

eolf-seql  
SEQUENCE LISTING

<110> sphingotec GmbH

<120> A method for predicting the risk of incidence of chronic kidney disease

<130> S75039W0

<140>

<141>

<160> 12

<170> PatentIn version 3.5

<210> 1

<211> 243

<212> PRT

<213> Homo sapiens

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Glu Cys Ser Gln Asp Cys Ala Thr Cys Ser Tyr Arg Leu Val Arg Pro  
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Ala Asp Ile Asn Phe Leu Ala Cys Val Met Glu Cys Glu Gly Lys Leu  
20 25 30

Pro Ser Leu Lys Ile Trp Glu Thr Cys Lys Glu Leu Leu Gln Leu Ser  
35 40 45

Lys Pro Glu Leu Pro Gln Asp Gly Thr Ser Thr Leu Arg Glu Asn Ser  
50 55 60

Lys Pro Glu Glu Ser His Leu Leu Ala Lys Arg Tyr Gly Gly Phe Met  
65 70 75 80

Lys Arg Tyr Gly Gly Phe Met Lys Lys Met Asp Glu Leu Tyr Pro Met  
85 90 95

Glu Pro Glu Glu Glu Ala Asn Gly Ser Glu Ile Leu Ala Lys Arg Tyr

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100

105

110

Gly Gly Phe Met Lys Lys Asp Ala Glu Glu Asp Asp Ser Leu Ala Asn  
115 120 125

Ser Ser Asp Leu Leu Lys Glu Leu Leu Glu Thr Gly Asp Asn Arg Glu  
130 135 140

Arg Ser His His Gln Asp Gly Ser Asp Asn Glu Glu Glu Val Ser Lys  
145 150 155 160

Arg Tyr Gly Gly Phe Met Arg Gly Leu Lys Arg Ser Pro Gln Leu Glu  
165 170 175

Asp Glu Ala Lys Glu Leu Gln Lys Arg Tyr Gly Gly Phe Met Arg Arg  
180 185 190

Val Gly Arg Pro Glu Trp Trp Met Asp Tyr Gln Lys Arg Tyr Gly Gly  
195 200 205

Phe Leu Lys Arg Phe Ala Glu Ala Leu Pro Ser Asp Glu Glu Gly Glu  
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Ser Tyr Ser Lys Glu Val Pro Glu Met Glu Lys Arg Tyr Gly Gly Phe  
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Met Arg Phe

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Glu Cys Ser Gln Asp Cys Ala Thr Cys Ser Tyr Arg Leu Val Arg Pro  
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Ala Asp Ile Asn Phe Leu Ala Cys Val Met Glu Cys Glu Gly Lys Leu  
20 25 30

Pro Ser Leu Lys Ile Trp Glu Thr Cys Lys Glu Leu Leu Gln Leu Ser  
35 40 45

Lys Pro Glu Leu Pro Gln Asp Gly Thr Ser Thr Leu Arg Glu Asn Ser  
50 55 60

Lys Pro Glu Glu Ser His Leu Leu Ala  
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Tyr Gly Gly Phe Met  
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Tyr Gly Gly Phe Leu  
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Met Asp Glu Leu Tyr Pro Met Glu Pro Glu Glu Glu Ala Asn Gly Ser

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Glu Ile Leu Ala  
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Asp Ala Glu Glu Asp Asp Ser Leu Ala Asn Ser Ser Asp Leu Leu Lys  
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Glu Leu Leu Glu Thr Gly Asp Asn Arg Glu Arg Ser His His Gln Asp  
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Gly Ser Asp Asn Glu Glu Glu Val Ser  
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Tyr Gly Gly Phe Met Arg Gly Leu  
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Ser Pro Gln Leu Glu Asp Glu Ala Lys Glu Leu Gln  
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Val Gly Arg Pro Glu Trp Trp Met Asp Tyr Gln  
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Phe Ala Glu Ala Leu Pro Ser Asp Glu Glu Gly Glu Ser Tyr Ser Lys  
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Glu Val Pro Glu Met Glu  
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Phe Ala Glu Ala Leu Pro Ser Asp Glu Glu Gly Glu Ser Tyr Ser Lys  
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Glu Val Pro Glu Met Glu Lys Arg Tyr Gly Gly Phe  
 20 25

<210> 12  
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Tyr Gly Gly Phe Met Arg Phe  
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