| Gene names | Protein names | Largest Cluster Res | Num in Largest |
|---|--|--------------------------|-------------------------|
| 1 ODC1 | Ornithine decarboxylase (ODC) (EC 4.1.1.17) | (kcal/mol) -10.880000 | Cluster 8.000000 |
| 2 AADAT KAT2 | Kynurenine/alpha-aminoadipate aminotransferase, mitochondrial (KAT/AadAT) (2-aminoadipate aminotransferase) (2-aminoadipate transaminase) (EC 2.6.1.39) (Alpha-aminoadipate aminotransferase) (AadAT) (Kynurenine aminotransferase II) (Kynurenineoxoglutarate transaminase 2) (EC 2.6.1.7) (Kynurenineoxoglutarate transaminase II) | -10.560000 | 30.000000 |
| 3 AGXT AGT1 SPAT | Serinepyruvate aminotransferase (SPT) (EC 2.6.1.51) (Alanineglyoxylate aminotransferase) (AGT) (EC 2.6.1.44) | -10.010000 | 28.000000 |
| 4 BCAT2 BCATM BCT2 ECA40 | Branched-chain-amino-acid aminotransferase, mitochondrial (BCAT(m)) (EC 2.6.1.42) (Placental protein 18) (PP18) | -9.800000 | 17.000000 |
| 5 GAD1 GAD GAD67 | Glutamate decarboxylase 1 (EC 4.1.1.15) (67 kDa glutamic acid decarboxylase) (GAD-67) (Glutamate decarboxylase 67 kDa isoform) | -9.480000 | 18.000000 |
| 6 SGPL1 KIAA1252 | Sphingosine-1-phosphate lyase 1 (S1PL) (SP-lyase 1) (SPL 1) (hSPL) (EC 4.1.2.27) (Sphingosine-1-phosphate aldolase) | -9.300000 | 30.000000 |
| 7 PYGL | Glycogen phosphorylase, liver form (EC 2.4.1.1) | -9.260000 | 17.000000 |
| 8 SEPSECS TRNP48 | O-phosphoseryl-tRNA(Sec) selenium transferase (EC 2.9.1.2) (Liver-pancreas antigen) (LP) (SLA-p35) (SLA/LP autoantigen) (Selenocysteine synthase) (Sec synthase) (Sec synthase) (Sep-tRNA:Sec-tRNA synthase) (SepSecS) (Soluble liver antigen) (SLA) (UGA suppressor tRNA-associated protein) (tRNA(Ser/Sec)-associated antigenic protein) | -8.950000 | 14.000000 |
| 9 KYAT3 CCBL2 KAT3 | Kynurenineoxoglutarate transaminase 3 (EC 2.6.1.7) (Cysteine-S-conjugate beta-lyase 2) (EC 4.4.1.13) (Kynurenine aminotransferase 3) (Kynurenine aminotransferase III) (KATIII) (Kynurenineglyoxylate transaminase) (EC 2.6.1.63) (Kynurenineoxoglutarate transaminase III) | -8.910000 | 32.000000 |
| 10 CTH 11 ACCS PHACS | Cystathionine gamma-lyase (EC 4.4.1.1) (Cysteine-protein sulfhydrase) (Gamma-cystathionase) 1-aminocyclopropane-1-carboxylate synthase-like protein 1 (ACC synthase-like protein 1) | -8.740000 -8.620000 | 28.000000 25.000000 |
| 12 OAT | Ornithine aminotransferase, mitochondrial (EC 2.6.1.13) (Ornithine delta-aminotransferase) (Ornithineoxo-acid aminotransferase) [Cleaved into: Ornithine aminotransferase, hepatic form; Ornithine aminotransferase, renal form] | -8.510000 | 15.000000 |
| $13_{\mathrm{ALT2}}^{\mathrm{GPT2}\ \mathrm{AAT2}}$ | Alanine aminotransferase 2 (ALT2) (EC 2.6.1.2) (Glutamate pyruvate transaminase 2) (GPT 2) (Glutamicalanine transaminase 2) (Glutamicpyruvic transaminase 2) | -8.360000 | 20.000000 |
| 14 SCLY SCL | Selenocysteine lyase (hSCL) (EC 4.4.1.16) | -8.310000 | 40.000000 |
| 15 ABAT GABAT | 4-aminobutyrate aminotransferase, mitochondrial (EC 2.6.1.19) ((S)-3-amino-2-methylpropionate transaminase) (EC 2.6.1.22) (GABA aminotransferase) (GABA-AT) (Gamma-amino-N-butyrate transaminase) (GABA transaminase) (GABA-T) (L-AIBAT) | -8.140000 | 61.000000 |
| 16 SHMT2 | Serine hydroxymethyltransferase, mitochondrial (SHMT) (EC 2.1.2.1) (Glycine hydroxymethyltransferase) (Serine methylase) | -8.100000 | 10.000000 |
| 17 SPTLC2 KIAA0526 LCB2 | Serine palmitoyltransferase 2 (EC 2.3.1.50) (Long chain base biosynthesis protein 2) (LCB 2) (Long chain base biosynthesis protein 2a) (LCB2a) (Serine-palmitoyl-CoA transferase 2) (SPT 2) | -7.990000 | 45.000000 |
| 18 ALAS2 ALASE ASB | 5-aminolevulinate synthase, erythroid-specific, mitochondrial (ALAS-E) (EC 2.3.1.37) (5-aminolevulinic acid synthase 2) (Delta-ALA synthase 2) (Delta-aminolevulinate synthase 2) | -7.970000 | 70.000000 |
| 19 GOT1 | Aspartate aminotransferase, cytoplasmic (cAspAT) (EC 2.6.1.1) (EC 2.6.1.3) (Cysteine aminotransferase, cytoplasmic) (Cysteine transaminase, cytoplasmic) (cCAT) (Glutamate oxaloacetate transaminase 1) (Transaminase A) | -7.800000 | 23.000000 |
| 20 GLDC GCSP | Glycine dehydrogenase (decarboxylating), mitochondrial (EC 1.4.4.2) (Glycine cleavage system P protein) (Glycine decarboxylase) (Glycine dehydrogenase (aminomethyl-transferring)) | -7.800000 | 18.000000 |
| 21 SDS SDH 22 SHMT1 | L-serine dehydratase/L-threonine deaminase (SDH) (EC 4.3.1.17) (L-serine deaminase) (L-threonine dehydratase) (TDH) (EC 4.3.1.19) Serine hydroxymethyltransferase, cytosolic (SHMT) (EC 2.1.2.1) (Glycine hydroxymethyltransferase) (Serine methylase) | -7.770000 -7.770000 | 15.000000 25.000000 |
| SPTLC3 23 C20orf38 | Serine palmitoyltransferase 3 (EC 2.3.1.50) (Long chain base biosynthesis protein 2b) (LCB2b) (Long chain base biosynthesis protein 3) (LCB 3) (Serine-palmitoyl-CoA transferase 3) (SPT | -7.560000 | 78.000000 |
| SPTLC2L ETNPPL AGXT2L1 | 3) Ethanolamine-phosphate phospho-lyase (EC 4.2.3.2) (Alanineglyoxylate aminotransferase 2-like 1) | -7.430000 | 43.000000 |
| AGXT2L1 25 CSAD CSD | Cysteine sulfinic acid decarboxylase (EC 4.1.1.29) (Aspartate 1-decarboxylase) (EC 4.1.1.11) (Cysteine-sulfinate decarboxylase) (Sulfinoalanine decarboxylase) | -7.390000 | 10.000000 |
| 26 DDC AADC | Aromatic-L-amino-acid decarboxylase (AADC) (EC 4.1.1.28) (DOPA decarboxylase) (DDC) | -7.240000 | 9.000000 |
| 27 KYAT1 CCBL1 | Kynurenineoxoglutarate transaminase 1 (EC 2.6.1.7) (Cysteine-S-conjugate beta-lyase) (EC 4.4.1.13) (Glutamine transaminase K) (GTK) (Glutaminephenylpyruvate transaminase) (EC 2.6.1.64) (Kynurenine aminotransferase 1) (Kynurenine aminotransferase I) (KATI) (Kynurenineoxoglutarate transaminase I) | -7.200000 | 41.000000 |
| ALAS1 ALAS3 28 ALASH OK/SW- cl.121 | 5-aminolevulinate synthase, nonspecific, mitochondrial (ALAS-H) (EC 2.3.1.37) (5-aminolevulinic acid synthase 1) (Delta-ALA synthase 1) (Delta-aminolevulinate synthase 1) | -7.170000 | 15.000000 |
| 29 GCAT KBL 30 SRR | 2-amino-3-ketobutyrate coenzyme A ligase, mitochondrial (AKB ligase) (EC 2.3.1.29) (Aminoacetone synthase) (Glycine acetyltransferase) Serine racemase (EC 5.1.1.18) (D-serine ammonia-lyase) (D-serine dehydratase) (EC 4.3.1.18) (L-serine ammonia-lyase) (EC 4.3.1.17) | -7.060000 -7.040000 | 21.000000 13.000000 |
| $31_{\mathrm{ECA39}}^{\mathrm{BCAT1}\ \mathrm{BCT1}}$ | Branched-chain-amino-acid aminotransferase, cytosolic (BCAT(c)) (EC 2.6.1.42) (Protein ECA39) | -6.990000 | 31.000000 |
| 32 TAT | Tyrosine aminotransferase (TAT) (EC 2.6.1.5) (L-tyrosine:2-oxoglutarate aminotransferase) | -6.920000 | 23.000000 |
| 33 HDC | Histidine decarboxylase (HDC) (EC 4.1.1.22) | -6.700000 | 28.000000 |
| 34 SDSL 35 ACCSL | Serine dehydratase-like (L-serine deaminase) (L-serine dehydratase/L-threonine deaminase) (L-threonine dehydratase) (TDH) (EC 4.3.1.19) (Serine dehydratase 2) (SDH 2) (EC 4.3.1.17) Probable inactive 1-aminocyclopropane-1-carboxylate synthase-like protein 2 (ACC synthase-like protein 2) | -6.680000 -6.550000 | 9.000000 11.000000 |
| 36 PYGM | Glycogen phosphorylase, muscle form (EC 2.4.1.1) (Myophosphorylase) | -6.540000 | 16.000000 |
| 37 MOCOS | Molybdenum cofactor sulfurase (MCS) (MOS) (MoCo sulfurase) (hMCS) (EC 2.8.1.9) (Molybdenum cofactor sulfurtransferase) | -6.430000 | 14.000000 |
| 38 GADL1 | Acidic amino acid decarboxylase GADL1 (Aspartate 1-decarboxylase) (ADC) (HuADC) (EC 4.1.1.11) (Cysteine sulfinic acid decarboxylase) (CSADC) (HuCSADC) (EC 4.1.1.29) (Glutamate decarboxylase-like protein 1) Threonine synthase-like 1 (TSH1) | -6.430000 | 17.000000 |
| 39 THNSL1 PHYKPL | Threofine Synthase-like 1 (15H1) | -6.260000 | 9.000000 |
| 40 AGXT2L2 PP9286 | 5-phosphohydroxy-L-lysine phospho-lyase (EC 4.2.3.134) (Alanineglyoxylate aminotransferase 2-like 2) | -6.230000 | 12.000000 |
| 41 GAD2 GAD65 | Glutamate decarboxylase 2 (EC 4.1.1.15) (65 kDa glutamic acid decarboxylase) (GAD-65) (Glutamate decarboxylase 65 kDa isoform) Aspartate aminotransferase, mitochondrial (mAspAT) (EC 2.6.1.1) (EC 2.6.1.7) (Fatty acid-binding protein) (FABP-1) (Glutamate oxaloacetate transaminase 2) (Kynurenine | -6.070000 | 5.000000 |
| 42 GOT2 | aminotransferase 4) (Kynurenine aminotransferase IV) (Kynurenineoxoglutarate transaminase 4) (Kynurenineoxoglutarate transaminase IV) (Plasma membrane-associated fatty acid-binding protein) (FABPpm) (Transaminase A) | -5.950000 | 12.000000 |
| 43 PLPBP PROSC 44 PYGB | Pyridoxal phosphate homeostasis protein (PLP homeostasis protein) (Proline synthase co-transcribed bacterial homolog protein) (Pyridoxal phosphate-binding protein) Glycogen phosphorylase, brain form (EC 2.4.1.1) | -5.940000 -5.900000 | 7.000000 17.000000 |
| 45 AGXT2 AGT2 | Alanineglyoxylate aminotransferase 2, mitochondrial (AGT 2) (EC 2.6.1.44) ((R)-3-amino-2-methylpropionatepyruvate transaminase) (EC 2.6.1.40) (Beta-ALAAT II) (Beta-alanine-pyruvate aminotransferase) (D-AIBAT) | -5.540000 | 17.000000 |
| 46 GPT AAT1 GPT1 | | -5.520000 | 47.000000 |
| 47 CBSL 48 PSAT1 PSA | Cystathionine beta-synthase-like protein (EC 4.2.1.22) (Beta-thionase) (Serine sulfhydrase) Phosphoserine aminotransferase (EC 2.6.1.52) (Phosphohydroxythreonine aminotransferase) (PSAT) | -5.470000 -5.270000 | 15.000000 7.000000 |
| 49 GOT1L1 | Putative aspartate aminotransferase, cytoplasmic 2 (EC 2.6.1.1) (Glutamate oxaloacetate transaminase 1-like protein 1) (Transaminase A-like protein 1) | -4.680000 | 7.000000 |
| $f{50}_{m{ m OAZIN}}^{m{ m AZIN1}\ m{ m OAZI}}$ | Antizyme inhibitor 1 (AZI) (AZI1) (Ornithine decarboxylase antizyme inhibitor) | -4.430000 | 7.000000 |
| 51 KYNU | Kynureninase (EC 3.7.1.3) (L-kynurenine hydrolase) | -3.970000 | 10.000000 |
| 52 THNSL2 | Threonine synthase-like 2 (TSH2) (EC 4.2.3) (Secreted osteoclastogenic factor of activated T-cells) (SOFAT) | -3.380000 | 14.000000 |

-3.380000

-3.350000

14.000000

9.000000

Threonine synthase-like 2 (TSH2) (EC 4.2.3.-) (Secreted osteoclastogenic factor of activated T-cells) (SOFAT)

Cysteine desulfurase, mitochondrial (EC 2.8.1.7)

52 THNSL2

53 NFS1 NIFS HUSSY-08