

## Targeted Genes and Methodology Details for Combined Humoral and Cell-Mediated Immunodeficiency Gene Panel

The following applies to COMID / Combined Humoral and Cell-Mediated Immunodeficiency Gene Panel, Varies. Testing is performed to evaluate for the presence of variants in coding regions and extending to +/- 10 base pairs of adjacent intronic sequence on either side of the coding exons of the genes analyzed. In addition, the analysis will cover select noncoding variants. Next-generation sequencing and/or a polymerase chain reaction-based quantitative method is performed to test for the presence of copy number variants (CNV) in the genes analyzed. Confirmation of select reportable variants may be performed by alternate methodologies based on internal laboratory criteria.

This list is current from July 2025 to the present. This document is intended to highlight additional evaluations for variants of high clinical interest as well as technical limitations. However, this document does not comprehensively reflect all genomic regions covered by this test. For questions regarding transcripts, genes or regions covered, contact the laboratory at 800-533-1710.

Genomic Build: GRCh37 (hg19) unless otherwise specified

Gene	Reference Transcript	Additional Evaluations	Technical Limitations
ADA	NM_000022.4	chr20:g.43249076C>T (c.976-34G>A); chr20:g.43248503A>T (c.1079-15T>A)	-
AK2	NM_001625.4	-	-
ARPC1B	NM_005720.4	-	-
ATM	NM_000051.3	chr11:g.108138753A>G (c.2639-384A>G); chr11:g.108139118_108139130del (c.2639-19_2639-7del); chr11: g.108141212_108141215del (c.2839-579_2839-576del); chr11:g.108164028A>G (c.4612-12A>G); chr11:g.108179837A>G (c.5763-1050A>G); chr11:g.108217993_108218019del (c.8585-13_8598del)	-
BCL10	NM_003921.5	-	-
BCL11B	NM_138576.4	-	-
BLM	NM_000057.4	-	-
CARD11	NM_032415.6	-	-
CD247	NM_198053.2	-	-
CD28	NM_006139.4	-	-
CD3D	NM_000732.4	-	-
CD3E	NM_000733.4	-	CNV analysis in exon 4 is not performed
CD3G	NM_000073.2	-	-
CD40	NM_001250.6	-	-
CD40LG	NM_000074.3	-	-
CD70	NM_001252.5	-	-
CD8A	NM_001768.6	-	-
CDCA7	NM_031942.5	-	-
CHD7	NM_017780.4	chr8:g.61763035 to chr8:g.61763039 (c.5405-17 to c.5405-13)	-
CIITA	NM_000246.3	-	-
CORO1A	NM_007074.3	-	CNV analysis in exon 11 is not performed
CTLA4	NM_005214.5	-	-

## Targeted Genes and Methodology Details for Combined Humoral and Cell-Mediated Immunodeficiency Gene Panel (continued)

Gene	Reference Transcript	Additional Evaluations	<b>Technical Limitations</b>
DCLRE1C	NM_001033855.3	-	-
DNMT3B	NM_006892.4	chr20:g.31395557G>A (c.2421-11G>A)	-
DOCK2	NM_004946.3	-	-
DOCK8	NM_203447.3	chr9:g.317025 to chr9:g.317028 (c.742-18 to c.742-15); chr9:g.372260_372348del (c.2083_2109+62del)	-
EPG5	NM_020964.3	-	-
ERCC6L2	NM_020207.5	-	-
EXTL3	NM_001440.4	-	-
FCHO1	NM_015122.3	-	-
FOXN1	NM_003593.2	-	-
GINS1	NM_021067.5	chr20:g.25388397A>G (c60A>G); chr20:g.25388409C>G (c48C>G);	-
HELLS	NM_018063.5	-	CNV analysis in exon 4 is not performed, sequence and CNV analysis in exon 7 is not performed
ICOS	NM_012092.4	-	-
ICOSLG	NM_015259.6	-	-
IKBKB	NM_001556.3	-	-
IKBKG	NM_001099857.3	chrX:g.153775960G>C (c16G>C); chrX:g.153788599A>T (c.519-23A>T); chrX:g.153787731C>T (c.518+866C>T); 11.7 kb deletion spanning <i>IKBKG</i> exons 4–10	CNV analysis in exons 3–10 is not performed with the exception of the common 11.7 kb deletion spanning exons 4–10
IKZF1	NM_001291845.2; NM_006060.6	-	CNV analysis in exons 4–8 is not performed
IL21	NM_021803.4	-	-
IL21R	NM_021798.4	-	-
IL2RA	NM_000417.3	-	-
IL2RB	NM_000878.5	-	-
IL2RG	NM_000206.2	chrX:g.70331494G>A (c105C>T); chrX:g.70330553T>C (c.270-15A>G); chrX:g.70329246_70329257del (c.595-15_595-4del)	-
IL7R	NM_002185.5	chr5:g.35867853G>A (c.379+288G>A)	-
ITK	NM_005546.3	-	-
ITPKB	NM_002221.3	-	-
JAK3	NM_000215.3	chr19:g.17946035C>T (c.1915-11G>A); chr19:g.17943239C>T (c.2680+89G>A)	-
KDM6A	NM_021140.3	-	-
KMT2A	NM_001197104.1	-	-
KMT2D	NM_003482.3	-	-
LAT	NM_001014987.2	-	-

Page 2 of 4 MC4091-449rev0625

## Targeted Genes and Methodology Details for Combined Humoral and Cell-Mediated Immunodeficiency Gene Panel (continued)

Gene	Reference Transcript	Additional Evaluations	<b>Technical Limitations</b>
LCK	NM_001042771.2	-	-
LCP2	NM_005565.5	-	CNV analysis in exon 8 is not performed
LIG1	NM_000234.3	-	-
LIG4	NM_002312.3	-	-
LRBA	NM_006726.4	-	CNV analysis in exons 20 and 39 is not performed
MAGT1	NM_032121.5	-	-
MALT1	NM_006785.4	-	CNV analysis in exon 8 is not performed
MAP3K14	NM_003954.5	-	-
МСМ10	NM_182751.3	-	-
MSN	NM_002444.3	-	-
MTHFD1	NM_005956.4	-	-
MYSM1	NM_001085487.3	-	-
NBN	NM_002485.4	-	CNV analysis in exon 16 is not performed
NFKBIA	NM_020529.2	-	-
NHEJ1	NM_024782.2	-	-
NSMCE3	NM_138704.4	-	-
ORAI1	NM_032790.3	-	-
PAX1	NM_006192.5	-	-
PGM3	NM_001199917.2	-	-
PNP	NM_000270.3	chr14:g.20942914G>A (c.286-18G>A)	-
POLD1	NM_002691.4	-	-
POLE	NM_006231.4	-	-
POLE2	NM_002692.4	-	CNV analysis in exons 8 and 19 is not performed
PRKDC	NM_006904.7	-	-
PSMB9	NM_002800.5	-	-
PTPRC	NM_002838.5	-	-
RAC2	NM_002872.5	-	-
RAG1	NM_000448.2	-	-
RAG2	NM_000536.4	-	-
RASGRP1	NM_005739.4	-	-
RBCK1	NM_031229.4	-	-
REL	NM_002908.4	-	CNV analysis in exon 9 is not performed
RELA	NM_021975.4	-	-
RELB	NM_006509.4	-	-

Page 3 of 4 MC4091-449rev0625

## Targeted Genes and Methodology Details for Combined Humoral and Cell-Mediated Immunodeficiency Gene Panel (continued)

Gene	Reference Transcript	Additional Evaluations	<b>Technical Limitations</b>
RFX5	NM_000449.3	-	-
RFXANK	NM_003721.4	chr19:g.19308305_19308330del (c.338-25_338del); chr19:g.19308411_19308468del (c.419_438+38del)	-
RFXAP	NM_000538.3	-	-
RHOH	NM_004310.5	-	-
RMRP (NME1)	NR_003051.3	5' UTR variants from n25 through the transcription initiation site	CNV analysis is not performed
RNF168	NM_152617.4	-	-
RNF31	NM_017999.5	-	-
RNU4ATAC	NR_023343.1	-	CNV analysis is not performed
SEC61A1	NM_013336.4	-	-
SEMA3E	NM_012431.3	-	-
SH2D1A	NM_002351.4	-	-
SKIV2L (SKIC2)	NM_006929.5	-	-
SMARCAL1	NM_014140.4	-	-
SP110	NM_004509.4	-	-
STAT3	NM_139276.2	chr17:g.40478306G>A (c.1282-89C>T)	-
STAT5B	NM_012448.4	-	CNV analysis in exons 6-8 is not performed
STIM1	NM_003156.3	-	-
STK4	NM_006282.5	-	-
TAP1	NM_000593.5	-	-
TAP2	NM_000544.3	-	-
TAPBP	NM_003190.4	-	-
TAZ (TAFAZZIN)	NM_000116.5	chrX:g.153641699G>A (c.284+110G>A)	-
TBX1	NM_080647.1	-	-
TFRC	NM_003234.4	-	-
TNFRSF4	NM_003327.4	-	-
TRAC	ENST00000611116.1	-	CNV analysis is not performed
TTC37 (SKIC3)	NM_014639.3	-	-
TTC7A	NM_020458.4	-	-
WAS	NM_000377.3	chrX:g.48547690_48547698delinsATCTGCAGACC (c.1339-19_1339-11delinsATCTGCAGACC)	-
WIPF1	NM_001077269.1	-	-
ZAP70	NM_001079.3	chr2:g.98349927G>A (c.838-80G>A); chr2:g.98354447G>A (c.1624-11G>A)	-
ZBTB24	NM_014797.2	-	-

Page 4 of 4 MC4091-449rev0625