

SEQUENCE LISTING

<110> Oy Jurilab Ltd

<120> The method for detecting a risk of cancer, coronary heart disease,
and stroke

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<170> PatentIn version 3.1

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 Glu Gln Arg Ala Ala Gln Lys Ala Asp Val Leu Thr Thr Gly Ala Gly
 20 25 30

aac cca gta gga gac aaa ctt aat gtt att aca gta ggg ccc cgt ggg 144
 Asn Pro Val Gly Asp Lys Leu Asn Val Ile Thr Val Gly Pro Arg Gly
 35 40 45

ccc ctt ctt gtt cag gat gtg gtt ttc act gat gaa atg gct cat ttt 192
 Pro Leu Leu Val Gln Asp Val Val Phe Thr Asp Glu Met Ala His Phe
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 Asp Arg Glu Arg Ile Pro Glu Arg Val Val His Ala Lys Gly Ala Gly

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gca aag gta ttt gag cat att gga aag aag act ccc atc gca gtt cgg Ala Lys Val Phe Glu His Ile Gly Lys Lys Thr Pro Ile Ala Val Arg 100 105 110				336
ttc tcc act gtt gct gga gaa tcg ggt tca gct gac aca gtt cgg gac Phe Ser Thr Val Ala Gly Glu Ser Gly Ser Ala Asp Thr Val Arg Asp 115 120 125				384
cct cgt ggg ttt gca gtg aaa ttt tac aca gaa gat ggt aac tgg gat Pro Arg Gly Phe Ala Val Lys Phe Tyr Thr Glu Asp Gly Asn Trp Asp 130 135 140				432
ctc gtt gga aat aac acc ccc att ttc ttc atc agg gat ccc ata ttg Leu Val Gly Asn Asn Thr Pro Ile Phe Phe Ile Arg Asp Pro Ile Leu 145 150 155 160				480
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325 330 335

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Ser Asn Met Pro Pro Gly Ile Glu Ala Ser Pro Asp Lys Met Leu Gln
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ggc cgc ctt ttt gcc tat cct gac act cac cgc cat cgc ctg gga ccc 1104
Gly Arg Leu Phe Ala Tyr Pro Asp Thr His Arg His Arg Leu Gly Pro
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Gly His Leu Lys Asp Ala Gln Ile Phe Ile Gln Lys Lys Ala Val Lys
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ttc tcc act gtt gct gga gaa tcg ggt tca gct gac aca gtt cgg gac 384
 Phe Ser Thr Val Ala Gly Glu Ser Gly Ser Ala Asp Thr Val Arg Asp
 115 120 125

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 Pro Arg Gly Phe Ala Val Lys Phe Tyr Thr Glu Asp Gly Asn Trp Asp
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ctc gtt gga aat aac acc ccc att ttc ttc atc agg gat ccc ata ttg 480
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 145 150 155 160

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ctg cat cag gtt tct ttc ttg ttc agt gat cgg ggg att cca gat gga							624
Leu His Gln Val Ser Phe Leu Phe Ser Asp Arg Gly Ile Pro Asp Gly	195		200		205		
cat cgc cac atg aat gga tat gga tca cat act ttc aag ctg gtt aat							672
His Arg His Met Asn Gly Tyr Gly Ser His Thr Phe Lys Leu Val Asn	210		215		220		
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Ala Asn Gly Glu Ala Val Tyr Cys Lys Phe His Tyr Lys Thr Asp Gln	225		230		235		240
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Gly Ile Lys Asn Leu Ser Val Glu Asp Ala Ala Arg Leu Ser Gln Glu	245		250		255		
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Asp Pro Asp Tyr Gly Ile Arg Asp Leu Phe Asn Ala Ile Ala Thr Gly	260		265		270		
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Lys Tyr Pro Ser Trp Thr Phe Tyr Ile Gln Val Met Thr Phe Asn Gln	275		280		285		
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Ala Glu Thr Phe Pro Phe Asn Pro Phe Asp Leu Thr Lys Val Trp Pro	290		295		300		
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Asn Pro Val Asn Tyr Phe Ala Glu Val Glu Gln Ile Ala Phe Asp Pro	325		330		335		
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Ser Asn Met Pro Pro Gly Ile Glu Ala Ser Pro Asp Lys Met Leu Gln	340		345		350		
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Asn Val Leu Asn Glu Glu Gln Arg Lys Arg Leu Cys Glu Asn Ile Ala
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Gly His Leu Lys Asp Ala Gln Ile Phe Ile Gln Lys Lys Ala Val Lys
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Asn Phe Thr Glu Val His Pro Asp Tyr Gly Ser His Ile Gln Ala Leu
 485 490 495

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