

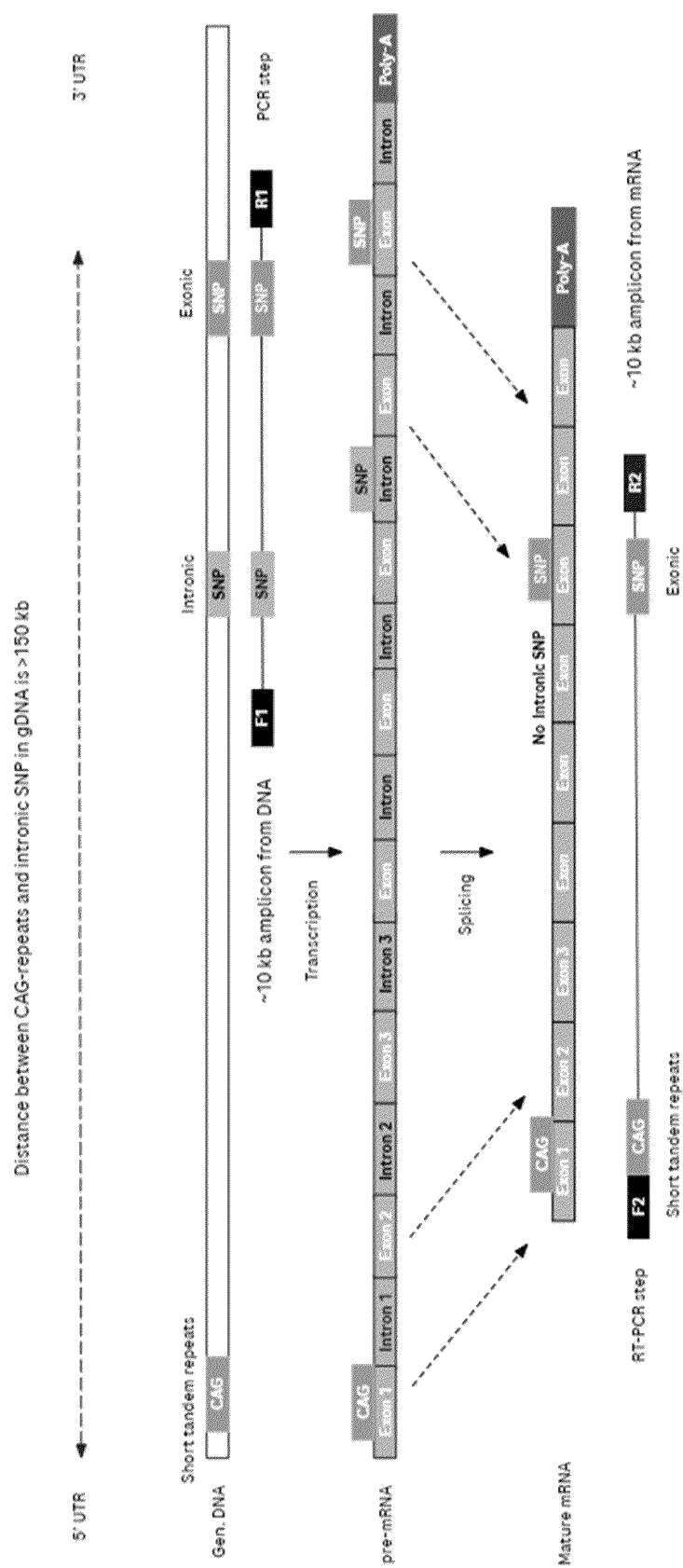


FIG. 1

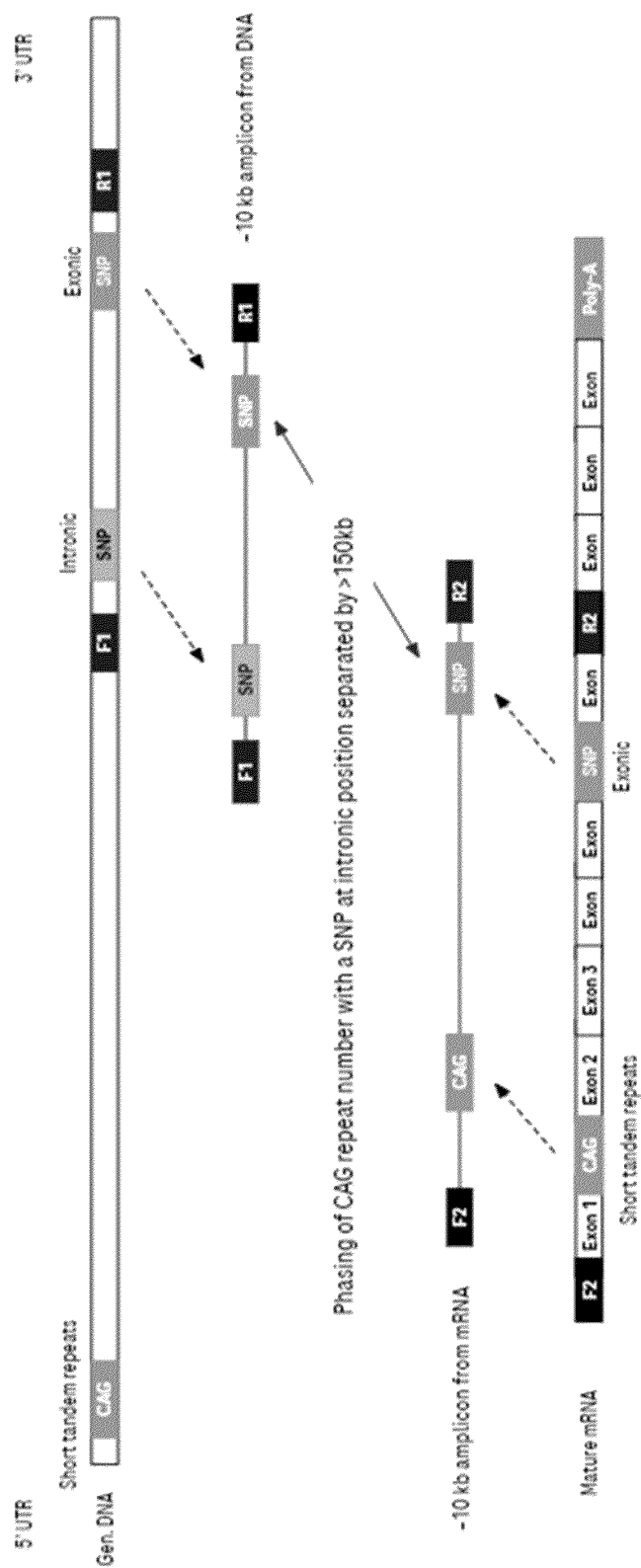


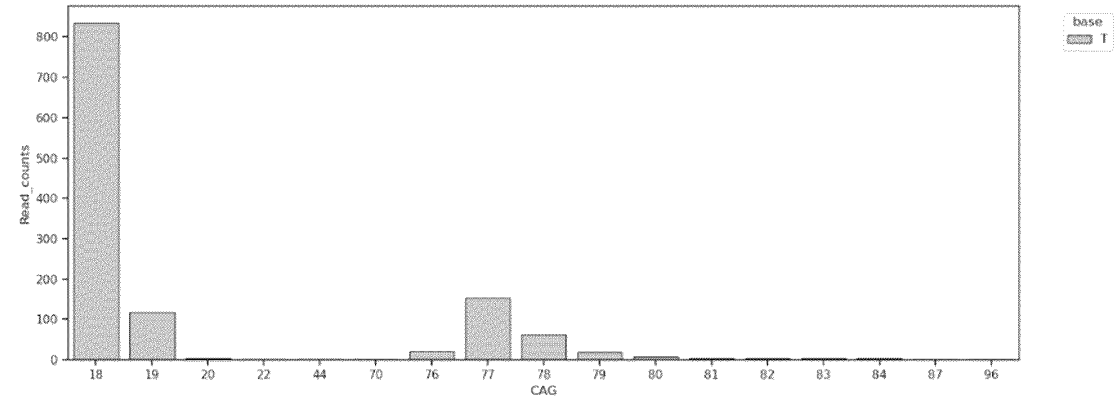
FIG. 2

FIG. 3

A

GM04282	Mean CAG repeat	Standard deviation	% reads	% Variant Exonic (C) SNP Allele	% Reference Exonic (T) SNP Allele	% Undefined
High CAG	77.63	2.88	22.74	0	95.25	4.75
Low CAG	18.13	0.36	77.26	0	95.51	4.49

B



C

GM04282	% Intronic (G) Allele	% Intronic (A) Allele
Exonic C Allele	NA	NA
Exonic T Allele	0	100

FIG.4

A

GM13503	Mean CAG repeat	Standard deviation	% reads	% Variant Exonic (C) SNP Allele	% Reference Exonic (T) SNP Allele	% Undefined
High CAG	46.29	0.6	48.14	0.7	99.3	0
Low CAG	18	0.08	51.86	99.67	0.16	0.16

FIG.4 (continued)

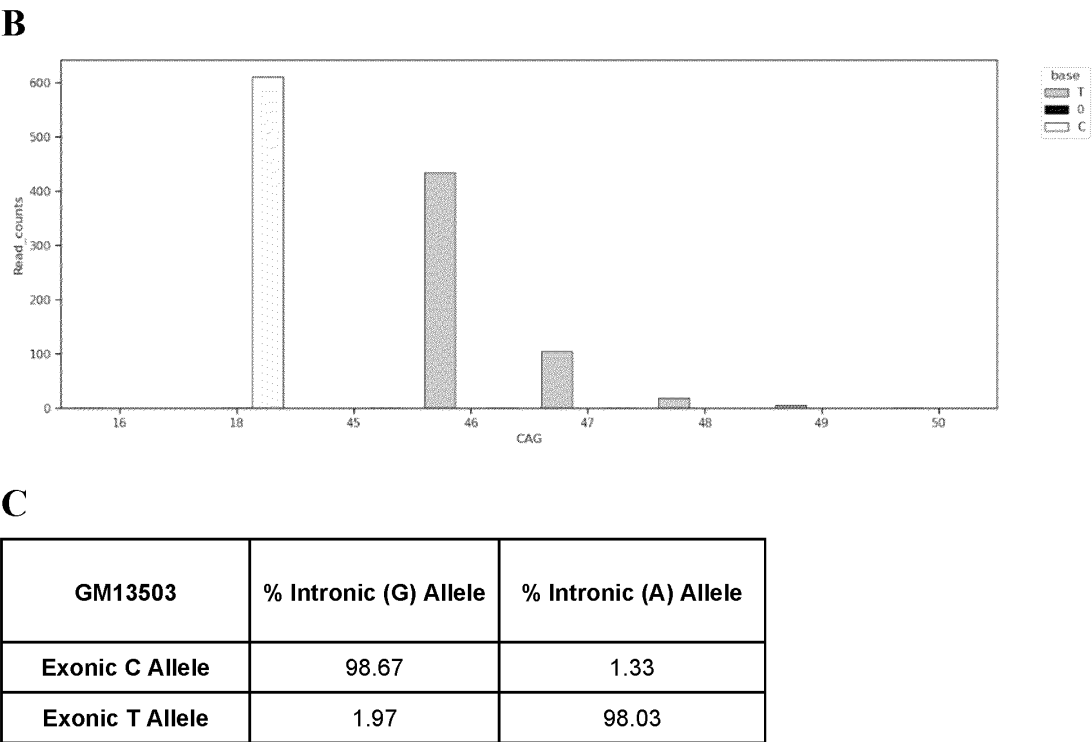
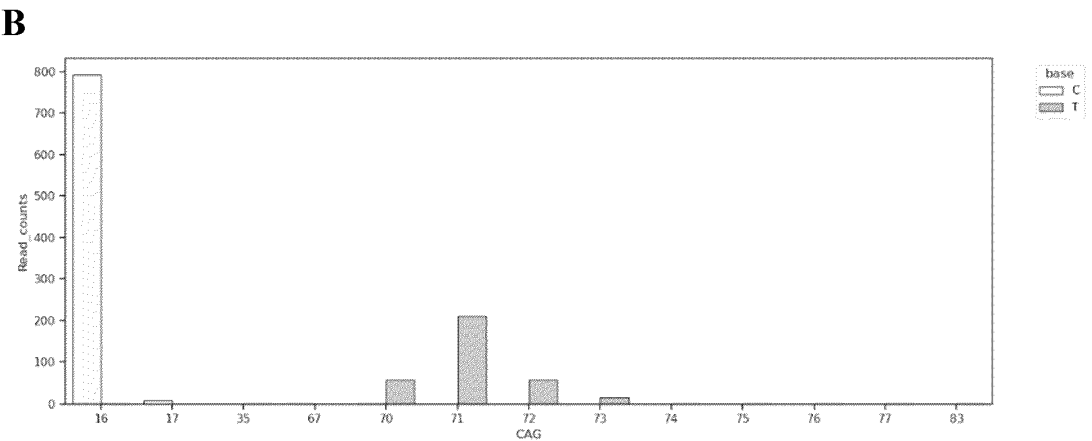


FIG.5

A

GM04724	Mean CAG repeat	Standard deviation	% reads	% Variant Exonic (C) SNP Allele	% Reference Exonic (T) SNP Allele	% Undefined
High CAG	71.21	1.18	30.67	0.55	96.68	2.77
Low CAG	16.03	0.67	69.33	98.16	0.25	1.59



C

GM04724	% Intronic (G) Allele	% Intronic (A) Allele
Exonic C Allele	97.54	2.46
Exonic T Allele	1.2	98.8

FIG.6

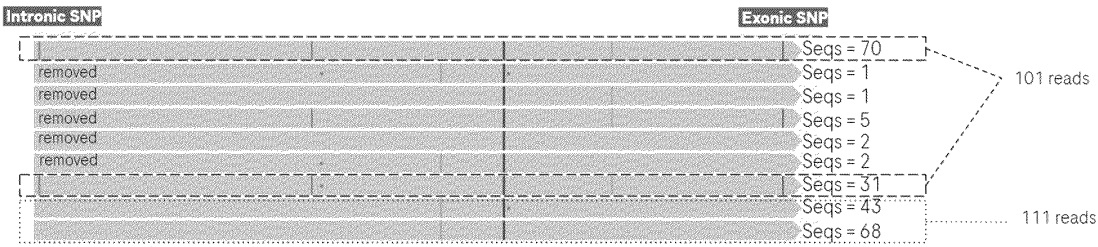


FIG.7

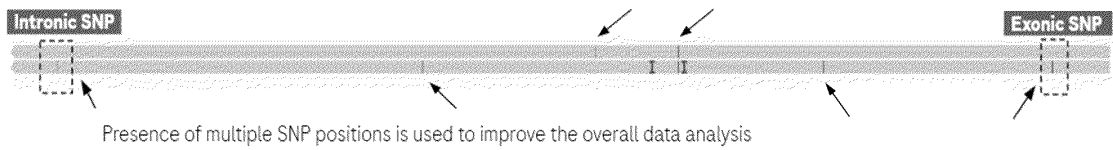


FIG. 8

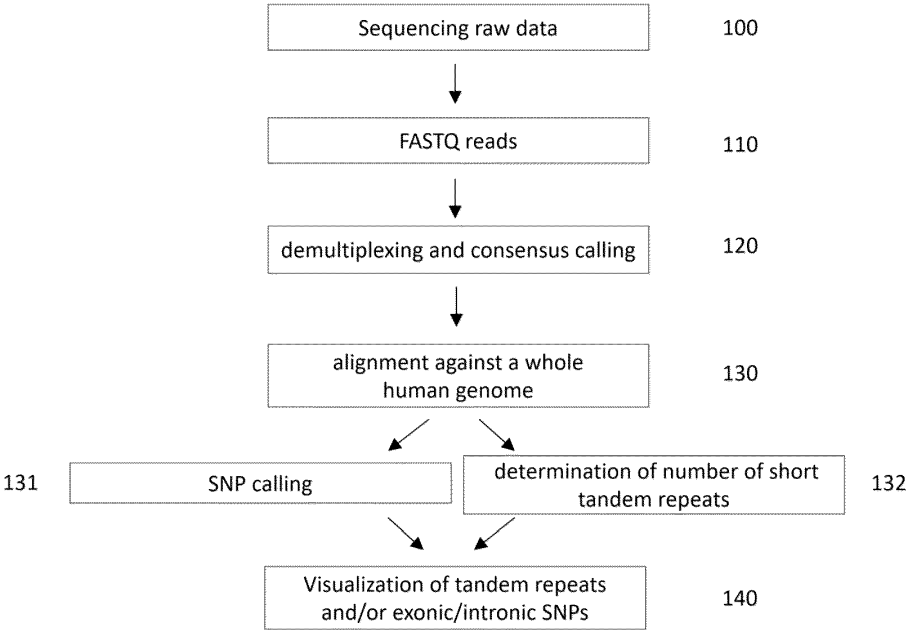


FIG. 9

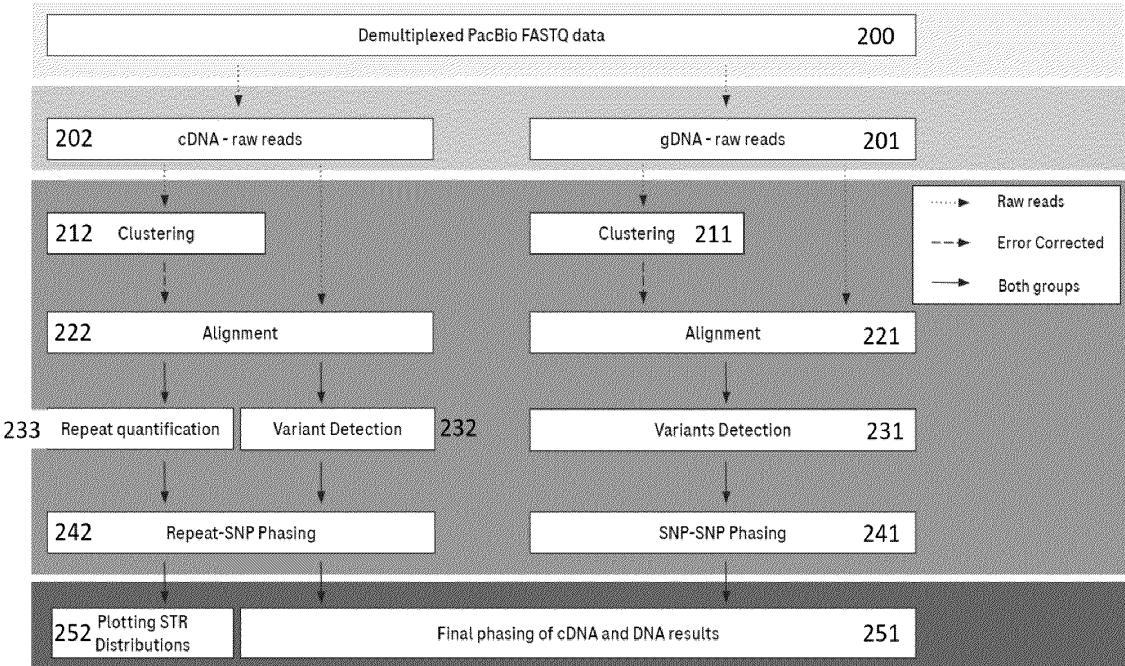


FIG. 10

