



Log₂ fold changes of probes

[Tyrosine metabolism; n = 13]

		$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$		
	K00812 aspB; aspartate aminotransferase [EC:2.6.1.1]							SAR116_2214	aminotransferase
	K00812 aspB; aspartate aminotransferase [EC:2.6.1.1]							SAR116_0974	aspartate aminotransferase
	K00813 aspC; aspartate aminotransferase [EC:2.6.1.1]							SAR116_0778	aspartate aminotransferase
	K00817 hisC; histidinol-phosphate aminotransferase [EC:2.6.1.9]							SAR116_0339	histidinol-phosphate aminotransferase
	K00817 hisC; histidinol-phosphate aminotransferase [EC:2.6.1.9]							SAR116_0442	histidinol-phosphate aminotransferase
	K00457 HPD; 4-hydroxyphenylpyruvate dioxygenase [EC:1.13.11.27]							SAR116_1810	putative 4-hydroxyphenylpyruvate dioxygenase protein
	K00451 HGD; homogentisate 1,2-dioxygenase [EC:1.13.11.5]							SAR116_1170	homogentisate 1,2-dioxygenase
	K16171 faaH; fumarylacetoacetate (FAA) hydrolase [EC:3.7.1.2]							SAR116_1171	putative aromatic amino acid degradation protein
	K00121 frmA; S-(hydroxymethyl)glutathione dehydrogenase / alcohol dehydrogenase [EC:1.1.1.284 1.1.1.1]							SAR116_2012	Alcohol dehydrogenase GroES-like protein
	K00483 hpaB; 4-hydroxyphenylacetate 3-monooxygenase [EC:1.14.14.9]							SAR116_0256	probable 4-hydroxyphenylacetic hydroxylase
	K10219 ligC; 2-hydroxy-4-carboxymuconate semialdehyde hemiacetal dehydrogenase [EC:1.1.1.312]							SAR116_0937	oxidoreductase domain protein
	K00135 gabD; succinate-semialdehyde dehydrogenase / glutarate-semialdehyde dehydrogenase [EC:1.2.1.16 1.2.1.79 1.2.1.20]							SAR116_1424	succinate-semialdehyde dehydrogenase protein
	K16165 nagK; fumarylpyruvate hydrolase [EC:3.7.1.20]							SAR116_1465	fumarylacetoacetate (FAA) hydrolase