Log₂ fold change -6 -4 -2 0 2 4 6

Log₂ fold changes of probes [Other Xenobiotic degradation; n = 5]

K01941 E6.3.4.6; urea carboxylase [EC:6.3.4.6]

K01782 fadJ; 3-hydroxyacyl-CoA dehydrogenase / enoyl-CoA hydratase / 3-hydroxybutyryl-CoA epimerase [EC:1.1.1.35 4.2.1.17 5.1.2.3]

K00121 frmA; S-(hydroxymethyl)glutathione dehydrogenase / alcohol dehydrogenase [EC:1.1.1.284 1.1.1.1]

K04100 ligA; protocatechuate 4,5-dioxygenase, alpha chain [EC:1.13.11.8]

K04101 ligB; protocatechuate 4,5-dioxygenase, beta chain [EC:1.13.11.8]

LL3Diel3DD3LL6Diel6DD6DD3LL3Diel3DD6LL6Diel6 SAR116_0936 protocatechuate 4,5-dioxygenase, beta chain

SAR116_1775 Allophanate hydrolase subunit 1 SAR116_0556 putative fatty acid oxidation complex alpha subunit SAR116_2012 Alcohol dehydrogenase GroES-like protein SAR116_0935 protocatechuate 4,5-dioxygenase, alpha chain