MTHFRSupport Covid Report

genome_Jamie_Hope_Full_20131107122754.zip

Report Summary

58
Total Variants

1
Categories

0
Pathway Figures

20
High Risk (+/+)

16
Moderate Risk (+/-)

22
Low Risk (-/-)

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Covid

58 variants found 20 16 22

Risk Summary: 20 high risk variants, 16 moderate risk variants - Consider discussing with healthcare provider

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SNP ID	SNP Name	Risk Allele	Your Allele	Result
rs689470	PTGS2/COX2 variant	A	GG	-/-
rs13306035	PTGS2/COX2 variant	G	AA	-/-
rs5276	PTGS2/COX2 variant	T	CC	-/-
rs2066826	PTGS2/COX2 variant	T	CC	-/-
rs5277	PTGS2/COX2 variant	G	CC	-/-
rs20417	PTGS2/COX2 variant	C	CC	+/+
rs20415	PTGS2/COX2 variant	T	CC	-/-
rs2027432	NLRP3 variant	G	GG	+/+
rs12048215	NLRP3 variant	A	AG	+/-
rs3806265	NLRP3 variant	T	CC	-/-
rs28937896	NLRP3 variant	T	TT	+/+
rs4612666	NLRP3 variant	C	TT	-/-
rs10754557	NLRP3 variant	G	AG	+/-
rs1539019	NLRP3 variant	C	AC	+/-
rs10157379	NLRP3 variant	T	CT	+/-
rs10754558	NLRP3 variant	C	GG	-/-
rs1143643	IL1B variant	T	TT	+/+
rs1143642	IL1B variant	G	GG	+/+
rs1143634	IL1B variant	G	GG	+/+
rs3136558	IL1B variant	A	AG	+/-
rs3917356	IL1B variant	T	TT	+/+
rs1143629	IL1B variant	G	AA	-/-
rs1143627	IL1B variant	G	AA	-/-
rs3087258	IL1B variant	G	GG	+/+
rs2536512	SOD3 A	G	AA	-/-
rs1799895	SOD3 R231G	G	CC	-/-
rs2069827	IL6 variant	T	GT	+/-
rs1800797	IL6 variant	G	AG	+/-
rs1800796	IL6 variant	G	GG	+/+
rs7802307	IL6 variant	T	AT	+/-
rs13447446	IL6 variant	G	GG	+/+
rs2069830	IL6 variant	C	CC	+/+
rs1524107	IL6 variant	T	CC	-/-
rs2066992	IL6 variant	T	GG	-/-
rs2069840	IL6 variant	C	CG	+/-
rs2069849	IL6 variant	C	CC	+/+
rs2069861	IL6 variant	C	CC	+/+
rs4496877	NOS3 variant	T	GT	+/-
rs3918226	NOS3 variant	T	CC	-/-
rs2853792	NOS3 variant	G	AG	+/-
rs3918227	NOS3 variant	C	CC	+/+

SNP ID	SNP Name	Risk Allele	Your Allele	Result
rs2853796	NOS3 variant	G	GT	+/-
rs743507	NOS3 variant	C	TT	-/-
rs580253	CASP1 variant	G	AG	+/-
rs1360485	HMGB1 variant	T	TT	+/+
rs1412125	HMGB1 variant	T	CT	+/-
rs4145277	HMGB1 variant	T	TT	+/+
rs4932178	FURIN variant	C	CC	+/+
rs17514846	FURIN variant	C	CC	+/+
rs4702	FURIN variant	G	GG	+/+
rs11538758	PRNP V210I	A	CC	-/-
rs1799990	PRNP M129V	G	GG	+/+
rs16990018	PRNP N171S	G	AA	-/-
rs28933385	PRNP E200K	A	GG	-/-
rs12626750	IFNAR1 variant	G	CC	-/-
rs2070788	TMPRSS2 variant	A	AG	+/-
rs12329760	TMPRSS2 variant	T	CC	-/-
rs2048683	ACE2 variant	G	GT	+/-

⚠ Important Disclaimer

Medical Disclaimer: This report is intended to translate your results into an easier to understand form. It is **not intended to diagnose or treat** any medical condition.

Professional Consultation: For diagnosis or treatment, please present this report to your qualified healthcare provider or find a practitioner on the MTHFRSupport™ website under "Find a Practitioner".

Genetic Interpretation: Genetic mutations are indicators that something *could* be affected, but they are **not a guarantee** that you are experiencing all or any of the associated issues.

Contributing Factors: Many factors influence whether genetic variants manifest as health issues, including:

- Environmental factors
- Ethnic background
- Diet and lifestyle
- Age and personal history
- Other genetic interactions