495	0.561518118	0.461230769	0.656198347	rs7104953	0.546144098	0.508	0.570247934
rs11645864	0.561488027	0.413846154	0.686501377	rs1985053	0.54613096	0.531692308	0.548760331
rs11141878	0.561460479	0.694461538	0.404407713	rs6030341	0.546130536	0.425230769	0.630853994
rs6695131	0.561441831	0.556615385	0.565289256	rs17113829	0.546109345	0.475384615	0.602203857
rs12243326	0.561438017	0.584923077	0.512947658	rs17082947	0.546106802	0.657538462	0.425895317
rs306593	0.561421488	0.671692308	0.432506887	rs9376092	0.54608985	0.516923077	0.564187328
rs2587427	0.561358762	0.540307692	0.572451791	rs2864668	0.54608773	0.406461538	0.660055096
rs7310929	0.5613541	0.508307692	0.587878788	rs1531581	0.546085611	0.586153846	0.498071625
rs1961460	0.561324009	0.648	0.482644628	rs818185	0.546073321	0.460307692	0.597245179
rs10055677	0.561304514	0.566153846	0.49476584	rs4281136	0.546053825	0.627384615	0.40661157
rs16908004	0.561299004	0.595076923	0.519559229	rs221668	0.546027548	0.646769231	0.425344353
rs2761201	0.561177368	0.537846154	0.585674931	rs11722715	0.546013562	0.625230769	0.45785124
rs7902673	0.561170587	0.608307692	0.509641873	rs591730	0.546002119	0.539384615	0.536088154
rs9874195	0.56114431	0.536307692	0.588980716	rs2255071	0.545999152	0.530769231	0.524517906
rs11098524	0.561138801	0.610769231	0.502479339	rs11256338	0.545982623	0.588615385	0.519559229
rs4592619	0.561066328	0.421538462	0.703581267	rs1028827	0.545978385	0.524923077	0.545454545
rs10963844	0.560981988	0.681230769	0.445179063	rs651141	0.545974995	0.641846154	0.402754821
rs7955714	0.560968426	0.609846154	0.506887032	rs2727692	0.545958042	0.54	0.564738292
rs2278729	0.560965459	0.648923077	0.462258953	rs2752153	0.545951685	0.632307692	0.462258953
rs9517723	0.560959949	0.548307692	0.52231405	rs10752589	0.545943208	0.617538462	0.488870523
rs1348469	0.560955711	0.601846154	0.509641873	rs2823577	0.545931341	0.665230769	0.472589532
rs3743560	0.560950625	0.491384615	0.614325069	rs7507442	0.545930918	0.508615385	0.581267218
rs7162038	0.560906124	0.678769231	0.431955923	rs3212335	0.545927951	0.632615385	0.412672176
rs869964	0.560905277	0.415076923	0.663911846	rs1216352	0.545916084	0.521538462	0.555922865
rs4798499	0.560894257	0.578153846	0.550413223	rs3919487	0.545907608	0.536923077	0.558126722
rs6850096	0.560856114	0.539076923	0.556473829	rs13227263	0.545898284	0.445538462	0.631955923
rs17277691	0.560835346	0.554461538	0.563636364	rs9529278	0.545897436	0.522153846	0.576859504
rs988213	0.560807798	0.525846154	0.490909091	rs10966901	0.545896588	0.452615385	0.620385675
rs11599327	0.560801441	0.632615385	0.485950413	rs12673360	0.545893198	0.632	0.449586777
rs2675717	0.560778555	0.413230769	0.694214876	rs4842266	0.545882602	0.618153846	0.457300275
rs1449302	0.560705658	0.400615385	0.6815427	rs1016740	0.545868616	0.569230769	0.525068871
rs203257	0.560670057	0.602153846	0.529476584	rs509242	0.545850816	0.632923077	0.450688705

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs4925109	0.560624709	0.624	0.462258953	rs13020302	0.545844035	0.483692308	0.581818182
rs12571964	0.560588684	0.446461538	0.67768595	rs17485868	0.545838949	0.555076923	0.515151515
rs11141992	0.560582327	0.405846154	0.684297521	rs6497072	0.545834711	0.632923077	0.459504132
rs2809247	0.560574698	0.673230769	0.428650138	rs7905804	0.545831744	0.649230769	0.443526171
rs11813508	0.560542488	0.546769231	0.554820937	rs6727754	0.545814791	0.417846154	0.651239669
rs11059100	0.56052723	0.689538462	0.402203857	rs2965160	0.545804196	0.623076923	0.418181818
rs9747501	0.560526807	0.515692308	0.596694215	rs2162159	0.545794448	0.413230769	0.690358127
rs1897472	0.560479763	0.601230769	0.504132231	rs1716170	0.545788091	0.562769231	0.523415978
rs13061242	0.560461115	0.5	0.61872782	rs936025	0.5457847	0.42	0.673829201
rs12186611	0.560447552	0.546769231	0.584022039	rs134748	0.545772409	0.515384615	0.568044077
rs549433	0.56041958	0.608	0.506336088	rs2363876	0.545768595	0.664	0.44077135
rs17553620	0.560414071	0.531076923	0.578512397	rs11219510	0.545757576	0.439076923	0.65785124
rs6742536	0.560401356	0.522461538	0.598347107	rs4775508	0.54575249	0.412923077	0.651239669
rs963666	0.560375079	0.646769231	0.468870523	rs1861619	0.54574698	0.414789231	0.661157025
rs1330106	0.560356008	0.686461538	0.433057851	rs799559	0.54572367	0.555692308	0.525068871
rs2512106	0.560304302	0.460615385	0.63399449	rs4127445	0.545704175	0.435692308	0.586225895
rs454583	0.560295825	0.703076923	0.425344353	rs10500707	0.545587646	0.6	0.478787879
rs12410581	0.560264887	0.711692308	0.423140496	rs9944879	0.545569845	0.452307692	0.636363636
rs7146427	0.560254715	0.476923077	0.617630854	rs17356414	0.5455669421	0.546461538	0.53553719
rs11946495	0.560248782	0.593846154	0.49201019	rs949492	0.545565183	0.472923077	0.507988981
rs2157605	0.560204704	0.407384615	0.705785124	rs403608	0.545566264	0.696	0.417630854
rs11640870	0.560189871	0.408923077	0.66446281	rs563573	0.54553933	0.530153846	0.52892562
rs4702534	0.560158932	0.522769231	0.596143251	rs12954678	0.545534668	0.561846154	0.511294766
rs7968902	0.560147065	0.469230769	0.650137741	rs6693833	0.545528735	0.492307692	0.587878788
rs2223814	0.560118245	0.459692308	0.643526171	rs1612695	0.545501187	0.406769231	0.688705234
rs8107912	0.560107226	0.632923077	0.451790634	rs894393	0.545593558	0.664307692	0.423140496
rs2046411	0.560046196	0.563076923	0.558677686	rs935622	0.545587201	0.504923077	0.575757576
rs16851037	0.560041534	0.484	0.637485565	rs6498482	0.545572367	0.536615385	0.478787879
rs10347091	0.560022039	0.493230769	0.485950413	rs7803908	0.545568976	0.546461538	0.521763085
rs4879930	0.55996609	0.577846154	0.548760331	rs10746902	0.545559229	0.632	0.46446281
rs1593519	0.559926679	0.595076923	0.510192837	rs9356383	0.545553719	0.407384615	0.64738292
rs12785018	0.559903369	0.500307692	0.597245179	rs4237677	0.545552448	0.429538462	0.658402204
rs2461796	0.559859292	0.430461538	0.656749311	rs4627591	0.545546514	0.585846154	0.485950413
rs4135168	0.559789362	0.547384615	0.553719008	rs10186393	0.545534223	0.424923077	0.67768595
rs10496417	0.559684679	0.680923077	0.420936639	rs9613519	0.545518542	0.655692308	0.435261708
rs1325266	0.559677898	0.564923077	0.43415978	rs9515828	0.54548506	0.607692308	0.470523416
rs942880	0.559579672	0.633230769	0.488705234	rs2583556	0.545484637	0.532307692	0.553719008
rs2949641	0.559505828	0.592923077	0.482093664	rs6881806	0.545482941	0.695076923	0.402754621
rs2164422	0.559500318	0.453230769	0.661707989	rs11081336	0.545479551	0.68	0.404958678
rs1004526	0.559486332	0.647692308	0.476584022	rs4077818	0.54547616	0.663692308	0.41322314
rs1458521	0.559473617	0.570769231	0.49476584	rs1546479	0.54547616	0.562461538	0.521763085
rs1434950	0.559424031	0.469230769	0.637465565	rs7091819	0.545447764	0.477538462	0.603856749

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs17730540	0.559358762	0.406461538	0.701377741	rs1907220	0.545446493	0.561538462	0.527272727
rs1869890	0.559349015	0.429538462	0.656198347	rs9317421	0.545438864	0.483692308	0.595041322
rs10511359	0.559294342	0.624615385	0.461707989	rs9306924	0.545424454	0.648307692	0.445730028
rs11556924	0.559235007	0.5	0.601101928	rs11633365	0.545423183	0.600615385	0.480991736
rs10043097	0.55919644	0.468307692	0.651790634	rs117131479	0.545421488	0.624615385	0.456198347
rs252026	0.559146853	0.570153846	0.540495868	rs12311525	0.545413859	0.616615385	0.454545455
rs12959290	0.559116338	0.585230769	0.492561983	rs17188630	0.545409197	0.54	0.536639118
rs7721185	0.559095571	0.624923077	0.493112948	rs4799084	0.545398178	0.430153846	0.646831956
rs941009	0.559087095	0.687692308	0.445730028	rs3782752	0.545397754	0.624615385	0.455647383
rs2144456	0.559087095	0.458769231	0.652341598	rs7118508	0.54539182	0.473230769	0.570247934
rs4750340	0.558994702	0.574769231	0.501377741	rs6824779	0.545386311	0.467692308	0.618181818
rs755723	0.558971816	0.688307692	0.424242424	rs9333211	0.545376139	0.617230769	0.463911846
rs10765400	0.558941725	0.508307692	0.61322314	rs2412692	0.545373172	0.523692308	0.548760331
rs2443970	0.558940453	0.648923077	0.461707989	rs2541522	0.545363001	0.601846154	0.482093664
rs7566559	0.558938334	0.664307692	0.428099174	rs203717	0.545363001	0.500615385	0.585674931
rs4980271	0.558892986	0.469846154	0.653443526	rs8081054	0.545361305	0.617846154	0.449586777
rs1641025	0.558883662	0.633538462	0.480991736	rs1415701	0.545355372	0.616615385	0.478787879
rs11013938	0.558847637	0.538769231	0.586126722	rs6487903	0.545324909	0.447076923	0.537190083
rs9977697	0.558813732	0.672307692	0.446280992	rs5018568	0.545315533	0.640615385	0.453443526
rs9825129	0.558807112	0.664307692	0.458402204	rs2837304	0.545313838	0.563384615	0.52892562
rs7083381	0.558784488	0.531692308	0.59338843	rs1498194	0.545307904	0.457846154	0.577410468
rs6079235	0.558782369	0.553538462	0.553719008	rs2015190	0.545307057	0.580307692	0.402203857
rs188324	0.558684043	0.577230769	0.539393939	rs833995	0.545287137	0.452615385	0.64077135
rs9496482	0.558669633	0.484	0.673140496	rs12707695	0.545281704	0.531692308	0.544352617
rs7201739	0.558632337	0.523692308	0.582369146	rs2796471	0.545264675	0.499384615	0.606060606
rs5025315	0.558625556	0.437846154	0.656198347	rs2979025	0.545258741	0.523692308	0.564738292
rs1917411	0.558590803	0.678769231	0.444628099	rs7731890	0.545249841	0.473538462	0.576859504
rs10919317	0.558564103	0.549230769	0.552066116	rs2968485	0.545240517	0.523692308	0.571349862
rs7238698	0.558561983	0.515384615	0.578512397	rs10739816	0.545239246	0.523076923	0.574655647
rs11265519	0.558542064	0.586153846	0.539393939	rs7656677	0.545221445	0.641846154	0.449035813
rs743019	0.55853825	0.465538462	0.584022039	rs10816459	0.545203221	0.686769231	0.409917355
rs1421681	0.558535707	0.465846154	0.634710744	rs7910586	0.545199407	0.483384615	0.618181818
rs9953249	0.558515363	0.594461538	0.409366391	rs9397338	0.545196864	0.62520769	0.465564738
rs2072960	0.558506039	0.423076923	0.660055096	rs96712	0.545192202	0.429846154	0.646280992
rs4463318	0.558501377	0.640615385	0.485399449	rs9526000	0.545191354	0.671076923	0.422589532
rs9873033	0.558482306	0.594153846	0.500826446	rs6031755	0.545191354	0.641846154	0.472589532
rs424232	0.558367451	0.609846154	0.468319559	rs1557183	0.54519093	0.461538462	0.634710744
rs778127	0.558366179	0.593846154	0.514049587	rs9558546	0.545184149	0.641230769	0.4345978
rs11107419	0.558331002	0.625230769	0.494214876	rs6755608	0.54514092	0.602153846	0.485399449
rs9789054	0.558298792	0.538461538	0.579063361	rs12644961	0.545138801	0.445230769	0.641322314
rs7157656	0.558219538	0.603384615	0.59553719	rs2669513	0.5451299	0.484	0.580716253
rs11150069	0.55821869	0.596615385	0.428650138	rs2839580	0.545121424	0.501538462	0.619834711

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs03679	0.558206823	0.493538462	0.614325069	rs4722071	0.545106167	0.531384615	0.542699725
rs782918	0.558170375	0.499384615	0.607162534	rs11212337	0.545101928	0.444615385	0.652892562
rs2252795	0.558165289	0.694769231	0.420385675	rs11604758	0.545099385	0.484307692	0.596694215
rs4233365	0.558147489	0.585846154	0.523415978	rs8005389	0.545076923	0.633846154	0.43415978
rs11882238	0.558143251	0.476923077	0.608264463	rs1110911	0.545076075	0.452615385	0.626957245
rs7804157	0.558137741	0.632615385	0.457300275	rs9401575	0.545073109	0.570769231	0.499724518
rs1462027	0.558127993	0.586461538	0.515702479	rs9514296	0.545047256	0.602769231	0.483195592
rs623876	0.558090273	0.632	0.472176309	rs17155270	0.545032422	0.539384615	0.554820937
rs6812055	0.558048739	0.548615385	0.530027548	rs4320749	0.544999788	0.415076923	0.683360882
rs2907334	0.558026277	0.484307692	0.637465565	rs41529847	0.544971816	0.436615385	0.650137741
rs6734275	0.558014834	0.495076923	0.60661157	rs9652942	0.544968426	0.555384615	0.52231405
rs11852815	0.557979657	0.585846154	0.53994904	rs317510	0.544930282	0.641230769	0.461157025
rs11775193	0.557957618	0.586769231	0.523415978	rs4627537	0.544926467	0.477230769	0.604407713
rs1390893	0.557927527	0.588615385	0.512396694	rs7315539	0.544922653	0.445846154	0.634710744
rs4448903	0.55791227	0.531692308	0.558126722	rs1579905	0.544915872	0.461230769	0.609917355
rs4833034	0.557906336	0.546153846	0.541046832	rs12829185	0.544906124	0.475384615	0.611570248
rs710377	0.557903793	0.593538462	0.49862259	rs2252523	0.544891714	0.594153846	0.480440771
rs3804622	0.557871583	0.536923077	0.494214876	rs519462	0.544889595	0.641230769	0.445179063
rs17443616	0.557794448	0.578461538	0.579476584	rs1859040	0.544863318	0.648	0.431404959
rs1447870	0.557780038	0.554153846	0.558677686	rs44848485	0.544848485	0.443076923	0.64738292
rs11235189	0.557778343	0.516	0.582369146	rs44845094	0.544845094	0.648	0.445179063
rs274627	0.557773257	0.695076923	0.43030303	rs7200813	0.544843823	0.476923077	0.607713499
rs9934007	0.557762238	0.492	0.596694215	rs9515613	0.544837889	0.601846154	0.462258953
rs12533855	0.557753338	0.540923077	0.548760331	rs4766687	0.544826446	0.470153846	0.614876033
rs6498142	0.557721975	0.462153846	0.640220386	rs6566222	0.544822208	0.642153846	0.441322314
rs1218831	0.557688493	0.498769231	0.598347107	rs2199487	0.544815003	0.54	0.554820937
rs7831651	0.55768595	0.633230769	0.450137741	rs17808461	0.544807798	0.499692308	0.575757576
rs10760689	0.557633821	0.642153846	0.45785124	rs4373399	0.544806327	0.624923077	0.452341598
rs1366733	0.557594829	0.655384615	0.44738292	rs7930788	0.544790422	0.453230769	0.64077135
rs7846606	0.557580843	0.624307692	0.477134986	rs2955737	0.544786184	0.590461538	0.453443526
rs12546419	0.557573638	0.467384615	0.636363636	rs17617721	0.544771774	0.605538462	0.404407713
rs2365669	0.557558381	0.648307692	0.412672176	rs10072866	0.544754397	0.608923077	0.436914601
rs786274	0.557552871	0.718769231	0.409917355	rs283566	0.54474804	0.538461538	0.478236915
rs2643217	0.557513456	0.515384615	0.604407713	rs7137096	0.544735749	0.656923077	0.430853994
rs7605657	0.557497351	0.436307692	0.636914601	rs8064352	0.544731087	0.611076923	0.47107438
rs266413	0.557482941	0.531076923	0.574655647	rs11603889	0.544706929	0.476307692	0.609366391
rs2238692	0.557482094	0.656923077	0.462809917	rs7654383	0.544698877	0.469230769	0.621487603
rs6500834	0.557459207	0.710769231	0.403856749	rs2059271	0.54468701	0.461230769	0.608264463
rs2379266	0.55741174	0.562461538	0.557575758	rs705998	0.544683619	0.563692308	0.521212121
rs11051756	0.557395635	0.547384615	0.572066612	rs12804325	0.544676838	0.529846154	0.559779614
rs9289218	0.557357067	0.608923077	0.507989881	rs7621056	0.544672176	0.507076923	0.58677686
rs868258	0.557353677	0.648615385	0.461157025	rs13143454	0.5446637	0.523692308	0.547107438

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs10774183	0.557311295	0.648307692	0.491460055	rs11111882	0.54457343	0.572	0.508539945
rs6481442	0.557301971	0.616615385	0.49807163	rs4831945	0.54454039	0.516923077	0.554269972
rs6782599	0.557279508	0.506769231	0.525068871	rs8142661	0.544639542	0.405230769	0.655096419
rs224125	0.557257046	0.556923077	0.555371901	rs326740	0.54453488	0.557846154	0.484297521
rs2113946	0.557198559	0.406461538	0.703030303	rs6460468	0.544631066	0.594461538	0.490909091
rs1265181	0.557162534	0.469538462	0.641873278	rs6953287	0.544626828	0.412923077	0.659504132
rs2066341	0.557147701	0.561846154	0.545454545	rs1548160	0.544625556	0.460307692	0.624793388
rs11733487	0.557133715	0.633230769	0.457300275	rs10485891	0.544585717	0.624307692	0.446280992
rs7938186	0.557112948	0.483692308	0.466666667	rs2342401	0.54548487	0.504923077	0.606060606
rs4334280	0.557072261	0.500615385	0.602754821	rs11058371	0.544576393	0.516307692	0.570247934
rs1545898	0.557068871	0.608615385	0.506336088	rs970797	0.544562407	0.401230769	0.605509642
rs236855	0.557029032	0.586153846	0.449586777	rs10841923	0.54454164	0.610153846	0.483195592
rs7078219	0.557012926	0.694769231	0.41322314	rs9292873	0.544530621	0.687076923	0.402203857
rs9875315	0.556998093	0.563076923	0.557575758	rs1450073	0.544517906	0.422153846	0.656198347
rs7552456	0.556987074	0.404923077	0.665013774	rs17112932	0.544509854	0.509230769	0.570798898
rs1937950	0.556977326	0.438461538	0.597796143	rs580209	0.544494596	0.671692308	0.413774105
rs6439655	0.556962068	0.438153846	0.649586777	rs10274367	0.544492901	0.632923077	0.446280992
rs187801	0.556961644	0.594461538	0.509090909	rs1974700	0.544486544	0.625230769	0.471625344
rs2209623	0.556945115	0.461230769	0.660055096	rs1810502	0.544476796	0.423384615	0.659504132
rs11081443	0.556930282	0.477230769	0.615426997	rs1939810	0.544472982	0.488615385	0.614325069
rs11765303	0.55689341	0.655692308	0.447933884	rs9890556	0.544463658	0.484307692	0.603305785
rs7546172	0.556887052	0.603384615	0.443526171	rs2862996	0.544459419	0.524307692	0.54600551
rs6128386	0.556877305	0.640615385	0.499173554	rs4313213	0.544455181	0.413538462	0.671625344
rs10519099	0.556867981	0.585538462	0.520661157	rs1028920	0.544444586	0.648615385	0.430853994
rs3931961	0.556782793	0.576923077	0.52231405	rs1583259	0.544430176	0.664615385	0.418732782
rs4548236	0.556728968	0.523384615	0.596143251	rs7647249	0.544423395	0.444307692	0.644628099
rs1498934	0.556727697	0.611384615	0.510192837	rs6025100	0.54441619	0.460923077	0.618732782
rs1789377	0.556726001	0.400307692	0.67768595	rs1168330	0.544414918	0.444923077	0.657300275
rs4459374	0.556632337	0.492615385	0.637465565	rs2755413	0.544406018	0.601846154	0.495867769
rs4810860	0.556614537	0.554153846	0.560881543	rs518556	0.544405594	0.593538462	0.445179063
rs11848300	0.556609027	0.408615385	0.655443526	rs13411031	0.544392032	0.657846154	0.436914601
rs7574426	0.556569612	0.541538462	0.571349862	rs11716910	0.544390337	0.484923077	0.582369146
rs2542612	0.556554779	0.46	0.631404959	rs4459584	0.544375927	0.445538462	0.641322314
rs1946384	0.556547574	0.489538462	0.586225895	rs8020218	0.544323797	0.609846154	0.476033058
rs7023221	0.556531045	0.493538462	0.614876033	rs4760636	0.544320831	0.649230769	0.448484848
rs7570390	0.556524687	0.507384615	0.6	rs923025	0.544305573	0.469846154	0.588429752
rs9416608	0.556448824	0.626153846	0.476594022	rs2800686	0.544296249	0.499692308	0.608815427
rs16950272	0.556399237	0.477230769	0.660606060	rs7320209	0.54428862	0.444307692	0.636914601
rs4937378	0.55638737	0.703076923	0.404958678	rs4870976	0.544276754	0.671384615	0.41046832
rs2291922	0.556384403	0.586769231	0.512947658	rs1368242	0.54424751	0.522769231	0.554269972
rs6482146	0.556350074	0.579384615	0.534435262	rs859202	0.544225895	0.593230769	0.516804408
rs702131	0.556292859	0.610153846	0.50137741	rs4233600	0.544221657	0.540923077	0.54214876

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs2275016	0.556282687	0.401538462	0.672727273	rs1279688	0.544219538	0.524615385	0.545454545
rs11637631	0.55626531	0.429538462	0.678787879	rs7306109	0.544207247	0.476	0.570798988
rs2125398	0.556211062	0.593846154	0.520661157	rs1382165	0.544190718	0.491692308	0.504132231
rs8028764	0.55620979	0.600923077	0.531129477	rs1452389	0.544185633	0.62520769	0.466115702
rs5748919	0.556209366	0.564	0.544903581	rs10792468	0.544174189	0.617846154	0.469421488
rs10346363	0.556177156	0.569846154	0.543801653	rs158773	0.544169527	0.475076923	0.597796143
rs685548	0.556171223	0.454461538	0.640220386	rs6917225	0.544166561	0.595384615	0.476584022
rs1400202	0.556169104	0.421538462	0.689256198	rs512342	0.544161051	0.484923077	0.609366391
rs1930687	0.556166561	0.443692308	0.632506887	rs1320665	0.544155541	0.532307692	0.566391185
rs12734731	0.556161475	0.547384615	0.563636364	rs8075406	0.544130536	0.586461538	0.478787879
rs11839700	0.556159356	0.605846154	0.485950413	rs485832	0.544120364	0.492615385	0.591735537
rs703277	0.556119093	0.687076923	0.419824711	rs4597906	0.544108074	0.446153846	0.619834711
rs9901757	0.556105107	0.458153846	0.601101928	rs1753579	0.544103836	0.546153846	0.573553719
rs804968	0.556102564	0.514461538	0.601652893	rs6013520	0.54410214	0.429538462	0.659504132
rs2312464	0.556097902	0.445230769	0.632506887	rs4685485	0.544091545	0.679076923	0.415426997
rs891035	0.556060606	0.586769231	0.516253444	rs6753046	0.544085188	0.423076923	0.611019284
rs2015372	0.556047044	0.674153846	0.439669421	rs377256	0.544073744	0.436	0.661707989
rs642407	0.556011443	0.580307692	0.456749311	rs9403498	0.544069082	0.577538462	0.536639118
rs27412	0.555980081	0.579076923	0.5322231405	rs6943595	0.544063149	0.453538462	0.63030303
rs4644181	0.555949989	0.452307692	0.642975207	rs626156	0.544055096	0.431692308	0.644628099
rs10193678	0.555910574	0.454153846	0.638953168	rs7046606	0.544039415	0.625230769	0.460055096
rs45114	0.555908879	0.585538462	0.51184573	rs7874443	0.544036872	0.530769231	0.539393939
rs2153110	0.555908455	0.562769231	0.538292011	rs485469	0.544035177	0.538461538	0.548760331
rs6965444	0.555898707	0.663692308	0.43415978	rs9297145	0.544030939	0.617538462	0.465564738
rs286588	0.555890231	0.438769231	0.659504132	rs4770755	0.544016529	0.506769231	0.551515152
rs9915378	0.555884298	0.516	0.581267218	rs6010861	0.54400551	0.656	0.426997245
rs272504	0.55586353	0.570153846	0.5184573	rs8029216	0.543977538	0.640307692	0.420385675
rs921720	0.555863107	0.444	0.651239669	rs7196953	0.543970757	0.570769231	0.508539945
rs6965295	0.555857597	0.521230769	0.400550964	rs1009170	0.543966518	0.515692308	0.554269972
rs1790313	0.555848273	0.680307692	0.425895317	rs200873	0.543957618	0.506769231	0.584573003
rs3190021	0.555844459	0.686461538	0.438016529	rs474829	0.543949989	0.532615385	0.529476584
rs9960	0.555825387	0.405846154	0.696969697	rs847396	0.543912693	0.428307692	0.662809917
rs4870981	0.555813944	0.421538462	0.67768595	rs7760476	0.543911846	0.444923077	0.64738292
rs11626893	0.555800805	0.648923077	0.462558953	rs2937545	0.543911846	0.672	0.403305785
rs10210419	0.555788515	0.648615385	0.466115702	rs932241	0.543909727	0.633230769	0.457300275
rs1357991	0.555749523	0.679384615	0.412121212	rs12357592	0.543908455	0.508923077	0.585674931
rs869286	0.555734266	0.438153846	0.669972452	rs13198677	0.543900826	0.438461538	0.649586777
rs10946444	0.555716465	0.466153846	0.54214876	rs7007235	0.54389786	0.525230769	0.560330579
rs11615777	0.555684255	0.671076923	0.436914601	rs3804945	0.543891079	0.64	0.430853994
rs1896974	0.555663912	0.611692308	0.489256198	rs12596008	0.543890655	0.490461538	0.578512397
rs1524760	0.555607544	0.632615385	0.476033058	rs7864598	0.543887688	0.419692308	0.425895317
rs1269342	0.55558932	0.453230769	0.654545455	rs8102662	0.543879212	0.404923077	0.659504132

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs11185393	0.55568976	0.642153846	0.414325069	rs7904670	0.543868616	0.624923077	0.449035813
rs12993761	0.55568553	0.414153846	0.684848485	rs10512080	0.543866073	0.663384615	0.415977961
rs7639226	0.555543971	0.679384615	0.433608815	rs4862122	0.543860564	0.442153846	0.599449036
rs6851231	0.555537614	0.625846154	0.488705234	rs9543763	0.543855054	0.602461538	0.45785124
rs10762543	0.555534647	0.508615385	0.607713499	rs10836236	0.543850816	0.461846154	0.632506887
rs2173196	0.555509642	0.438461538	0.644628099	rs7530588	0.543849968	0.600615385	0.482093664
rs231915	0.555503285	0.445846154	0.652341598	rs17151293	0.543846154	0.499076923	0.593939394
rs10007790	0.555500742	0.461230769	0.643526171	rs16877238	0.543839373	0.618461538	0.456749311
rs4573083	0.555487603	0.469230769	0.637465565	rs4764446	0.543838101	0.655692308	0.475344353
rs6673531	0.555478279	0.400307692	0.668319559	rs1335515	0.543832168	0.641230769	0.468870523
rs4704417	0.55547616	0.688	0.4723140496	rs4732776	0.543816487	0.604307692	0.401652893
rs13041126	0.555457936	0.601538462	0.521212121	rs2013594	0.543798686	0.608923077	0.451790634
rs2180955	0.555424454	0.694769231	0.407162534	rs1489129	0.543795719	0.570153846	0.510743802
rs7745131	0.555422335	0.548615385	0.584573003	rs1700392	0.543789786	0.617230769	0.47768595
rs10093110	0.555388006	0.468923077	0.684297521	rs2052413	0.543769866	0.664615385	0.411570248
rs4937339	0.555351134	0.655692308	0.446280992	rs9379248	0.543768595	0.525230769	0.580716253
rs800318	0.555348167	0.584923077	0.516804408	rs4662248	0.543757576	0.516923077	0.473829201
rs11052011	0.555343929	0.499384615	0.595449036	rs6848074	0.543755457	0.484	0.602754821
rs4516626	0.555305785	0.673538462	0.408264463	rs2775139	0.543752914	0.602153846	0.480991736
rs769960	0.555278661	0.554769231	0.55261708	rs6757928	0.543718584	0.524923077	0.556473829
rs7934351	0.555252384	0.554153846	0.545454545	rs11577034	0.543680017	0.640923077	0.450137741
rs7127986	0.555237127	0.554461538	0.54600551	rs718617	0.543672388	0.522789231	0.565289256
rs41514548	0.555232888	0.468307692	0.635812672	rs4917817	0.54366815	0.491692308	0.544352617
rs9610510	0.55521085	0.602461538	0.504683196	rs7554231	0.543657555	0.686461538	0.414876033
rs235665	0.555206188	0.484307692	0.607713499	rs1423904	0.543656283	0.485230769	0.58953168
rs11978979	0.555178216	0.633538462	0.476033058	rs936750	0.543654588	0.429846154	0.653443526
rs1408953	0.55515834	0.561538462	0.534435262	rs1153589	0.543649078	0.433846154	0.628650138
rs10064772	0.555133291	0.672	0.416528926	rs492083	0.543638907	0.430153846	0.642424242
rs2082412	0.555129477	0.485538462	0.628099174	rs2766533	0.543634245	0.442769231	0.580716253
rs878263	0.555093809	0.479384615	0.564738292	rs10050439	0.543634245	0.508307692	0.581267218
rs4875727	0.55509769	0.577846154	0.507438017	rs7501420	0.543632973	0.585846154	0.480440771
rs1448963	0.55508879	0.553230769	0.548760331	rs10906317	0.543631278	0.655076923	0.426446281
rs673693	0.555063785	0.493538462	0.607162534	rs1721823	0.54362704	0.664307692	0.407713499
rs260452	0.555062089	0.473230769	0.601101928	rs10509199	0.543621954	0.485230769	0.57630854
rs10774840	0.555029879	0.679692308	0.422589532	rs17060577	0.543591439	0.413230769	0.672176309
rs6105944	0.555014198	0.639384615	0.462809917	rs1207692	0.543586353	0.633230769	0.444628099
rs716417	0.55499004	0.577538462	0.544352617	rs1018109	0.543558805	0.476307692	0.588429752
rs1663517	0.554978478	0.423076923	0.656749311	rs8032013	0.543554567	0.635358462	0.440220386
rs1576938	0.554975478	0.64	0.473829201	rs10176316	0.543526171	0.616	0.473829201
rs10792757	0.554947235	0.421846154	0.695316804	rs17178783	0.543522356	0.655692308	0.435261708
rs1365461	0.554938334	0.680615385	0.404407713	rs9518182	0.543515999	0.603384615	0.424793388
rs3829881	0.554932401	0.481846154	0.580165289	rs6465273	0.543509642	0.655692308	0.44077135

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs1813526	0.554909091	0.655692308	0.49035813	rs2013600	0.543495656	0.687692308	0.400550964
rs3800775	0.554909091	0.453846154	0.650137741	rs965974	0.543491418	0.507076923	0.579614325
rs11239821	0.554887052	0.561230769	0.510192837	rs2300937	0.54345285	0.603692308	0.452892562
rs4245150	0.554875185	0.483692308	0.626997245	rs2489720	0.543438017	0.5	0.579614325
rs7210514	0.554865861	0.665538462	0.440220386	rs1544028	0.543427845	0.484615385	0.595041322
rs450587	0.554843823	0.468615385	0.630853994	rs10470321	0.543421911	0.502769231	0.517355372
rs17084983	0.554822632	0.523076923	0.574104683	rs79880	0.543421911	0.429230769	0.63415978
rs1866524	0.554801017	0.500615385	0.61707989	rs342214	0.543417249	0.648615385	0.44077135
rs4797242	0.554754821	0.601538462	0.498071625	rs9616730	0.543409197	0.688	0.400550964
rs289027	0.554752702	0.546461538	0.562534435	rs304051	0.54340284	0.46	0.60407713
rs7077100	0.554730239	0.562769231	0.560330579	rs12568559	0.543400297	0.491384615	0.591184573
rs2584468	0.55472812	0.585538462	0.47768595	rs4758441	0.543386311	0.532	0.552066116
rs2164576	0.554726001	0.470461538	0.642975207	rs2023718	0.543367663	0.539076923	0.555371901
rs2459602	0.554711591	0.679076923	0.42369146	rs6862758	0.543349015	0.516615385	0.558126722
rs12997174	0.554685739	0.413230769	0.679338843	rs9311026	0.543337995	0.515692308	0.581818182
rs4568144	0.554668786	0.658769231	0.415426997	rs181652	0.543326552	0.664923077	0.400550964
rs804445	0.554629371	0.577230769	0.517906336	rs41329345	0.543320819	0.421538462	0.66446281
rs920483	0.554593346	0.474153846	0.56969697	rs882731	0.543320195	0.461538462	0.637465565
rs465318	0.554589108	0.525230769	0.595041322	rs1348232	0.543310447	0.528307692	0.51184573
rs2998367	0.55454164	0.594461538	0.427548209	rs488429	0.543290951	0.429846154	0.65785124
rs3010055	0.554508582	0.484615385	0.607162534	rs541945	0.543276965	0.551076923	0.504132231
rs6923826	0.554443738	0.5	0.573553719	rs3218791	0.543263403	0.452615385	0.631404959
rs1809181	0.554400932	0.624923077	0.477134986	rs3117229	0.543259165	0.648615385	0.452341598
rs12938215	0.554392456	0.655384615	0.442975207	rs2824981	0.543258317	0.545846154	0.537741047
rs10774863	0.554386099	0.586461538	0.533333333	rs6505780	0.543254503	0.672	0.415426997
rs1424372	0.554386099	0.515384615	0.44738292	rs1328758	0.543246874	0.444615385	0.641873278
rs4559153	0.554380165	0.556	0.555922865	rs13001751	0.543240941	0.522769231	0.56584022
rs11630094	0.554342445	0.546461538	0.559779614	rs12821085	0.543236279	0.525846154	0.40407713
rs1766059	0.554304726	0.656923077	0.428099174	rs1516655	0.54323416	0.405846154	0.687052342
rs813188	0.554302606	0.416615385	0.679889807	rs6563501	0.543231193	0.5	0.584022039
rs4943777	0.554291163	0.624615385	0.480991736	rs4897574	0.543226955	0.656	0.435812672
rs10504705	0.55427572	0.429538462	0.691460055	rs2164560	0.543218478	0.578769231	0.472727273
rs4240495	0.554266582	0.548307692	0.538842975	rs11534064	0.543196864	0.602153846	0.456198347
rs1144283	0.554261072	0.483692308	0.604958678	rs4872343	0.543191354	0.600307692	0.45785124
rs9999484	0.554251748	0.656615385	0.412121212	rs9807802	0.543185844	0.639076923	0.429201102
rs2282367	0.554246662	0.63230769	0.460055096	rs10224342	0.543173554	0.407076923	0.59338843
rs17013965	0.55424412	0.531076923	0.561983471	rs1245910	0.543171911	0.554461538	0.491460055
rs328506	0.554207671	0.593230769	0.519559229	rs11027288	0.543145158	0.438461538	0.642975207
rs11827486	0.554203857	0.412923077	0.691460055	rs16955101	0.543128205	0.624	0.460055096
rs7983427	0.554190718	0.509230769	0.515151515	rs6493620	0.543114643	0.601230769	0.4815427
rs1861373	0.554183513	0.525538462	0.519008264	rs6049096	0.543109133	0.522769231	0.54600551
rs6478204	0.554176732	0.663384615	0.409917355	rs17089365	0.543107014	0.483384615	0.611570248

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs1.106945	0.554167832	0.608923077	0.47768595	rs2017941	0.543101081	0.414153846	0.672176309
rs13194498	0.554156813	0.569538462	0.525619835	rs11159175	0.543068447	0.609538462	0.444628099
rs10736136	0.554154694	0.485230769	0.615426997	rs6110206	0.543067175	0.578153846	0.507988981
rs3748060	0.554082221	0.468	0.632568887	rs11002535	0.543057094	0.461230769	0.626446281
rs6029516	0.554067811	0.404923077	0.663911846	rs3811873	0.543051918	0.679076923	0.408815427
rs883275	0.554060606	0.610461538	0.509641873	rs501378	0.543034965	0.476923077	0.58953168
rs2049128	0.554052977	0.469230769	0.63415978	rs1027275	0.543034541	0.611076923	0.488705234
rs41129967	0.554049587	0.530461538	0.554820937	rs7803642	0.543031574	0.663692308	0.407162534
rs4246740	0.554040263	0.436923077	0.667217631	rs17364645	0.543026912	0.577846154	0.48815427
rs8087137	0.554037296	0.625538462	0.471625344	rs6944259	0.543020131	0.522461538	0.517906336
rs2165418	0.554008476	0.531692308	0.570247934	rs3731843	0.543018012	0.617846154	0.435647383
rs6571925	0.554003814	0.592923077	0.508539945	rs4686203	0.543013774	0.602769231	0.467217631
rs2076188	0.554	0.670769231	0.439669421	rs11135768	0.543013774	0.468307692	0.611570248
rs11052241	0.553963128	0.531076923	0.577961433	rs11708888	0.543013774	0.624923077	0.466666667
rs11038993	0.553956347	0.492307692	0.580165289	rs16957962	0.543002755	0.618153846	0.496418733
rs10118031	0.553950413	0.492	0.596143251	rs9957668	0.543002331	0.501230769	0.598898072
rs2570558	0.553943208	0.586153846	0.516804408	rs1438145	0.542982835	0.531692308	0.556473829
rs330109	0.553923289	0.534461538	0.508539945	rs2192838	0.542962068	0.581230769	0.412672176
rs754296	0.553918627	0.673230769	0.431955923	rs13099515	0.542944692	0.421846154	0.668870523
rs11735551	0.553888112	0.516307692	0.611019284	rs2581790	0.54293452	0.517230769	0.487603306
rs17769363	0.553858868	0.483384615	0.612672176	rs1040271	0.542915024	0.615076923	0.456198347
rs10751506	0.55380801	0.640923077	0.461157025	rs6540992	0.542911634	0.499076923	0.584022039
rs10770847	0.5537936	0.458769231	0.596143251	rs6416796	0.542895953	0.413846154	0.59490358
rs11833867	0.553751218	0.625846154	0.489256198	rs1890035	0.542854418	0.518153846	0.580165289
rs6543570	0.553747828	0.554461538	0.536639118	rs215529	0.542853994	0.436923077	0.635261708
rs2037816	0.553736385	0.701230769	0.412672176	rs5997228	0.542851875	0.640307692	0.436914601
rs10514200	0.553725366	0.624615385	0.485950413	rs2825684	0.54284467	0.593230769	0.453443526
rs1283390	0.553716889	0.420307692	0.69476584	rs500467	0.542837042	0.547384615	0.531129477
rs10994437	0.55371138	0.538153846	0.568595041	rs2727330	0.542827294	0.641538462	0.428099174
rs12812172	0.553703751	0.523076923	0.580165289	rs1230094	0.542825175	0.577538462	0.513498623
rs12152254	0.553633821	0.514769231	0.606060606	rs4751013	0.542820089	0.466153846	0.56584022
rs464457	0.55358381	0.421230769	0.680991736	rs7962583	0.54281246	0.649230769	0.430853994
rs11748914	0.553548633	0.654461538	0.427548209	rs1086047	0.542790846	0.523384615	0.571900826
rs6901053	0.553514728	0.491692308	0.626446281	rs2929726	0.542773893	0.610153846	0.432506887
rs7406717	0.553485908	0.563076923	0.526170799	rs984802	0.542767535	0.664923077	0.405509642
rs2223252	0.553454969	0.671076923	0.420385675	rs6043283	0.542765416	0.468923077	0.630853994
rs5018958	0.553453698	0.602153846	0.494214876	rs7105777	0.542753973	0.462461538	0.607162534
rs7938467	0.553450307	0.438769231	0.661157025	rs2996060	0.542750583	0.517538462	0.548209366
rs7021930	0.553426573	0.524	0.564187328	rs17106478	0.542746768	0.500615385	0.585123967
rs6573648	0.553413011	0.561538462	0.548209366	rs936012	0.542746345	0.438769231	0.644077135
rs6482066	0.553411164	0.481538462	0.579614325	rs11054800	0.542745921	0.547076923	0.538842975
rs4034	0.553405806	0.640615385	0.453443526	rs2343231	0.542725154	0.436615385	0.649586777

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs1045539	0.55338277	0.458461538	0.568595041	rs12899490	0.542693791	0.482153846	0.615977961
rs32489	0.553367663	0.586153846	0.521212121	rs890067	0.542680653	0.406769231	0.646280992
rs6078285	0.55335622	0.493230769	0.600550964	rs1572400	0.542669633	0.593846154	0.508539945
rs9522321	0.553351981	0.54	0.563636364	rs10063810	0.54266709	0.445846154	0.618181818
rs2414003	0.553318924	0.548	0.554269972	rs9291428	0.542647595	0.577230769	0.517355372
rs1930069	0.553318924	0.681230769	0.400550964	rs4527728	0.542645476	0.447076923	0.602203857
rs7159120	0.553300699	0.632307692	0.465013774	rs10882097	0.542638271	0.618153846	0.489256198
rs12902848	0.553297309	0.484615385	0.608264463	rs2720721	0.542631914	0.476	0.627548209
rs1679806	0.553291799	0.640615385	0.448484848	rs2207549	0.542626828	0.517538462	0.534986226
rs7299940	0.553290528	0.414461538	0.6815427	rs1264704	0.542610723	0.586461538	0.508539945
rs12414475	0.553290528	0.483692308	0.628650138	rs652451	0.542609875	0.578461538	0.50523416
rs35804423	0.553278237	0.444923077	0.667217631	rs255561	0.542585293	0.414769231	0.583471074
rs6038317	0.553268065	0.456923077	0.596694215	rs2365259	0.542578936	0.549538462	0.490909091
rs17821405	0.553242212	0.515692308	0.558677686	rs6733823	0.542571307	0.489230769	0.570798898
rs17060468	0.55322865	0.703076923	0.409917355	rs8037962	0.542564103	0.437230769	0.635812672
rs2720010	0.553207035	0.453538462	0.64738292	rs12145963	0.542552659	0.484923077	0.583471074
rs4776123	0.553191354	0.516	0.575757576	rs2737191	0.542543335	0.563076923	0.522865014
rs7863054	0.55315321	0.650153846	0.458402204	rs8061801	0.542514092	0.553846154	0.519559229
rs12542047	0.553137953	0.672307692	0.424242424	rs2173074	0.54250053	0.476615385	0.60661157
rs6926482	0.553121424	0.491384615	0.592286501	rs10748349	0.542498411	0.643384615	0.421487603
rs2334282	0.553101081	0.577230769	0.533834298	rs7493522	0.542483153	0.658153846	0.407713499
rs10793169	0.553091604	0.468	0.638016529	rs2076655	0.542474253	0.562461538	0.502479339
rs1416390	0.553090061	0.429538462	0.67768595	rs4693266	0.542454334	0.507076923	0.576859504
rs888821	0.553089214	0.578461538	0.50523416	rs591062	0.542446281	0.476	0.601652893
rs1149648	0.55308328	0.485846154	0.577410468	rs300970	0.54244501	0.509846154	0.479338843
rs11206244	0.553066751	0.545538462	0.532231405	rs447986	0.542444586	0.489538462	0.549862259
rs6734368	0.553047256	0.438461538	0.665013774	rs4951147	0.542438652	0.617538462	0.448484848
rs9855092	0.553037932	0.484923077	0.612672176	rs17079281	0.542415766	0.483692308	0.592286501
rs1244359	0.552996398	0.641230769	0.450688705	rs11583969	0.542407713	0.438461538	0.561432507
rs2793701	0.552990464	0.671692308	0.429201102	rs8123577	0.542404323	0.656615385	0.422038567
rs2939739	0.552930706	0.546153846	0.559779614	rs11248332	0.542402628	0.581846154	0.479889807
rs3779127	0.552925196	0.664	0.409366391	rs4626910	0.542398813	0.672307692	0.408264463
rs17110099	0.552924348	0.617846154	0.469972452	rs9527815	0.54239288	0.544615385	0.48429521
rs954580	0.552913329	0.633538462	0.455647383	rs1676235	0.54238737	0.679692308	0.401652893
rs8017720	0.552909515	0.462769231	0.650688705	rs699517	0.542385675	0.542385675	0.52892562
rs10166894	0.552908667	0.477846154	0.602754821	rs11096623	0.542383737	0.465953846	0.605509642
rs154483	0.552854418	0.484	0.601652893	rs129423	0.54237847	0.422153846	0.658402204
rs10511813	0.552828142	0.703076923	0.405509642	rs2225623	0.542377199	0.461538462	0.639118457
rs4659339	0.552797203	0.496923077	0.591735537	rs4686671	0.542354736	0.609846154	0.46464281
rs1387754	0.552767112	0.422153846	0.678787879	rs634554	0.542353041	0.585538462	0.503030303
rs1919484	0.552766688	0.624923077	0.465013774	rs11770967	0.54234626	0.412	0.597245179
rs7581601	0.552759907	0.563307692	0.527272727	rs3757340	0.542339479	0.579076923	0.495867769

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs1559677	0.552757364	0.588923077	0.429201102	rs2050616	0.542355241	0.526769231	0.521763085
rs2797180	0.552739987	0.423076923	0.662258953	rs4702966	0.542332698	0.502153846	0.575757576
rs974671	0.552739987	0.466461538	0.627548209	rs11096905	0.542327188	0.406153846	0.661157025
rs4074748	0.552734901	0.545846154	0.579063361	rs9545074	0.542321678	0.617230769	0.467217631
rs11080490	0.552730239	0.624615385	0.468319559	rs1949133	0.542317016	0.538769231	0.55261708
rs11906748	0.55270142	0.617230769	0.467217631	rs1160031	0.542308964	0.438153846	0.609366391
rs10514202	0.552679381	0.586461538	0.527227272	rs4308106	0.542300064	0.635692308	0.423140496
rs1470627	0.552676838	0.476615385	0.619283747	rs10507218	0.542292435	0.461538462	0.61046832
rs2838738	0.552648019	0.54	0.491460055	rs111966212	0.542292011	0.507384615	0.567493113
rs10511576	0.552591227	0.525846154	0.51184573	rs7283473	0.542280144	0.579076923	0.483195592
rs6667450	0.552586141	0.570769231	0.528374656	rs7709626	0.54226531	0.4735538462	0.455096419
rs2940371	0.552536131	0.578153846	0.525068871	rs928844	0.542261072	0.501538462	0.597796143
rs7852014	0.552530197	0.408923077	0.69476584	rs1898605	0.542237338	0.556615385	0.41322314
rs10974944	0.552528078	0.506769231	0.570798898	rs7036106	0.542231829	0.491692308	0.600550964
rs1371864	0.552504344	0.467692308	0.635812672	rs11881904	0.542226743	0.633230769	0.450688705
rs1201110	0.552494172	0.562153846	0.538292011	rs11587512	0.542213605	0.561538462	0.510743802
rs4890575	0.552481882	0.656	0.442424242	rs10028441	0.542211909	0.616615385	0.512396694
rs7129128	0.55248061	0.563846154	0.533333333	rs1653748	0.542206823	0.401538462	0.61046832
rs279880	0.552463658	0.459692308	0.621487603	rs8090547	0.542164442	0.423076923	0.65399449
rs2427341	0.552459419	0.641230769	0.461157025	rs12474591	0.542162323	0.555692308	0.531129477
rs964279	0.552456029	0.632923077	0.471625344	rs5009567	0.542159356	0.437538462	0.64628099
rs565042	0.552443314	0.664615385	0.438016529	rs232569	0.54214876	0.616923077	0.453443526
rs405792	0.552376775	0.406461538	0.6815427	rs1446046	0.542142827	0.647384615	0.436363636
rs6750230	0.552372537	0.586461538	0.532231405	rs17102450	0.542102988	0.617230769	0.450688705
rs17081444	0.552350498	0.623076923	0.469421468	rs4542744	0.542101716	0.415384615	0.669421488
rs7777181	0.552344988	0.657230769	0.405509642	rs111141473	0.542100445	0.599384615	0.43030303
rs13106655	0.552327612	0.491076923	0.614325069	rs1957583	0.542097054	0.564307692	0.503581267
rs490570	0.552310659	0.579076923	0.485399449	rs9614202	0.542092392	0.594153846	0.489807163
rs1366765	0.552310235	0.561230769	0.531129477	rs6973953	0.542091969	0.461846154	0.616528926
rs1975145	0.552307268	0.476307692	0.634710744	rs1944331	0.542083916	0.641230769	0.409917355
rs7713338	0.552294978	0.625538462	0.476033058	rs1453240	0.542077135	0.570153846	0.512947658
rs391070	0.552290316	0.602461538	0.495867769	rs1157500	0.542074592	0.624923077	0.449586777
rs7297457	0.552283958	0.681538462	0.444628099	rs16838780	0.542067811	0.535692308	0.500826446
rs11696306	0.552277601	0.460923077	0.639669421	rs2018070	0.542052554	0.473846154	0.547658402
rs11258686	0.552235643	0.672923077	0.424793388	rs13422730	0.542041958	0.600307692	0.424242424
rs2002311	0.552233948	0.556	0.534986226	rs1030975	0.542027548	0.548923077	0.523966942
rs10815721	0.552233524	0.405538462	0.689256198	rs164448	0.542023734	0.515384615	0.547107438
rs1370480	0.552231405	0.515384615	0.591735537	rs17050395	0.542018848	0.446153846	0.631404959
rs4618346	0.552222505	0.501230769	0.581818182	rs543224	0.542018224	0.600615385	0.471625344
rs1535060	0.552222505	0.451076923	0.642424242	rs12434656	0.542010595	0.569538462	0.467217631
rs983649	0.552220386	0.414153846	0.677134986	rs949912	0.541988557	0.664923077	0.409366391
rs6921510	0.552208943	0.650769231	0.404958678	rs7960156	0.541972452	0.5	0.561983471

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs1062948	0.552189023	0.529230769	0.556126722	rs10519247	0.541967366	0.515076923	0.574104683
rs10910204	0.552164018	0.483384615	0.603856749	rs11505418	0.541961009	0.413846154	0.667768595
rs2016840	0.552159356	0.594461538	0.505785124	rs7546194	0.54192117	0.649846154	0.415977961
rs668596	0.552154694	0.476615385	0.600550964	rs12586410	0.541907608	0.430461538	0.656749311
rs4714899	0.552151727	0.539076923	0.462809917	rs6593217	0.541899131	0.610153846	0.472727273
rs2846581	0.552128841	0.610153846	0.478787879	rs4599356	0.541889383	0.434153846	0.620385675
rs3924649	0.552110617	0.576923077	0.534986226	rs10502069	0.541884298	0.648615385	0.436363636
rs10833748	0.552095359	0.672923077	0.418732782	rs4679621	0.541873278	0.443384615	0.60050964
rs312058	0.552077559	0.469230769	0.619283747	rs17791802	0.541864378	0.634153846	0.441873278
rs2110152	0.552056792	0.616923077	0.475482094	rs6928012	0.541855902	0.428923077	0.663911846
rs1635627	0.552039415	0.656307692	0.44738292	rs1686166	0.541850816	0.554461538	0.512396694
rs4756347	0.552020767	0.663384615	0.441873278	rs1671413	0.541846154	0.561230769	0.514049587
rs561970	0.55201441	0.461230769	0.63415978	rs4633574	0.541844035	0.602153846	0.462809917
rs701815	0.551963552	0.608615385	0.504683196	rs2111112	0.541805467	0.484	0.581818182
rs4451390	0.551960161	0.492	0.598898072	rs41801653	0.541801653	0.422153846	0.661515025
rs10851954	0.551957194	0.484923077	0.612121212	rs16937996	0.541792329	0.404923077	0.668319559
rs2975696	0.551947023	0.577538462	0.514600551	rs4741815	0.541787243	0.454153846	0.622038567
rs10502983	0.551942784	0.657230769	0.446831956	rs11047765	0.541785548	0.641538462	0.426446281
rs7142050	0.551872007	0.561846154	0.537190083	rs4144418	0.541781733	0.405538462	0.653443526
rs10196764	0.551866497	0.554769231	0.543801653	rs7100333	0.541774105	0.431384615	0.603305785
rs3915852	0.551855783	0.569846154	0.517906336	rs7933723	0.541772409	0.406153846	0.651790634
rs1429573	0.551852087	0.639384615	0.449035813	rs2025509	0.541755033	0.672	0.413774105
rs11632275	0.551838525	0.469538462	0.608264663	rs3858620	0.541736809	0.461230769	0.62369146
rs350753	0.551827082	0.452615385	0.638567493	rs3807531	0.541735113	0.625230769	0.446780992
rs10466270	0.551813096	0.514461538	0.501928375	rs11251508	0.541721551	0.490153846	0.544903581
rs1321	0.551768171	0.467692308	0.639118457	rs6453423	0.541719432	0.609538462	0.474931129
rs888342	0.551753761	0.640307692	0.468870523	rs6055922	0.54170926	0.444923077	0.652892562
rs10808167	0.551737656	0.459076923	0.583471074	rs12366859	0.541691036	0.540615385	0.546556474
rs4633561	0.551695698	0.664615385	0.438016529	rs9508485	0.541690189	0.602769231	0.496418733
rs6581515	0.551690612	0.697846154	0.409917355	rs7129430	0.541689341	0.546153846	0.521763085
rs953239	0.551683831	0.414461538	0.638567493	rs2830859	0.541685527	0.460615385	0.618181818
rs1508289	0.551683831	0.539692308	0.569146006	rs2038819	0.541680865	0.561846154	0.503030303
rs11972945	0.551669421	0.461846154	0.644628099	rs6767825	0.54167705	0.648307692	0.435812672
rs11900348	0.551628735	0.508307692	0.592286501	rs7737692	0.541675779	0.5	0.572451791
rs17652091	0.551620682	0.695692308	0.412121212	rs4275737	0.541672812	0.578461538	0.477134986
rs6360	0.551614749	0.626153846	0.469421488	rs6759648	0.54166815	0.461538462	0.607162534
rs4368032	0.551611782	0.601230769	0.463360882	rs4241213	0.541667726	0.500615385	0.561983471
rs2072691	0.551608273	0.516307692	0.551515152	rs2355087	0.541660097	0.485846154	0.560330579
rs17813652	0.551605849	0.610461538	0.491460055	rs6115865	0.541658402	0.532	0.537741047
rs1552549	0.551565162	0.572	0.408264463	rs10135795	0.541657131	0.405846154	0.6815427
rs10184532	0.551543971	0.555076923	0.538842975	rs640060	0.541633821	0.528	0.469421488
rs1436094	0.551534223	0.671384615	0.431955923	rs987724	0.54162704	0.429846154	0.658953168

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs13246428	0.551523628	0.577230769	0.517906336	rs4336756	0.541624497	0.632	0.460055096
rs9471595	0.551515152	0.405538462	0.689256198	rs9890937	0.541621954	0.515692308	0.598898072
rs7400076	0.551480822	0.664	0.444077135	rs2705608	0.541621106	0.499384615	0.590082645
rs10192834	0.551477008	0.491384615	0.595592287	rs12547784	0.541618987	0.406789231	0.677134986
rs2880781	0.55146175	0.656	0.444077135	rs7489480	0.541615173	0.664923077	0.415426997
rs358493	0.551460479	0.579076923	0.523415978	rs7111924	0.54161263	0.492615385	0.482646428
rs10502255	0.551460055	0.617230769	0.478236915	rs17170619	0.541595253	0.438153846	0.652892562
rs7827545	0.551456665	0.569230769	0.515151515	rs630849	0.54159271	0.457846154	0.585674931
rs2360667	0.551427421	0.437846154	0.6959504132	rs11617085	0.541592287	0.438461538	0.646831956
rs7269546	0.551417249	0.686769231	0.41046832	rs2675638	0.541592287	0.406461538	0.649035813
rs7918405	0.551403263	0.625230769	0.467217631	rs628118	0.54157491	0.529538462	0.555371901
rs1497427	0.551401144	0.594769231	0.492011019	rs9302698	0.541565586	0.476307692	0.604958678
rs12378835	0.551393939	0.632	0.469972452	rs9966360	0.541560924	0.538769231	0.55922865
rs8069452	0.551353253	0.480923077	0.579614325	rs803765	0.541532104	0.469230769	0.608264463
rs11950065	0.551341386	0.452615385	0.665564738	rs10162279	0.541521509	0.421538462	0.633057851
rs1048466	0.551336724	0.554153846	0.530578512	rs1152522	0.541518966	0.624923077	0.447933884
rs2811174	0.551312566	0.444923077	0.657300275	rs2722171	0.541506251	0.437538462	0.642975207
rs4970722	0.551310871	0.695384615	0.401652893	rs276219	0.541497351	0.522769231	0.554269972
rs6881733	0.551299004	0.422769231	0.646831956	rs2172289	0.541489722	0.515692308	0.54600551
rs7795448	0.55129519	0.547076923	0.549311295	rs7192086	0.541482094	0.492307692	0.584573003
rs3739034	0.551292223	0.460923077	0.644077135	rs10018606	0.541477855	0.514461538	0.55261708
rs4922231	0.551269337	0.540923077	0.560330579	rs2072115	0.541467684	0.515076923	0.560881543
rs9688110	0.551268913	0.540615385	0.533333333	rs1046896	0.541463446	0.617230769	0.444077135
rs9345216	0.551254927	0.617230769	0.484848485	rs12914158	0.541459707	0.616923077	0.462809917
rs511198	0.551240093	0.570153846	0.53553719	rs6537336	0.541453698	0.438153846	0.638016529
rs7267275	0.551222293	0.584615385	0.495724518	rs12588287	0.541453274	0.506461538	0.58292011
rs158688	0.551216783	0.531692308	0.570247934	rs17484600	0.54144946	0.593230769	0.467217631
rs155568	0.551198983	0.532307692	0.542699725	rs11054692	0.541433355	0.499692308	0.584573003
rs6798464	0.551180335	0.516615385	0.581818182	rs2325065	0.541413859	0.454461538	0.632506887
rs955196	0.551176944	0.579692308	0.529476584	rs11162830	0.541404959	0.466153846	0.574104683
rs6910043	0.551129053	0.695076923	0.406060606	rs4952810	0.541393516	0.430153846	0.621487603
rs16986718	0.551124391	0.617846154	0.478787879	rs1867073	0.541387158	0.624615385	0.48815427
rs1939710	0.551105743	0.516307692	0.564738292	rs17662089	0.541379106	0.588923077	0.495316804
rs938291	0.551082857	0.663384615	0.429752066	rs1601422	0.541375715	0.617230769	0.460606061
rs1343995	0.551064209	0.610769231	0.485950413	rs39182	0.541364696	0.5986153846	0.495867769
rs2432755	0.551034541	0.507384615	0.597245179	rs12521931	0.541363848	0.548307692	0.53553719
rs618897	0.551021403	0.608307692	0.488705234	rs6691548	0.541357915	0.485230769	0.493663912
rs7095891	0.550993431	0.687076923	0.407162534	rs3817494	0.5413363	0.506461538	0.525068871
rs494374	0.550990888	0.538461538	0.550413223	rs6912658	0.5413363	0.406461538	0.617630854
rs4618385	0.550984107	0.539692308	0.539944904	rs11734306	0.541327824	0.554461538	0.4815427
rs10274707	0.550964611	0.687384615	0.413774105	rs930167	0.541320195	0.608307692	0.452892562
rs10899449	0.550962068	0.648923077	0.416528926	rs8074770	0.541313838	0.571384615	0.479889807

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs6992513	0.550948082	0.540615385	0.509641187	rs1317625	0.541301123	0.656615385	0.419834711
rs7359525	0.550943844	0.539384615	0.496969697	rs2053954	0.541300275	0.595846154	0.506336088
rs1335017	0.550942573	0.571076923	0.54214876	rs4241112	0.541290528	0.616923077	0.441322314
rs11624936	0.550939182	0.592923077	0.492011019	rs23242	0.541282051	0.61384615	0.420936639
rs6801579	0.550936639	0.453846154	0.644628099	rs9298555	0.541269761	0.624923077	0.457300275
rs12518350	0.550927315	0.610153846	0.467768595	rs1431114	0.541249417	0.647384615	0.438016529
rs4128913	0.550917567	0.460923077	0.634710744	rs7117375	0.541244331	0.500307692	0.566942149
rs208418	0.550913329	0.671076923	0.411570248	rs3845743	0.541230345	0.524	0.572451791
rs1079748	0.550910786	0.634769231	0.443526171	rs1544355	0.541229922	0.515076923	0.564187328
rs1560061	0.550895105	0.522153846	0.578512397	rs6437794	0.541225683	0.642461538	0.418181818
rs6531130	0.550879424	0.508	0.595041322	rs6417338	0.54121085	0.632615385	0.430853994
rs32198	0.55086459	0.575692308	0.480440771	rs2315625	0.541210002	0.515692308	0.572451791
rs4077685	0.550848061	0.687384615	0.407713499	rs12448859	0.541200254	0.626769231	0.463911846
rs7110439	0.550833651	0.626461538	0.452892562	rs1909475	0.541192202	0.563384615	0.520110193
rs12204218	0.55083238	0.633538462	0.466666667	rs754100	0.541153634	0.657230769	0.413774105
rs11249045	0.550823056	0.641538462	0.466666667	rs551244	0.54114092	0.601538462	0.478787879
rs16958069	0.550814156	0.405846154	0.641873278	rs4143952	0.541140072	0.446153846	0.628099174
rs972671	0.55080907	0.405230769	0.644628099	rs1845430	0.541138377	0.532615385	0.520661157
rs3836501	0.550797627	0.656	0.432506887	rs4085921	0.541119729	0.499384615	0.567493113
rs771236	0.550794236	0.650769231	0.412121212	rs11259283	0.541116338	0.532692308	0.537741047
rs1921083	0.550761602	0.672923077	0.414876033	rs1548991	0.54110659	0.414461538	0.677134986
rs1584717	0.550726425	0.571076923	0.523415978	rs17330287	0.541106167	0.462153846	0.618732782
rs2177123	0.550717949	0.641230769	0.49035813	rs10787308	0.541101928	0.632615385	0.438016529
rs7312475	0.550673872	0.429538462	0.656749311	rs9878321	0.541095147	0.546769231	0.516804408
rs235770	0.550670905	0.500307692	0.588429752	rs11199742	0.541087519	0.617230769	0.463360882
rs362794	0.550636999	0.640615385	0.455096419	rs1424883	0.541081585	0.610153846	0.480440771
rs1014137	0.550622166	0.634769231	0.428650138	rs7738604	0.541072261	0.578461538	0.492561983
rs9809784	0.55061708	0.464	0.528374656	rs654894	0.541064832	0.422769231	0.656749311
rs4765033	0.550613689	0.523692308	0.554269972	rs4645434	0.541061242	0.468	0.592837466
rs7902136	0.55060267	0.476615385	0.626446281	rs7152965	0.54105937	0.665538462	0.429752066
rs7629156	0.550594194	0.531692308	0.586225895	rs2237268	0.54104768	0.64	0.476446281
rs7846875	0.55059377	0.525846154	0.580165289	rs4406360	0.541029879	0.459384615	0.608264463
rs765238	0.55055605	0.630461538	0.449035813	rs7973207	0.541023522	0.588307692	0.480991736
rs10869758	0.55054715	0.695076923	0.405509642	rs3103731	0.541019784	0.452307692	0.627548209
rs750597	0.550537826	0.579384615	0.487603306	rs6488805	0.541011231	0.647384615	0.431955923
rs11886398	0.550492901	0.648615385	0.414325069	rs10431863	0.541004874	0.647076923	0.47107438
rs2240398	0.55049163	0.563384615	0.526170799	rs2036417	0.541002331	0.460307692	0.616528926
rs6069909	0.550488668	0.570461538	0.512336694	rs4932221	0.540999788	0.635338462	0.465013774
rs4875388	0.550461538	0.563384615	0.53553719	rs716829	0.540997669	0.579692308	0.431404959
rs7865592	0.550447976	0.428923077	0.672727273	rs11240341	0.540985802	0.477230769	0.585123967
rs10879959	0.550435262	0.664307692	0.439669421	rs10759254	0.540985378	0.413538462	0.67107438
rs10833511	0.550428481	0.687692308	0.411570248	rs7570469	0.540984954	0.632	0.438016529

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs7076937	0.550416614	0.624923077	0.47768595	rs603745	0.540963764	0.523692308	0.557024793
rs20832253	0.55040178	0.406769231	0.687036664	rs4674980	0.540960373	0.428307692	0.643526171
rs73132235	0.550391185	0.656307692	0.445730028	rs12731852	0.540958254	0.593846154	0.476033058
rs12915009	0.5503884827	0.546461538	0.525619885	rs16893344	0.540945539	0.624615385	0.424793388
rs436282	0.550382708	0.547384615	0.5344435262	rs6709176	0.540939606	0.490769231	0.590082645
rs954237	0.550376351	0.523692308	0.567493113	rs13180	0.540937063	0.612923077	0.429752066
rs17002700	0.55037296	0.405230769	0.696969697	rs7338828	0.540920534	0.405538462	0.651790634
rs3893060	0.550365332	0.579692308	0.467217631	rs2160321	0.540917143	0.453230769	0.626997245
rs2417848	0.550343717	0.455384615	0.621487603	rs2333423	0.540917143	0.514461538	0.54600551
rs297012	0.550342022	0.656307692	0.435812672	rs1403728	0.540910362	0.438769231	0.63195523
rs7801794	0.550312778	0.469538462	0.622038567	rs9861286	0.540899343	0.672307692	0.415426997
rs20455	0.55031193	0.484615385	0.613774105	rs473295	0.540895105	0.601538462	0.489807163
rs10865676	0.550301759	0.631692308	0.493663912	rs2070464	0.540878576	0.405230769	0.66446281
rs17437778	0.55029964	0.563076923	0.523415978	rs2692894	0.540868828	0.563076923	0.500275482
rs530008	0.550295825	0.420615385	0.661157025	rs1885894	0.540865438	0.655692308	0.418181818
rs7951928	0.550258953	0.599692308	0.493112948	rs633724	0.540856537	0.625846154	0.402754821
rs11102782	0.550205128	0.688307692	0.401652893	rs1428669	0.540854418	0.523076923	0.550964187
rs2925337	0.550202161	0.413846154	0.688705234	rs11052229	0.540842128	0.602769231	0.498071625
rs7241641	0.550194957	0.516	0.576859504	rs4770603	0.540834499	0.664	0.415977961
rs1484032	0.550192837	0.672307692	0.444071735	rs1049687	0.540834075	0.414153846	0.665013774
rs747912	0.550172494	0.546461538	0.564187328	rs1408343	0.540833651	0.530153846	0.54214876
rs6668917	0.55015427	0.572	0.538292011	rs2561667	0.540825599	0.414461538	0.633608815
rs944868	0.550153422	0.625538462	0.483195592	rs333948	0.540824327	0.430769231	0.646280992
rs4984407	0.55014876	0.469538462	0.636363636	rs971626	0.540786607	0.483692308	0.568044077
rs7245865	0.550128417	0.492	0.61046832	rs7910587	0.54077135	0.476	0.6
rs230884	0.550125874	0.625230769	0.468319559	rs1585535	0.540760331	0.421230769	0.634710744
rs6496259	0.55011316	0.476923077	0.640220386	rs1631619	0.540757364	0.557846154	0.460606061
rs4689642	0.550102988	0.577538462	0.513438623	rs10498494	0.540742106	0.524923077	0.519008264
rs11042786	0.550100021	0.570461538	0.519559229	rs9908583	0.540732359	0.537538462	0.552782369
rs4745816	0.550097478	0.435384615	0.58677686	rs7076095	0.540725577	0.537538462	0.498071625
rs767755	0.550065268	0.569538462	0.539393939	rs1540052	0.540719644	0.508307692	0.571900826
rs2279605	0.55005552	0.469230769	0.627548209	rs4450898	0.540717101	0.547384615	0.524517906
rs1915960	0.550047044	0.464923077	0.597796143	rs1529570	0.540713711	0.406461538	0.672727273
rs17470134	0.550040687	0.64	0.450137741	rs459756	0.540712015	0.595538462	0.493663912
rs4812571	0.550036448	0.500307692	0.587878788	rs4334676	0.540704881	0.539384615	0.538842975
rs2157598	0.550036448	0.516615385	0.570738898	rs2271829	0.540702691	0.534153846	0.469972452
rs1724932	0.550030091	0.562769231	0.516804408	rs7415063	0.540698029	0.548923077	0.50137741
rs11845632	0.550025429	0.626461538	0.447933884	rs17333131	0.540688281	0.656	0.41322314
rs6702764	0.550021615	0.515692308	0.571349862	rs7331570	0.540674295	0.636923077	0.408264463
rs2807552	0.549995762	0.647692308	0.451239669	rs4873195	0.540667514	0.600307692	0.465013774
rs1544056	0.549993643	0.430769231	0.665564738	rs772908	0.5406637	0.508	0.564738292
rs976244	0.549978385	0.485846154	0.609366391	rs1420361	0.540661157	0.546153846	0.529476584

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs2560828	0.549973299	0.468923077	0.624793388	rs11010067	0.5405819	0.633230769	0.436363636
rs10862477	0.549922865	0.626461538	0.459504132	rs1004704	0.540635304	0.476615385	0.597796143
rs17689863	0.549906336	0.508615385	0.584573003	rs1401317	0.540627675	0.431076923	0.644628099
rs12525800	0.549875397	0.5	0.580165289	rs468080	0.540621318	0.438769231	0.631955923
rs2191323	0.549873702	0.624615385	0.461157025	rs11157655	0.540615385	0.500615385	0.579063361
rs6981569	0.549866014	0.648923077	0.444077135	rs400042	0.540613266	0.421230769	0.658402204
rs4932165	0.549826234	0.491692308	0.608815427	rs2796771	0.540607756	0.600615385	0.461707989
rs12456284	0.549812672	0.639692308	0.456138347	rs9641113	0.540602246	0.552	0.47107438
rs201017	0.549801229	0.522461538	0.571349862	rs1203651	0.540596737	0.408	0.615426997
rs10836643	0.549798686	0.546461538	0.537139083	rs11196939	0.540580631	0.408	0.563663664
rs4937732	0.549788515	0.437538462	0.665013774	rs10773456	0.54057936	0.522461538	0.553168044
rs4839412	0.549785972	0.712	0.402754821	rs2807822	0.540575546	0.454769231	0.626446281
rs769163	0.549768171	0.649230769	0.415977961	rs4611418	0.540565798	0.422153846	0.658953168
rs11861717	0.549762662	0.688307692	0.407713499	rs3818638	0.540564103	0.539076923	0.56969697
rs3779370	0.549757576	0.548	0.542699725	rs2156411	0.540555202	0.639692308	0.432506887
rs4867263	0.549747828	0.670769231	0.430853994	rs9882790	0.540538673	0.460307692	0.612672176
rs17063391	0.549743166	0.594461538	0.498071625	rs5010979	0.54052384	0.453230769	0.631955923
rs2390478	0.549739775	0.640615385	0.457306275	rs10869290	0.540523416	0.493384615	0.573002755
rs3937693	0.549728332	0.445846154	0.621487603	rs13076727	0.540516211	0.564923077	0.470523416
rs852499	0.54971477	0.444923077	0.638016529	rs13329545	0.540512397	0.586461538	0.479898807
rs16913638	0.549702479	0.539384615	0.526721763	rs1286814	0.540507311	0.539384615	0.519559229
rs9314727	0.549677705	0.593846154	0.502479339	rs12670377	0.540484001	0.516	0.549311295
rs12892711	0.549653316	0.438153846	0.640220386	rs731803	0.540483153	0.518769231	0.510743802
rs1925757	0.549618987	0.404923077	0.667768595	rs7161192	0.540460691	0.444	0.525068871
rs11990945	0.549618563	0.406153846	0.620936639	rs4145201	0.540449672	0.506461538	0.562534435
rs761793	0.549608815	0.679692308	0.434710744	rs6911630	0.540438228	0.578461538	0.494214876
rs6428686	0.549604153	0.601230769	0.47107438	rs8074097	0.540433399	0.569846154	0.493663912
rs3861076	0.549593558	0.492923077	0.597245179	rs10495337	0.540433399	0.608615385	0.453443526
rs440156	0.549591015	0.560615385	0.527823691	rs9318052	0.5404306	0.626153846	0.479338843
rs1476698	0.549581267	0.493538462	0.604958678	rs1388613	0.5404217	0.460923077	0.61322314
rs2769916	0.549575605	0.562769231	0.525619835	rs2368069	0.540414495	0.575692308	0.409917355
rs4573070	0.5495558381	0.525538462	0.560330579	rs2191926	0.540411952	0.509538462	0.529476584
rs9520691	0.549554567	0.507384615	0.571349862	rs1609474	0.540411528	0.452923077	0.601652893
rs4890980	0.549547362	0.468307692	0.648484848	rs732996	0.540409833	0.514461538	0.564187328
rs918480	0.549506675	0.413538462	0.685399449	rs1808529	0.540408561	0.548	0.517355372
rs6508899	0.549488451	0.632	0.426997245	rs72991	0.540390761	0.625230769	0.468319559
rs964403	0.549484213	0.461230769	0.598347107	rs17533457	0.540386523	0.429538462	0.639669421
rs2238090	0.549465141	0.648923077	0.422589532	rs6804595	0.540382284	0.476307692	0.593939394
rs7529292	0.549445645	0.444	0.65399449	rs10985529	0.54036957	0.64	0.426446281
rs233203	0.549435474	0.488615385	0.577410468	rs2792954	0.540369146	0.4832153846	0.556473829
rs2535415	0.549424878	0.480307692	0.551515152	rs874427	0.540368722	0.619076923	0.41046832
rs16831243	0.549409621	0.406461538	0.683746556	rs7709129	0.540367875	0.422153846	0.643526171

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs6478108	0.549386311	0.561230769	0.537190083	rs1317443	0.54035516	0.430461538	0.649586777
rs934778	0.549360034	0.536923077	0.538292011	rs566416	0.540352193	0.477538462	0.607713499
rs10898868	0.549333029	0.536923077	0.54214876	rs9544950	0.540350498	0.458153846	0.575206612
rs4815191	0.549333291	0.514461538	0.577961433	rs7038823	0.540336512	0.523692308	0.527272727
rs226456	0.549324433	0.569846154	0.534986226	rs10824478	0.540328459	0.641538462	0.438016529
rs11777456	0.549323162	0.571692308	0.53537719	rs10929965	0.54032634	0.533230769	0.499724518
rs17040762	0.549306209	0.646769231	0.44738292	rs2525743	0.54032634	0.483384615	0.58953168
rs1895207	0.549292223	0.477846154	0.614876033	rs12471726	0.540317864	0.545846154	0.502479339
rs28881	0.549281627	0.415076923	0.679338843	rs468308	0.540311083	0.664615385	0.424793388
rs918233	0.549276118	0.648615385	0.436363636	rs4608395	0.540310659	0.429846154	0.628099174
rs1236860	0.549274423	0.617230769	0.472176309	rs17831865	0.540308116	0.476615385	0.614325069
rs641778	0.549267218	0.453230769	0.635261708	rs7710314	0.540299216	0.438461538	0.631955923
rs2228375	0.549265099	0.564923077	0.495336804	rs11291881	0.540297097	0.456615385	0.588429752
rs7992054	0.549261284	0.617846154	0.482644628	rs4963924	0.540296249	0.593230769	0.440220386
rs4405339	0.549247722	0.662153846	0.443526171	rs10954412	0.540283535	0.559692308	0.427548209
rs670728	0.549244755	0.469538462	0.611019284	rs1072264	0.540282687	0.594769231	0.478236915
rs557564	0.549231617	0.509538462	0.58292011	rs1502641	0.540278025	0.428615385	0.64738292
rs4619182	0.549226531	0.523076923	0.562534435	rs4644590	0.540277177	0.469846154	0.597796143
rs3734165	0.549213816	0.465846154	0.595592287	rs7166692	0.540277939	0.489230769	0.575206612
rs7122693	0.549209578	0.429846154	0.640220386	rs4627442	0.540271668	0.436	0.650688705
rs4298326	0.54919644	0.578461538	0.528374656	rs136343	0.540269125	0.704	0.402754821
rs2562784	0.549185844	0.485230769	0.609917355	rs1188349	0.540268701	0.609230769	0.459504132
rs2082871	0.549163806	0.502153846	0.551515152	rs1833868	0.540246662	0.657230769	0.422038567
rs7181054	0.549128205	0.467384615	0.635812672	rs10744391	0.540239034	0.593230769	0.507988981
rs10173591	0.54912651	0.605846154	0.455647383	rs614968	0.540230981	0.476615385	0.547107438
rs11789818	0.549124815	0.672307692	0.438567493	rs208798	0.540222505	0.404923077	0.663911846
rs480700	0.549106167	0.413846154	0.651790634	rs6959745	0.540209366	0.657230769	0.429201102
rs10760969	0.549087095	0.687692308	0.40661157	rs953288	0.54019538	0.421538462	0.668319559
rs7145856	0.549078618	0.477846154	0.599449036	rs10449236	0.540193685	0.664	0.408815427
rs10764780	0.549077685	0.625538462	0.444628099	rs2905986	0.540170799	0.529230769	0.505785124
rs6946119	0.54903327	0.609846154	0.487052342	rs6136898	0.540168256	0.484	0.596143251
rs10904949	0.549001907	0.567384615	0.422589532	rs10062843	0.540157661	0.522461538	0.549311295
rs10256021	0.54900106	0.68	0.415426997	rs10777182	0.540156389	0.538461538	0.546556474
rs1232487	0.548988769	0.490769231	0.612121212	rs11070561	0.540141979	0.578153846	0.478787879
rs9911992	0.548953592	0.632923077	0.4950504132	rs10509316	0.540134774	0.445846154	0.640220386
rs4583067	0.548929434	0.562461538	0.496418733	rs4733071	0.540124603	0.624307692	0.462809917
rs1535548	0.548928587	0.461230769	0.645730028	rs4427133	0.54011655	0.640923077	0.445730028
rs1410364	0.548920534	0.562769231	0.554269972	rs974995	0.54011104	0.578153846	0.477134986
rs803413	0.548900191	0.515384615	0.585123967	rs11256439	0.540105107	0.609230769	0.471625344
rs16866695	0.548871795	0.593230769	0.50137741	rs137686	0.540099597	0.507384615	0.566942149
rs17763853	0.54885908	0.640615385	0.437485565	rs4895410	0.540097902	0.429846154	0.637465565
rs2158917	0.548842975	0.522769231	0.570798898	rs9582124	0.540091121	0.580923077	0.450137741

ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe	ILMNID	mean_cv_auc	mean_cv_sen	mean_cv_spe
rs487438	0.548839161	0.570461538	0.511294766	rs10805561	0.540091121	0.484307652	0.584573003
rs9805018	0.548837889	0.657230769	0.427548209	rs857424	0.54006442	0.499384615	0.573002755
rs10922605	0.548820937	0.601538462	0.487603306	rs6577097	0.540031786	0.554153846	0.509090909
rs10489022	0.548812884	0.562153846	0.523966942	rs683471	0.540018648	0.521846154	0.567493113
rs1886049	0.548809494	0.572307692	0.469421488	rs7173885	0.540010172	0.484615385	0.611019284
rs4812893	0.548801441	0.664923077	0.408815427	rs12316815	0.540005933	0.460923077	0.614325069
rs418692	0.548791541	0.507076923	0.577961433	rs8023364	0.540002543	0.523384615	0.558677686
				rs29664	0.540001695	0.671692308	0.402203857

WHAT IS CLAIMED IS:

1. A kit for determining methylation status of at least one CpG dinucleotide
5 and a genotype of at least one single-nucleotide polymorphism (SNP), the kit comprising:
at least one first nucleic acid primer at least 8 nucleotides in length that is
complementary to a bisulfite-converted nucleic acid sequence comprising a first CpG
dinucleotide at a GC locus selected from the group consisting of cg19693031,
cg17901584, cg06500161, cg13375427, and cg21139312 or at a second CpG dinucleotide
10 in linkage disequilibrium with the first CpG dinucleotide at a GC locus selected from the
group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and
cg21139312, wherein the linkage disequilibrium has a value of $R > 0.3$, wherein the at
least one first nucleic acid primer detects a methylated or unmethylated CpG
dinucleotide, and
15 at least one second nucleic acid primer at least 8 nucleotides in length that
is complementary to a DNA sequence or a bisulfite-converted DNA sequence of a first
SNP selected from the group consisting of rs1834967, rs41480347, rs2141638,
rs6684261, and rs1423855 or a second SNP in linkage disequilibrium with the first SNP
selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and
20 rs1423855, wherein the linkage disequilibrium has a value of $R > 0.3$.
2. The kit of claim 1, wherein the at least one first nucleic acid primer detects
the unmethylated CpG dinucleotide.
- 25 3. The kit of claim 1, wherein the at least one first nucleic acid primer detects
the methylated CpG dinucleotide.
4. The kit of any of claims 1-3, wherein the at least one first nucleic acid
primer comprises one or more nucleotide analogs.
30
5. The kit of any of claims 1-4, wherein the at least one first nucleic acid
primer comprises one or more synthetic or non-natural nucleotides.

6. The kit of any one of claims 1-5, further comprising a solid substrate to which the at least one first nucleic acid primer is bound.

7. The kit of claim 6, wherein the substrate is a polymer, glass,
5 semiconductor, paper, metal, gel or hydrogel.

8. The kit of claim 6, wherein the solid substrate is a microarray or microfluidics card.

10 9. The kit of any one of claims 1-8, further comprising a detectable label.

10. The kit of any one of claims 1-9, further comprising at least a third nucleic acid primer at least 8 nucleotides in length that is complementary to a nucleic acid sequence upstream of the CpG dinucleotide.

15

11. The kit of any one of claims 1-9, further comprising at least a third nucleic acid primer at least 8 nucleotides in length that is complementary to a nucleic acid sequence downstream of the CpG dinucleotide.

20 12. A method of determining the presence of biomarkers associated with diabetes in a biological sample from a subject, comprising:

(a) providing a first portion of the biological sample and a second portion of the biological sample, wherein the nucleic acid from at least the first portion is bisulfite converted;

25 (b) contacting the first portion of the biological sample with a first oligonucleotide primer at least 8 nucleotides in length that is complementary to a sequence that comprises a first CpG dinucleotide at a GC locus selected from the group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312, or a second CpG dinucleotide in linkage disequilibrium with the first CpG dinucleotide at a
30 GC locus selected from the group consisting of cg19693031, cg17901584, cg06500161, cg13375427, and cg21139312, wherein the linkage disequilibrium has a value of $R > 0.3$, wherein the first nucleic acid primer detects a methylated or unmethylated CpG dinucleotide; and

(c) contacting the second portion of the biological sample with a nucleic acid primer at least 8 nucleotides in length that is complementary to a DNA sequence or a bisulfite-converted DNA sequence of a first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855 or a second SNP in linkage disequilibrium with a first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855, wherein the linkage disequilibrium has a value of $R > 0.3$,

wherein the percentage of methylation of the CpG dinucleotide at the GC locus selected from the group consisting of cg06500161, cg21139312, cg19693031, cg17901584, and cg13375427 and the identity of the nucleotide in the first SNP selected from the group consisting of rs1834967, rs41480347, rs2141638, rs6684261, and rs1423855 or the second SNP in linkage disequilibrium with the first SNP are biomarkers associated with diabetes in the subject.

13. A method of determining the presence of a biomarker associated with diabetes in a subject sample, the method comprising:

(a) isolating nucleic acid sample from the subject sample,
 (b) performing a genotyping assay on a first portion of the nucleic acid sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain genotype data; and/or

(c) bisulfite converting the nucleic acid in a second portion of the nucleic acid and performing methylation assessment on a second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data; and

(d) entering the genotype data from step (b) and/or methylation data from step (c) into an algorithm that accounts for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect, wherein the algorithm is a machine learning algorithm capable of accounting for linear and non-linear effects.

30

14. The method of claim 13, wherein the at least one interaction effect is selected from the group consisting of a gene-environment interaction (SNPxCpG) effect,

a gene-gene interaction (SNP×SNP) effect, and an environment-environment interaction (CpG×CpG) effect.

15 15. The method of claim 14, wherein the at least one interaction effect is a gene-environment interaction effect (SNP×CpG) between a CpG site from Appendix A or a CpG site that is collinear ($R>0.3$) with a CpG site from Appendix A and a SNP from Appendix C or a SNP within moderate linkage disequilibrium ($R>0.3$) from a SNP from Appendix C.

10 16. The method of claim 14, wherein the at least one interaction effect is an environment-environment interaction effect (CpG×CpG) between at least two CpG sites from Appendix A.

15 17. The method of claim 16, wherein one or both of the at least two CpG sites are collinear ($R>0.3$) with one or both of the at least two CpG sites from Appendix A.

 18. The method of claim 14, wherein the at least one interaction effect is a gene-gene interaction effect (SNP×SNP) between at least two SNPs from Appendix C.

20 19. The method of claim 18, wherein one or both of the at least two SNPs are collinear ($R>0.3$) with one or both of the at least two SNPs from Appendix C.

 20. The method of any of claims 13-19, wherein the biological sample is a saliva sample.

25

 21. A system for determining methylation status of at least one CpG dinucleotide and a genotype of at least one single-nucleotide polymorphism (SNP), the system comprising:

 a nucleic acid isolation module configured to isolate a nucleic acid sample from a subject sample;

30 a genotyping assay module configured to perform a genotyping assay on a first portion of the nucleic acid sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage

disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain genotype data;

a methylation assay module configured to bisulfite convert the nucleic acid in a second portion of the nucleic acid and perform a methylation assessment on a second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data; and

an identification system configured to account for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on the genotype data from step (b) and/or methylation data from step (c).

22. The system of claim 21, wherein the algorithm is a machine learning algorithm capable of accounting for linear and non-linear effects.

23. The system of claim 21 or 22, further comprising an output module configured to provide an output based on an identification by the identification system, wherein the identification accounts for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on the genotype data from step (b) and/or methylation data from step (c).

24. A non-transitory computer-readable medium storing instructions executable by a processing device to perform operations comprising:

accounting for at least one SNP main effect and/or at least one CpG main effect and/or at least one interaction effect based on genotype data and/or methylation data, wherein:

the genotype data is based on a genotyping assay on a first portion of a nucleic acid sample isolated from a subject sample to detect the presence of at least one SNP, wherein the at least one SNP is a first SNP from Appendix C and/or is a second SNP in linkage disequilibrium ($R > 0.3$) with a first SNP from Appendix C to obtain the genotype data; and

the methylation data is based on a methylation assay on a bisulfite converted nucleic acid in a second portion of the nucleic acid sample to detect methylation status of at least one CpG site from Appendix A and/or a CpG site collinear ($R > 0.3$) with a CpG from Appendix A to obtain methylation data.

25. The non-transitory computer-readable medium of claim 24, wherein the operations further comprise providing an output based on the accounting.

- 5 26. The non-transitory computer-readable medium of claim 25, wherein the output comprises one or more of storing a report based on the accounting to another non-transitory computer-readable medium, modifying a display based on the accounting, triggering an audible alert based on the accounting, triggering a haptic or vibratory alert based on the accounting, triggering the printing of a report based on the accounting, or
- 10 triggering the delivery of a therapeutic based on the accounting.