



Welcome to the future of health and human potential

NAME: TEST ACCOUNT

DATE: SEP 26, 2022



myDNA is an accredited laboratory with NATA approved Lab services based in Melbourne Australia and a CLIA certified and CAP accredited Lab in Houston Texas.

Each sample is run on parallel arrays that have been analytically validated.

This analysis does not provide information on genetic carrier risk profiles and is not intended to diagnose a disease.

For further information please contact Lab Services partner NutriPATH on 1300 688 522

The following table lists the genes covered in the myDNA Comprehensive Health Report.

| Gene | Wild Type | Heterozygous | Homozygous |
|---|-----------|--------------|------------|
| 5-HT2A | GG | | |
| | CC | | |
| | GG | | |
| 9p21 | | AG | |
| ABCG2 (Q141K) | GG | | |
| ACAT1-02 | GG | | |
| ACE1 G2350A | | AG | |
| ACE2 A8790G | AA | | |
| ACSL1 | | | AA |
| ACTN3 | | | TT |
| ADD1 | | GT | |
| ADIPOQ | TT | | |
| ADRB2 | GG | | |
| | | CG | |
| AGTR1 | AA | | |
| ALDH2 | GG | | |
| ANKK1 | | AG | |
| APB1 | CC | | |
| APOA2 | AA | | |
| APOE You have the ApoE e3/e3 genotype, improving cholesterol transport and the maintenance of brain neurons. The ApoE e3 allele improves cognitive fitness, HDL and LDL profiles, viral protection, and the response to plant bioactive compounds. | TT | | |
| | | | CC |
| ARMS2 | GG | | |
| ATM D1853N | | | AA |
| BCMO1 A379V | CC | | |
| BCMO1 R267S | | | TT |
| BDNF | CC | | |

| Gene | Wild Type | Heterozygous | Homozygous |
|-----------------|-----------|--------------|------------|
| CAT C-262T | | CT | |
| CBS | | CT | |
| CBS 191150T | GG | | |
| CBS A13637G | CC | | |
| CFH | | | CC |
| COL1A1 | CC | | |
| COMT | | AG | |
| | | CT | |
| COQ2 | | CT | |
| CTH | GG | | |
| CYP1A1 | TT | | |
| CYP1A2 | | | AA |
| CYP1B1*6 L432V | GG | | |
| CYP2C19*17 | | | TT |
| CYP2C9*3 A1075C | AA | | |
| CYP2D6 T100C | GG | | |
| CYP2E1 | CC | | |
| CYP2R1 | | | GG |
| CYP3A4*1B | TT | | |
| DAO C2029G | | CG | |
| DHFR A20965G | TT | | |
| DHFR C19483A | GG | | |
| DI01 | | AC | |
| DI02 | TT | | |
| ESR2 | AA | | |
| F5 | CC | | |
| FAAH | CC | | |
| FADS1 | CC | | |

| Gene | Wild Type | Heterozygous | Homozygous |
|---------------|-----------|--------------|------------|
| FADS2 | AA | | |
| | CC | | |
| FTO | | | AA |
| | | | GG |
| FUT2 | | AG | |
| GAD1 | | AG | |
| | TT | | |
| | | CT | |
| | | AG | |
| | GG | | |
| GATA3 | | GT | |
| GPX1 | | | AA |
| GSTM1 | AA | | |
| GSTP1 C341T | CC | | |
| GSTP1 I105V | AA | | |
| HFE-C282Y | GG | | |
| HLA DQ2.5 | | CT | |
| HLA-DQ8 | TT | | |
| HNMT | | AG | |
| HNMT C314T | | CT | |
| IL6 | | | GG |
| LCT | | AG | |
| LPA | TT | | |
| LZTFL1 | | | AA |
| MAO-A | | | GG |
| MDM2 | TT | | |
| MLH1 | GG | | |
| MTHFD1 G1958A | | AG | |
| MTHFR 1298 | TT | | |
| MTHFR 677 | | AG | |
| MTNR1B | | CG | |

| Gene | Wild Type | Heterozygous | Homozygous |
|------------|-----------|--------------|------------|
| MTR A2756G | AA | | |
| MTRR A66G | | | GG |
| NAT2 | | AG | |
| NBPF3 | | | CC |
| NOS1 | CC | | |
| NOS2 | | AG | |
| PEMT | | CT | |
| PON1 | | CT | |
| PPAR-alpha | CC | | |
| PPARGC1A | | CT | |
| PPCDC | | CT | |
| SELENBP1 | TT | | |
| SHBG | GG | | |
| | GG | | |
| | CC | | |
| SIRT1 | AA | | |
| SLC17A7 | | | GG |
| SLC23A1 | CC | | |
| SOD2 | | AG | |
| SOD3 | CC | | |
| TCF7L2 | | CT | |
| TCN2 C766G | | | GG |
| TFR2 | | | CC |
| TMPRSS2 | | | AA |
| TP53 | | CG | |
| VKORC1*2 | CC | | |
| XRCC3 | | AG | |