

--Liver Detox - Phase I --				
Gene	rsID	risk	your Allele	Results
BMP2 C282Y	rs235756	G	GG	+/+
CYP1A1*2C A4889G	rs1048943	C	CT	+/-
CYP1A2 C164A	rs762551	C	AC	+/-
CYP1B1 N453S	rs1800440	T	TT	+/+
CYP1B1 R48G	rs10012	C	CC	+/+
CYP2D6 S486T	rs1135840	C	CC	+/+
GSTM1 7730C>T	rs1056806	T	CT	+/-
GSTP1 A114V	rs1138272	T	CT	+/-
GSTP1 I105V	rs1695	G	AG	+/-
PON1 Q192R	rs662	C	CT	+/-
SOD3 489 C>T	rs2855262	T	CT	+/-
UGT1A1 C175181T	rs887829	T	CT	+/-
UGT1A1 C179920T	rs4148325	T	CT	+/-
UGT1A1 G179250T	rs6742078	T	GT	+/-
--Liver Detox - Phase II --				
Gene	rsID	risk	your Allele	Results
ABP1/DAO C47T	rs10156191	T	CT	+/-
ACAT1 G22670A	rs3741049	A	AG	+/-
ACE G2328A	rs4343	G	AG	+/-
ADA G22A	rs73598374	T	CT	+/-
AGT M235T/C4072T	rs699	G	GG	+/+
AMT T5998G	rs4855873	C	CC	+/+
APOC3 G34G	rs4520	T	CT	+/-
BHMT-02 C13813T	rs567754	T	CT	+/-
BHMT-08 C6457T	rs651852	T	TT	+/+
CAT A12175G	rs480575	G	AG	+/-
CAT T5070C	rs1049982	C	CT	+/-
COMT/TXNRD2 A4251G	rs737866	C	CT	+/-
COMT/TXNRD2 T4239C	rs737865	G	AG	+/-
DAO A24464G	rs3918347	G	GG	+/+
DHFR A16352G	rs1643649	C	CT	+/-
DHFR A20965G	rs1643659	C	CT	+/-
DHFR C19483A	rs1677693	T	GT	+/-
DHFR/MSH T-473A	rs1650697	A	AG	+/-
DISC1 C14853T	rs823162	C	CT	+/-
--Liver Detox - Phase II --				
Gene	rsID	risk	your Allele	Results
DISC1 R264Q	rs3738401	A	AG	+/-
FOLR2 G-1316A	rs651933	A	AG	+/-
FUT2 C12376T	rs281377	T	CT	+/-

GAD1 C34281T	rs701492	T	TT	+/+
GAD1 G25509C	rs769407	C	CC	+/+
GAD1 G3992T	rs3791878	T	TT	+/+
GAD1 T30473C	rs3791851	C	CC	+/+
GART A9979G	rs4817579	T	TT	+/+
GGH C6699T	rs3780126	A	AG	+/-
GGT1 T17549C	rs5751901	T	TT	+/+
GSR A43851G	rs2551715	C	CT	+/-
--Liver Detox - Phase II --				
Gene	rsID	risk	your Allele	Results
GSTP1 A114V	rs1138272	T	CT	+/-
GSTP1 I105V	rs1695	G	AG	+/-
HNMT A47507G	rs1378321	G	AG	+/-
HNMT T939C	rs1050891	G	AG	+/-
HRH4 T3537649C	rs16940765	C	CC	+/+
HTR2A T64185C	rs7997012	G	GG	+/+
MAT1A C1131T	rs2993763	A	AG	+/-
MAT1A C15656T	rs4934028	A	AG	+/-
MAT1A T*1297C	rs1985908	G	AG	+/-
MAT2B A7755681G	rs4869089	G	GG	+/+
MMAB/MVK A-818G	rs12314392	A	AG	+/-
MTHFD1 G1958A	rs2236225	A	AA	+/+
MTHFD1L T31397C	rs17349743	C	CT	+/-
MTHFD2 C8503T	rs1667627	C	CT	+/-
MTHFS - ST20 MTHFS G396	rs6495446	C	CT	+/-
MTHFS - ST20 MTHFS G560	rs2733103	T	CT	+/-
NAT2 C481T	rs1799929	T	CT	+/-
NAT2 I114T	rs1801280	C	CT	+/-
NAT2 K268R	rs1208	G	AG	+/-
--Liver Detox - Phase II --				
Gene	rsID	risk	your Allele	Results
NQO1 C494+	rs10517	A	AG	+/-
NQO1 C609T	rs1800566	A	AG	+/-
NQO1 G13070C	rs689452	C	CG	+/-
NQO1 T6314C	rs2917669	A	AG	+/-
NQO1 T7706C	rs1437135	G	AG	+/-
NRG1 C3314T	rs6994992	T	CT	+/-
SLC6A2 A40223G	rs4564560	G	AG	+/-
SLC6A2 C41517T	rs11568324	T	CT	+/-
SLC6A2 C45295T	rs3785157	T	CT	+/-
SLC6A2 C51371T	rs2242447	C	CT	+/-
SLC6A3 C17719T	rs460000	T	TT	+/+

SLC6A3 G37899A	rs27048	C	CC	+/+
SLC6A3 G45996T	rs6869645	T	CT	+/-
SLC6A3 G48964A	rs11133767	T	CT	+/-
SLC6A3 G56022A	rs27072	C	CC	+/+
SLC6A3 T55729C	rs1042098	G	AG	+/-
SULT1C3 A108878711G	rs1470874	A	AG	+/-
SULT1C3 M194T	rs6722745	T	TT	+/+
SULT2A1 A20117G	rs296366	T	CT	+/-
--Liver Detox - Phase II --				
Gene	rsID	risk	your Allele	Results
SULT2A1 G9598T	rs2547231	C	AC	+/-
SULT2A1 G9696A	rs4149449	T	CT	+/-
--Yeast/Alcohol Metabolism --				
Gene	rsID	risk	your Allele	Results
ALDH2 T12488C	rs2238151	T	CT	+/-
--Methylation & Methionine/Homocysteine Pathways --				
Gene	rsID	risk	your Allele	Results
BHMT-02 C13813T	rs567754	T	CT	+/-
BHMT-08 C6457T	rs651852	T	TT	+/+
CBS A360A	rs1801181	A	AG	+/-
CBS C*351T	rs706209	A	AA	+/+
CBS C699T	rs234706	A	AG	+/-
CBS T*330C	rs706208	G	GG	+/+
CTH A11886G	rs1145920	A	AG	+/-
CTH T8763C	rs681475	T	TT	+/+
DHFR A16352G	rs1643649	C	CT	+/-
DHFR A20965G	rs1643659	C	CT	+/-
DHFR C19483A	rs1677693	T	GT	+/-
DHFR/MSH T-473A	rs1650697	A	AG	+/-
FOLR2 G-1316A	rs651933	A	AG	+/-
MAT1A C1131T	rs2993763	A	AG	+/-
MAT1A C15656T	rs4934028	A	AG	+/-
MAT1A T*1297C	rs1985908	G	AG	+/-
MAT2B A7755681G	rs4869089	G	GG	+/+
MTHFD1 G1958A	rs2236225	A	AA	+/+
--Methylation & Methionine/Homocysteine Pathways --				
Gene	rsID	risk	your Allele	Results
MTHFD1L T31397C	rs17349743	C	CT	+/-
MTHFD2 C8503T	rs1667627	C	CT	+/-
MTHFR 03 P39P	rs2066470	A	AG	+/-
MTHFR A*372C	rs4846049	T	GT	+/-
MTHFR A1298C	rs1801131	G	GT	+/-

MTHFR A1572G	rs17367504	G	AG	+/-
MTHFR C10318T	rs17037390	A	AG	+/-
MTHFR G1793A (R594Q)	rs2274976	T	CT	+/-
MTHFR G18861A	rs1476413	T	CT	+/-
MTHFS - ST20 MTHFS G396	rs6495446	C	CT	+/-
MTHFS - ST20 MTHFS G560	rs2733103	T	CT	+/-
MTR A50417C	rs2789352	A	AC	+/-
MTR C62048T	rs2275568	T	CT	+/-
MTR G35489C	rs12060570	G	GG	+/+
MTR G74984T	rs3768142	G	GT	+/-
MTR G94982A	rs2275566	G	AG	+/-
MTR G95096T	rs2275565	T	GT	+/-
MTR T9195C	rs10925235	T	CT	+/-
MTRR -11 A664A	rs1802059	A	AG	+/-
MTRR A66G	rs1801394	G	AG	+/-
MTRR C32295T	rs3776455	T	CT	+/-
MTRR C524T	rs1532268	T	CT	+/-
MTRR G*541A	rs9332	A	AG	+/-
MTRR G12099A	rs3776467	G	AG	+/-
MTRR K350A	rs162036	G	AG	+/-
MTRR T*662A	rs8659	A	AT	+/-
--Methylation & Methionine/Homocysteine Pathways --				
Gene	rsID	risk	your Allele	Results
MTRR T12072C	rs326121	C	CT	+/-
NOS1 T2202C	rs2293054	A	AG	+/-
NOS3 A6251T	rs1800783	A	AT	+/-
NOS3 C19635T	rs3918188	A	AC	+/-
NOS3 G6797A	rs1800779	G	AG	+/-
PEMT G634A	rs7946	T	TT	+/+
PEMT T17020543A	rs4646406	A	AA	+/+
SULT1C3 A108878711G	rs1470874	A	AG	+/-
SULT1C3 M194T	rs6722745	T	TT	+/+
SULT2A1 A20117G	rs296366	T	CT	+/-
SULT2A1 G9598T	rs2547231	C	AC	+/-
SULT2A1 G9696A	rs4149449	T	CT	+/-
--Trans-Sulfuration Pathway --				
Gene	rsID	risk	your Allele	Results
CBS A360A	rs1801181	A	AG	+/-
CBS C*351T	rs706209	A	AA	+/+
CBS C699T	rs234706	A	AG	+/-
CBS T*330C	rs706208	G	GG	+/+
CTH A11886G	rs1145920	A	AG	+/-

CTH T8763C	rs681475	T	TT	+/+
MTHFS - ST20 MTHFS G396	rs6495446	C	CT	+/-
MTHFS - ST20 MTHFS G560	rs2733103	T	CT	+/-
SULT1C3 A108878711G	rs1470874	A	AG	+/-
SULT1C3 M194T	rs6722745	T	TT	+/+
SULT2A1 A20117G	rs296366	T	CT	+/-
SULT2A1 G9598T	rs2547231	C	AC	+/-
SULT2A1 G9696A	rs4149449	T	CT	+/-

--Neurotransmitter Pathway: Serotonin & Dopamine --

Gene	rsID	risk	your Allele	Results
AANAT C10236T	rs11077820	T	TT	+/+
ALDH2 T12488C	rs2238151	T	CT	+/-
ANK3 G658454A	rs9804190	T	CT	+/-
ANKK1 E713K	rs1800497	A	AG	+/-
ANKK1 G318R	rs11604671	A	AG	+/-
COMT A 26166G	rs2239393	G	AG	+/-
COMT A-1324G	rs6269	G	AG	+/-
COMT A309G	rs174675	T	CT	+/-
COMT C28914T	rs174696	C	CT	+/-
COMT C31430T	rs9332377	T	CT	+/-
COMT G*522A	rs165599	G	AG	+/-
COMT T13376G	rs5993883	G	GT	+/-
COMT T24075C	rs4646312	C	CT	+/-
COMT T26501G	rs740601	T	GT	+/-
DBH A1410G	rs77905	A	AA	+/+
DBH A486G	rs1108580	A	AG	+/-
DBH C12599T	rs1611123	T	CT	+/-
DBH C12828T	rs1611125	C	CT	+/-

--Neurotransmitter Pathway: Serotonin & Dopamine --

Gene	rsID	risk	your Allele	Results
DBH C18813T	rs2283123	C	CC	+/+
DBH G15032A	rs1541332	G	AG	+/-
DBH T13150C	rs2519152	C	CT	+/-
DBH T15791C	rs2519154	C	CT	+/-
DBH T16031C	rs2797853	T	CT	+/-
DBH T7335C	rs2007153	C	CC	+/+
DBH T8114C	rs2519155	C	CT	+/-
DBH T9160C	rs2873804	C	CT	+/-
DDC A14870G	rs921451	T	CT	+/-
DDC A19551G	rs10499695	C	CT	+/-
DDC A85104G	rs1451371	T	TT	+/+
DDC C166017G	rs1470750	C	CC	+/+

DDC G1385A	rs11575542	T	CT	+/-
DDC G15443T	rs1451375	A	AC	+/-
DHFR A16352G	rs1643649	C	CT	+/-
DHFR A20965G	rs1643659	C	CT	+/-
DHFR C19483A	rs1677693	T	GT	+/-
DHFR/MSH T-473A	rs1650697	A	AG	+/-
DRD1 C7464T	rs686	A	AG	+/-
DRD1 G6014A	rs4532	T	CT	+/-
DRD1 T5262C	rs265981	G	AG	+/-
DRD2 A32594G	rs4936270	C	CC	+/+
DRD2 A33029G	rs4245146	T	CT	+/-
DRD2 A53817G	rs1125394	C	CT	+/-
DRD2 A61820G	rs1079727	T	CT	+/-
DRD2 C31550T	rs4274224	G	AG	+/-
DRD2 C33935T	rs17529477	A	AG	+/-
DRD2 C36639T	rs4648319	A	AG	+/-
--Neurotransmitter Pathway: Serotonin & Dopamine --				
Gene	rsID	risk	your Allele	Results
DRD2 C957T	rs6277	G	AG	+/-
DRD2 G52663A	rs2471857	T	CT	+/-
DRD2 G54383A	rs1079596	T	CT	+/-
DRD2 G54716A	rs1079597	T	CT	+/-
DRD2 G65466T	rs2283265	C	AC	+/-
DRD2 G67314T	rs1076560	A	AC	+/-
DRD2 T21228G	rs7131056	A	AC	+/-
DRD2 T55093G	rs1076563	C	AC	+/-
DRD2 T64501C	rs2734838	G	AG	+/-
DRD3 C26625T	rs167771	G	AG	+/-
DRD3 G17248T	rs10934256	A	AC	+/-
DRD3 T14368C	rs1486009	G	AG	+/-
DRD3 T21277C	rs324029	A	AA	+/+
DRD4 C8887A	rs11246226	A	AC	+/-
GCH1 A14340G	rs3783642	C	CC	+/+
GCH1 A14404T	rs3783641	A	AA	+/+
GCH1 A30528G	rs12147422	C	CT	+/-
GCH1 A53980G	rs4411417	C	CT	+/-
GCH1 C12707T	rs8017210	A	AA	+/+
GCH1 C15878T	rs7147286	A	AA	+/+
GCH1 C36378991T	rs8007267	T	TT	+/+
GCH1 C37668A	rs7492600	T	GT	+/-
GCH1 G26425A	rs3783637	T	CT	+/-
GCH1 G55306457C	rs10483639	C	CG	+/-

GCH1 G62974A	rs752688	T	CT	+/-
--Neurotransmitter Pathway: Serotonin & Dopamine --				
Gene	rsID	risk	your Allele	Results
MAOA A16535C	rs5906883	A	A	+/+
MAOA A85020G	rs2235186	A	A	+/+
MAOA C42794T	rs909525	C	C	+/+
MAOA G3638A	rs5953210	G	G	+/+
MAOB A118723G	rs1799836	C	C	+/+
NOS1 T2202C	rs2293054	A	AG	+/-
NOS3 A6251T	rs1800783	A	AT	+/-
NOS3 C19635T	rs3918188	A	AC	+/-
NOS3 G6797A	rs1800779	G	AG	+/-
PAH C35625G	rs1522305	G	GG	+/+
PAH C45188T	rs1718301	A	AA	+/+
PAH G735A	rs1042503	T	TT	+/+
PEMT G634A	rs7946	T	TT	+/+
PEMT T17020543A	rs4646406	A	AA	+/+
PNMT G-184A	rs876493	A	AA	+/+
SLC6A2 A40223G	rs4564560	G	AG	+/-
SLC6A2 C41517T	rs11568324	T	CT	+/-
SLC6A2 C45295T	rs3785157	T	CT	+/-
SLC6A2 C51371T	rs2242447	C	CT	+/-
--Neurotransmitter Pathway: Serotonin & Dopamine --				
Gene	rsID	risk	your Allele	Results
SLC6A3 C17719T	rs460000	T	TT	+/+
SLC6A3 G37899A	rs27048	C	CC	+/+
SLC6A3 G45996T	rs6869645	T	CT	+/-
SLC6A3 G48964A	rs11133767	T	CT	+/-
SLC6A3 G56022A	rs27072	C	CC	+/+
SLC6A3 T55729C	rs1042098	G	AG	+/-
TH V112M	rs6356	T	CT	+/-
--Neurotransmitter Pathway: Glutamate & GABA --				
Gene	rsID	risk	your Allele	Results
GAD1 C34281T	rs701492	T	TT	+/+
GAD1 G25509C	rs769407	C	CC	+/+
GAD1 G3992T	rs3791878	T	TT	+/+
GAD1 T30473C	rs3791851	C	CC	+/+
--COMT Activity --				
Gene	rsID	risk	your Allele	Results
ALDH2 T12488C	rs2238151	T	CT	+/-
ANK3 G658454A	rs9804190	T	CT	+/-
ANKK1 E713K	rs1800497	A	AG	+/-

ANKK1 G318R	rs11604671	A	AG	+/-
COMT A 26166G	rs2239393	G	AG	+/-
COMT A-1324G	rs6269	G	AG	+/-
COMT A309G	rs174675	T	CT	+/-
COMT C28914T	rs174696	C	CT	+/-
COMT C31430T	rs9332377	T	CT	+/-
COMT G*522A	rs165599	G	AG	+/-
COMT T13376G	rs5993883	G	GT	+/-
COMT T24075C	rs4646312	C	CT	+/-
COMT T26501G	rs740601	T	GT	+/-
CYP1A1*2C A4889G	rs1048943	C	CT	+/-
CYP1B1 N453S	rs1800440	T	TT	+/+
CYP1B1 R48G	rs10012	C	CC	+/+
DBH A1410G	rs77905	A	AA	+/+
DBH A486G	rs1108580	A	AG	+/-
DBH C12599T	rs1611123	T	CT	+/-
DBH C12828T	rs1611125	C	CT	+/-
DBH C18813T	rs2283123	C	CC	+/+
DBH G15032A	rs1541332	G	AG	+/-
DBH T13150C	rs2519152	C	CT	+/-
DBH T15791C	rs2519154	C	CT	+/-
DBH T16031C	rs2797853	T	CT	+/-
DBH T7335C	rs2007153	C	CC	+/+
DBH T8114C	rs2519155	C	CT	+/-
DBH T9160C	rs2873804	C	CT	+/-
DRD1 C7464T	rs686	A	AG	+/-
DRD1 G6014A	rs4532	T	CT	+/-
DRD1 T5262C	rs265981	G	AG	+/-
DRD2 A32594G	rs4936270	C	CC	+/+
DRD2 A33029G	rs4245146	T	CT	+/-
DRD2 A53817G	rs1125394	C	CT	+/-
DRD2 A61820G	rs1079727	T	CT	+/-
DRD2 C31550T	rs4274224	G	AG	+/-
DRD2 C33935T	rs17529477	A	AG	+/-
DRD2 C36639T	rs4648319	A	AG	+/-
DRD2 C957T	rs6277	G	AG	+/-
DRD2 G52663A	rs2471857	T	CT	+/-
DRD2 G54383A	rs1079596	T	CT	+/-
DRD2 G54716A	rs1079597	T	CT	+/-
DRD2 G65466T	rs2283265	C	AC	+/-
DRD2 G67314T	rs1076560	A	AC	+/-
DRD2 T21228G	rs7131056	A	AC	+/-

DRD2 T55093G	rs1076563	C	AC	+/-
DRD2 T64501C	rs2734838	G	AG	+/-
DRD3 C26625T	rs167771	G	AG	+/-
DRD3 G17248T	rs10934256	A	AC	+/-
DRD3 T14368C	rs1486009	G	AG	+/-
DRD3 T21277C	rs324029	A	AA	+/+
DRD4 C8887A	rs11246226	A	AC	+/-
MAOA A16535C	rs5906883	A	A	+/+
MAOA A85020G	rs2235186	A	A	+/+
MAOA C42794T	rs909525	C	C	+/+
MAOA G3638A	rs5953210	G	G	+/+
MAOB A118723G	rs1799836	C	C	+/+
PAH C35625G	rs1522305	G	GG	+/+
PAH C45188T	rs1718301	A	AA	+/+
PAH G735A	rs1042503	T	TT	+/+
PNMT G-184A	rs876493	A	AA	+/+
TH V112M	rs6356	T	CT	+/-
--Mitochondrial Function --				
Gene	rsID	risk	your Allele	Results
ALDH2 T12488C	rs2238151	T	CT	+/-
GCH1 A14340G	rs3783642	C	CC	+/+
GCH1 A14404T	rs3783641	A	AA	+/+
GCH1 A30528G	rs12147422	C	CT	+/-
GCH1 A53980G	rs4411417	C	CT	+/-
GCH1 C12707T	rs8017210	A	AA	+/+
GCH1 C15878T	rs7147286	A	AA	+/+
GCH1 C36378991T	rs8007267	T	TT	+/+
GCH1 C37668A	rs7492600	T	GT	+/-
GCH1 G26425A	rs3783637	T	CT	+/-
GCH1 G55306457C	rs10483639	C	CG	+/-
GCH1 G62974A	rs752688	T	CT	+/-
MAT1A C1131T	rs2993763	A	AG	+/-
MAT1A C15656T	rs4934028	A	AG	+/-
MAT1A T*1297C	rs1985908	G	AG	+/-
MAT2B A7755681G	rs4869089	G	GG	+/+
MTHFD1 G1958A	rs2236225	A	AA	+/+
MTHFD1L T31397C	rs17349743	C	CT	+/-
MTHFD2 C8503T	rs1667627	C	CT	+/-
MTRR -11 A664A	rs1802059	A	AG	+/-
MTRR A66G	rs1801394	G	AG	+/-
MTRR C32295T	rs3776455	T	CT	+/-
MTRR C524T	rs1532268	T	CT	+/-

MTRR G*541A	rs9332	A	AG	+/-
MTRR G12099A	rs3776467	G	AG	+/-
MTRR K350A	rs162036	G	AG	+/-
MTRR T*662A	rs8659	A	AT	+/-
MTRR T12072C	rs326121	C	CT	+/-
NDUFS7	rs2332496	A	AG	+/-
NDUFS7	rs1142530	T	CT	+/-
NDUFS7	rs7258846	T	GT	+/-
NDUFS8	rs2075626	C	CT	+/-
NOS1 T2202C	rs2293054	A	AG	+/-
NOS3 A6251T	rs1800783	A	AT	+/-
NOS3 C19635T	rs3918188	A	AC	+/-
NOS3 G6797A	rs1800779	G	AG	+/-
PEMT G634A	rs7946	T	TT	+/+
PEMT T17020543A	rs4646406	A	AA	+/+
SOD2 406+816G>T	rs2758331	A	AC	+/-
SOD2 A16V	rs4880	G	AG	+/-
--Alzheimers/Cardio/Lipid --				
Gene	rsID	risk	your Allele	Results
ABCA7 T1046520G	rs3764650	G	GT	+/-
ADAM12 G63103C	rs3740199	C	CG	+/-
APBB2 G41002946C	rs13133980	C	CG	+/-
APOE IVS1+69	rs440446	G	CG	+/-
APP Glu647Asp	rs63750363	C	CC	+/+
APP Ala674Gly	rs63750671	G	GG	+/+
BDNF T64089C	rs11030104	A	AG	+/-
BDNF V81M	rs6265	T	CT	+/-
CETP I405V	rs5882	A	AA	+/+
CETP C4402A	rs1800775	C	AC	+/-
CR1 A27577G	rs6656401	A	AG	+/-
F2 C494T	rs5896	T	CT	+/-
GAB2 G42719T	rs2373115	A	AC	+/-
GP6 Pro219Ser	rs1613662	G	GG	+/+
HRG Pro204Ser	rs9898	T	TT	+/+
IDE G91606A	rs6583817	T	CT	+/-
IL6 A6262G	rs2069837	G	AG	+/-
KLC1 C.*396G	rs8702	G	GG	+/+
LRP6 V1062I	rs2302685	T	TT	+/+
NR1I2 C6709A	rs1523127	C	CC	+/+
SERPINC1 G5301A	rs2227589	T	CT	+/-
SORL1 C163447T	rs726601	T	CT	+/-
SORL1 G146128A	rs10892759	A	AA	+/+

SORL1 C164978A	rs1784931	C	CC	+/+
SORL1 T121434428C	rs11218342	C	CT	+/-
SORL1 G130294A	rs3781835	A	AG	+/-
SORL1 A123560G	rs1792124	A	AG	+/-
SORL1 G106922A	rs2298814	A	AG	+/-
SORL1 G108082A	rs6589885	A	AG	+/-
SORL1 G105680A	rs7946599	A	AG	+/-
THBD A7681G	rs1042580	C	CC	+/+
TOMM40 G45395266A	rs157580	G	AG	+/-
--IgE --				
Gene	rsID	risk	your Allele	Results
CD14	rs2569191	C	CC	+/+
SOCS-1 -820G>T	rs33977706	A	AC	+/-
FCGR2A	rs1801274	A	AG	+/-
--IgA --				
Gene	rsID	risk	your Allele	Results
HLA-DPB2 / COL11A2P	rs1883414	A	AA	+/+
HLA-DQA2	rs9275224	A	AA	+/+
IRF5	rs4728142	A	AA	+/+
MTC03P1	rs9275596	C	CC	+/+
TRAF1	rs3761847	G	AG	+/-
--Clotting Factors --				
Gene	rsID	risk	your Allele	Results
CETP C4402A	rs1800775	C	AC	+/-
GP6 Pro219Ser	rs1613662	G	GG	+/+
HRG Pro204Ser	rs9898	T	TT	+/+
NR112 C6709A	rs1523127	C	CC	+/+
SERPINC1 G5301A	rs2227589	T	CT	+/-
--Celiac Disease/Gluten I --				
Gene	rsID	risk	your Allele	Results
CTLA4	rs231775	G	AG	+/-
HLA DQA1	rs2187668	T	TT	+/+
HLA-DQA2	rs9275224	A	AA	+/+
--Thyroid --				
Gene	rsID	risk	your Allele	Results
CTLA4	rs231775	G	AG	+/-
--Eye Health --				
Gene	rsID	risk	your Allele	Results
BCMO1	rs4889294	C	CT	+/-
BCMO1 R267S	rs12934922	T	AT	+/-
--Other Immune Factors --				
Gene	rsID	risk	your Allele	Results

ATG16L1	rs10210302	C	CT	+/-
GSDMB	rs7216389	T	TT	+/+
IL5	rs2069812	A	AA	+/+
MeFV E148Q	rs3743930	G	CG	+/-
STAT4	rs10181656	G	CG	+/-