| Gene     | RefSeq NM_ | Coding position | Protein           |
|----------|------------|-----------------|-------------------|
| FAS      | NM_000043.6| c.766G>C        | p.Glu256Gln       |
| FAS      | NM_000043.6| c.819G>T        | p.Gln273His       |
| CTLA4    | NM_005214.4| c.168_171delTG  | p.Cys585fs*13     |
| IKBKG    | NM_001099856.6| c.373G>A  | p.Glu125Lys       |
| CARD11   | NM_032415.7| c.2899C>T       | p.Arg967Cys       |
| ADA2     | NM_177405.3| c.559A>C        | p.Thr187Pro       |
| ADA2     | NM_177405.3| c.563T>C        | p.Leu188Pro       |
| STAT3    | NM_213662.2| c.1973A>G       | p.Lys658Arg       |
| LIG4     | NM_001098268.2| c.833G>A  | p.Arg278His homo  |
| LRBA     | NM_001367550.1| c.1963C>T  | p.Arg655Ter homo  |
| FAS      | NM_000043.6| c.385T>C        | p.Cys129Arg       |
| STAT3    | NM_001369512.1| c.454C>T   | p.Arg152Trp       |
| CARD11   | NM_032415.7| c.3025G>A       | p.Val1009Ile      |
| RAG1     | NM_001377280.1| c.1520G>A  | p.Arg507Gln homo  |
| TNFRSF13B| NM_012452.3| c.581_582delCCinsAA| p.Ser194Ter     |
| TNFRSF13B| NM_012452.3| c.605G>A        | p.Arg202His       |
| TNFRSF13B| NM_012452.3| c.171G>C        | p.Gln57His        |
| CASP10   | NM_032977.4| c.1228G>A       | p.Val410Ile       |

Coding positions of the found variants.