

**Table S1. 45 well characterized genes that were affected by a high-impact SNP.**

Effect Types: 1: Stop gained; 2: Splice site acceptor; 3: Splice site donor; 4: Stop lost; 5: Start lost.

| Symbol   | Descriptions                                   | Chr | Disorder/Disease   | Function  | Pathway   | Effect |
|----------|--|-----|--|---|---|--------|
| ABCA9    | ATP-binding cassette A9                        | 17  | Pseudoxanthoma elasticum   | Monocyte differentiation; Lipid homeostasis                 | ABC transporters  | 1      |
| ADCK3    | aaRF domain containing kinase 3                | 1   | Spinocerebellar ataxia   | Protein serine/threonine kinase activity                    |   | 1      |
| ANKRD35  | Ankyrin repeat containing protein 35           | 1   |  | Protein binding   |   | 1      |
| AQP7P3   | Aquaporin 7 pseudogene 3                       | 9   |  | Water transport   |   | 2      |
| CAD      | CAD trifunctional protein                      | 2   | Fibrosarcoma   | Aspartate carbamoyltransferase activity                     | Pyrimidine metabolism; Transcription/Ligand-dependent activation of ESR1/SP pathway   | 1      |
| CAST     | Calpain inhibitor                              | 5   | Vascular disease; Alzheimer's disease  | Endopeptidase inhibitor activity                            |   | 2      |
| CDC27    | cell division cycle 27                         | 17  |  | Cell cycle checkpoint                                       | Cell cycle_Regulation of G1/S transition  | 1      |
| CES1     | Carboxylesterase 1                             | 16  | CES 1 deficiency   | carboxylesterase activity                                   | Drug metabolism; Cholesterol and Sphingolipids transport  | 3      |
| CLTCL1   | Clathrin, heavy chain-like 1                   | 22  | Tetralogy of Fallot; Ventricular septal defect; triventricular septal defect; Pulmonary valve stenosis | Receptor-mediated endocytosis                               | Lysosome; Endocytosis; Endocrine-regulated calcium reabsorption; Huntington's disease; Bacterial invasion of epithelial cells | 3      |
| DEFB108B | Beta-defensin 8                                | 11  |  | defense response to bacterium                               |   | 5      |
| DPRX     | Divergent-paired related homeobox              | 19  |  | Sequence-specific DNA binding transcription factor activity |   | 1      |
| DST      | Dystonia musculorum protein                    | 6   | Bullous pemphigoid; Epidermolysis bullosa  | Microtubule cytoskeleton organization                       | Cytoskeleton remodeling, Keratin/Neuro filaments  | 2      |
| FAM104B  | Family with sequence similarity 104, member B  | X   |  |   |   | 3      |
| FRG1     | FSHD region gene-1                             | 4   | Facioscapulohumeral muscular dystrophy; Suppurative lymphadenitis                                      | Nuclear mRNA splicing via spliceosome                       |   | 3      |
| FRG1B    | FSHD region gene 1 family, member B            | 20  |  |   |   | 2      |
| FRG2C    | FSHD region gene 2 family, member C            | 3   |  |   |   | 1      |
| GIMAP6   | GTPase, IMAP family member 6                   | 7   |  | GTP binding   |   | 1      |
| HEATR7B2 | HEAT repeat family member 7B2                  | 5   |  | Binding   |   | 2      |
| HSPBAP1  | 27 kDa heat shock protein-associated protein 1 | 3   | Intractable epilepsy; Renal carcinoma  | Cellular stress response                                    |   | 1      |
| HTR2C    | 5-hydroxytryptamine receptor 1C                | X   | Schizophrenia; Migraine; Prader-Willi syndrome; Attention deficit hyperactivity                        | Phosphatidylinositol phospholipase C activity               | Calcium signaling pathway; Neuroactive ligand-receptor interaction  | 1      |

|          |   |    |  |  |  |   |
|----------|---|----|--|--|--|---|
| KBTBD3   | BTB and kelch domain-containing protein 3   | 11 |  | Protein binding  |  | 1 |
| KIAA0430 | Limkain-b1                                  | 16 |  | Nucleotide binding                                     |  | 2 |
| KRTAP2-2 | Keratin-associated protein 2.2              | 17 |  | Keratin filament                                       |  | 1 |
| MLL3     | Myeloid/lymphoid leukemia 3                 | 7  | Leukemia   | Methyltransferase activity                             | Lysine degradation   | 1 |
| MST1P2   | Macrophage stimulating 1 pseudogene 2       | 1  |  |  |  | 2 |
| MYT1     | Myelin transcription factor I               | 20 | Dysembryoplastic neuroepithelial tumor; Periventricular leukomalacia | Oligodendrocyte lineage development                    |  | 1 |
| NBPF1    | Neuroblastoma breakpoint family member 1    | 1  | Shipyards eye; Gastroenteritis                                       | Cytoplasm  |  | 4 |
| NBPF15   | Neuroblastoma breakpoint family, member 15  | 1  |  | Cytoplasm  |  | 2 |
| NOTCH2NL | Notch 2 N-terminal like                     | 1  | Alagille syndrome; Hajdu-Cheney syndrome; Glioma                     | Calcium ion binding                                    | Notch signaling pathway  | 3 |
| NUP153   | Nuclear pore complex protein                | 6  | Infectious mononucleosis; Hemophagocytic lymphohistiocytosis         | DNA binding  | Host Interactions of HIV factors                                       | 3 |
| OR8U8    | Olfactory receptor 8U8                      | 11 |  | G-protein coupled olfactory receptor activity          | Olfactory transduction   | 3 |
| PCNT     | Pericentrin                                 | 21 | Seckel syndrome; Microcephaly  | M transition of mitotic cell cycle                     | Centrosome maturation  | 1 |
| PPP1R32  | Protein phosphatase 1 regulatory subunit    | 11 |  | Phosphatase binding                                    |  | 2 |
| PPP2R2B  | Protein phosphatase 2, regulatory subunit B | 5  | Spinocerebellar ataxia   | Apoptotic process                                      | mRNA surveillance pathway; Tight junction; Regulation of CFTR activity | 1 |
| PROSER1  | Proline and serine rich 1                   | 13 |  |  |  | 1 |
| QRSL1    | Glutamyl-amidotransferase subunit A         | 6  | Lymphogranuloma venereum; Intestinal volvulus; Trachoma              | Carbon-nitrogen ligase activity                        |  | 2 |
| SAFB2    | Scaffold attachment factor B2               | 19 | Breast Cancer  | Nucleotide binding                                     |  | 2 |
| TBCK     | TBC1 domain containing kinase               | 4  |  |  |  | 1 |
| TCP10L2  | T-complex 10-like protein 2                 | 6  | Spina bifida   | Cytosol  |  | 1 |
| TECTA    | Tectorin alpha                              | 11 | Nonsyndromic deafness; Scotoma; Sensorineural hearing loss           | Cell-matrix adhesion                                   |  | 1 |
| TFAP2B   | Transcription factor AP-2 beta              | 6  | Patent ductus arteriosus; Skeletal muscle neoplasm                   | Cellular ammonia/urea/creatinine homeostasis           |  | 1 |
| TRIP12   | Thyroid receptor-interacting protein 12     | 2  | Chronic dacryocystitis; Heart disease                                | Ubiquitin-protein ligase activity                      | Ubiquitin mediated proteolysis   | 3 |
| WASH2P   | Protein FAM39B                              | 2  | Fission of tubules   |  |  | 3 |
| XIAP     | X-linked inhibitor of apoptosis protein     | X  | Leukemia; Lymphoma   | Caspases, apoptosis regulation; inflammatory signaling | Ubiquitin mediated proteolysis; SMAC-mediated apoptotic r.             | 1 |
| ZNF778   | Zinc finger protein 778                     | 16 | KBG syndrome; Learning disability                                    | Zinc ion binding                                       |  | 1 |