

Genes annotated to mmu00860

11656	Alas2; aminolevulinic acid synthase 2, erythroid [KO:K00643] [EC:2.3.1.37]
11655	Alas1; aminolevulinic acid synthase 1 [KO:K00643] [EC:2.3.1.37]
67417	Ears2; glutamyl-tRNA synthetase 2 (mitochondrial)(putative) [KO:K01885] [EC:6.1.1.17]
107508	Eprs; glutamyl-prolyl-tRNA synthetase [KO:K14163] [EC:6.1.1.15 6.1.1.17]
17025	Alad; aminolevulinate, delta-, dehydratase [KO:K01698] [EC:4.2.1.24]
15288	Hmbs; hydroxymethylbilane synthase [KO:K01749] [EC:2.5.1.61]
22276	Uros; uroporphyrinogen III synthase [KO:K01719] [EC:4.2.1.75]
22275	Urod; uroporphyrinogen decarboxylase [KO:K01599] [EC:4.1.1.37]
12892	Cpox; coproporphyrinogen oxidase [KO:K00228] [EC:1.3.3.3]
19044	Ppox; protoporphyrinogen oxidase [KO:K00231] [EC:1.3.3.4]
14151	Fech; ferrochelatase [KO:K01772] [EC:4.99.1.1]
70383	Cox10; cytochrome c oxidase assembly protein 10 [KO:K02257] [EC:2.5.1.-]
226139	Cox15; cytochrome c oxidase assembly protein 15 [KO:K02259]
77697	Mmab; methylmalonic aciduria (cobalamin deficiency) type B homolog (human) [KO:K00798] [EC:2.5.1.17]
15368	Hmox1; heme oxygenase (decycling) 1 [KO:K00510] [EC:1.14.99.3]
15369	Hmox2; heme oxygenase (decycling) 2 [KO:K00510] [EC:1.14.99.3]
109778	Blvra; biliverdin reductase A [KO:K00214] [EC:1.3.1.24]
233016	Blvrb; biliverdin reductase B (flavin reductase (NADPH)) [KO:K05901] [EC:1.5.1.30 1.3.1.24]
22238	Ugt2b5; UDP glucuronosyltransferase 2 family, polypeptide B5 [KO:K00699] [EC:2.4.1.17]
22236	Ugt1a2; UDP glucuronosyltransferase 1 family, polypeptide A2 [KO:K00699] [EC:2.4.1.17]
94284	Ugt1a6a; UDP glucuronosyltransferase 1 family, polypeptide A6A [KO:K00699] [EC:2.4.1.17]
94215	Ugt2a1; UDP glucuronosyltransferase 2 family, polypeptide A1 [KO:K00699] [EC:2.4.1.17]
394434	Ugt1a9; UDP glucuronosyltransferase 1 family, polypeptide A9 [KO:K00699] [EC:2.4.1.17]
394430	Ugt1a10; UDP glycosyltransferase 1 family, polypeptide A10 [KO:K00699] [EC:2.4.1.17]
394432	Ugt1a7c; UDP glucuronosyltransferase 1 family, polypeptide A7C [KO:K00699] [EC:2.4.1.17]
394433	Ugt1a5; UDP glucuronosyltransferase 1 family, polypeptide A5 [KO:K00699] [EC:2.4.1.17]
72094	Ugt2a3; UDP glucuronosyltransferase 2 family, polypeptide A3 [KO:K00699] [EC:2.4.1.17]
552899	Ugt2a2; UDP glucuronosyltransferase 2 family, polypeptide A2 [KO:K00699] [EC:2.4.1.17]
71773	Ugt2b1; UDP glucuronosyltransferase 2 family, polypeptide B1 [KO:K00699] [EC:2.4.1.17]
394435	Ugt1a6b; UDP glucuronosyltransferase 1 family, polypeptide A6B [KO:K00699] [EC:2.4.1.17]
394436	Ugt1a1; UDP glucuronosyltransferase 1 family, polypeptide A1 [KO:K00699] [EC:2.4.1.17]
100727	Ugt2b34; UDP glucuronosyltransferase 2 family, polypeptide B34 [KO:K00699] [EC:2.4.1.17]
231396	Ugt2b36; UDP glucuronosyltransferase 2 family, polypeptide B36 [KO:K00699] [EC:2.4.1.17]
100559	Ugt2b38; UDP glucuronosyltransferase 2 family, polypeptide B38 [KO:K00699] [EC:2.4.1.17]
112417	Ugt2b37; UDP glucuronosyltransferase 2 family, polypeptide B37 [KO:K00699] [EC:2.4.1.17]
243085	Ugt2b35; UDP glucuronosyltransferase 2 family, polypeptide B35 [KO:K00699] [EC:2.4.1.17]
110006	Gusb; glucuronidase, beta [KO:K01195] [EC:3.2.1.31]
15159	Hccs; holocytochrome c synthetase [KO:K01764] [EC:4.4.1.17]
12870	Cp; ceruloplasmin [KO:K13624] [EC:1.16.3.1]
14319	Fth1; ferritin heavy chain 1 [KO:K00522] [EC:1.16.3.2]
67634	Ftmt; ferritin mitochondrial [KO:K00522] [EC:1.16.3.2]
100862446	ferritin light chain 1-like; K13625 ferritin light chain [KO:K13625]
14325	Ft11; ferritin light chain 1 [KO:K13625]