Log$_2$ fold changes of probes
[Pyruvate metabolism; n = 33]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K01895 ACSS1_2; acetyl–CoA synthetase [EC:6.2.1.1]							SAR116_1878 Acetate––CoA ligase
K00161 PDHA; pyruvate dehydrogenase E1 component alpha subunit [EC:1.2.4.1]							SAR116_0205 Pyruvate dehydrogenase (acetyl–transferring)
K00162 PDHB; pyruvate dehydrogenase E1 component beta subunit [EC:1.2.4.1]							SAR116_0206 Pyruvate dehydrogenase beta subunit
K00627 DLAT; pyruvate dehydrogenase E2 component (dihydrolipoamide acetyltransferase) [EC:2.3.1.12]							SAR116_0207 Dihydrolipoamide acetyltransferase
K00382 DLD; dihydrolipoamide dehydrogenase [EC:1.8.1.4]							SAR116_1836 dihydrolipoamide dehydrogenase
K00382 DLD; dihydrolipoamide dehydrogenase [EC:1.8.1.4]							SAR116_0208 Pyruvate/2–oxoglutarate dehydrogenase complex
K00121 frmA; S–(hydroxymethyl)glutathione dehydrogenase / alcohol dehydrogenase [EC:1.1.1.284 1.1.1.1]							SAR116_2012 Alcohol dehydrogenase GroES–like protein
K00625 E2.3.1.8; phosphate acetyltransferase [EC:2.3.1.8]							SAR116_2025 Phosphate acetyltransferase
K00873 PK; pyruvate kinase [EC:2.7.1.40]							SAR116_1437 pyruvate kinase
K01962 accA; acetyl–CoA carboxylase carboxyl transferase subunit alpha [EC:6.4.1.2 2.1.3.15]							SAR116_1916 Acetyl–CoA carboxylase alpha subunit
K02160 accB; acetyl–CoA carboxylase biotin carboxyl carrier protein							SAR116_1039 acetyl–CoA carboxylase
K02160 accB; acetyl–CoA carboxylase biotin carboxyl carrier protein							SAR116_1773 biotin/lipoyl attachment
K01961 accC; acetyl–CoA carboxylase, biotin carboxylase subunit [EC:6.4.1.2 6.3.4.14]							SAR116_1040 acetyl–CoA carboxylase
K01961 accC; acetyl–CoA carboxylase, biotin carboxylase subunit [EC:6.4.1.2 6.3.4.14]							SAR116_1774 acetyl CoA carboxylase
K01963 accD; acetyl–CoA carboxylase carboxyl transferase subunit beta [EC:6.4.1.2 2.1.3.15]							SAR116_1258 Acetyl–CoA carboxylase carboxyl transferase
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_0754 aldehyde dehydrogenase family protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_1715 Betaine–aldehyde dehydrogenase
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_2169 Betaine–aldehyde dehydrogenase
K01026 pct; propionate CoA–transferase [EC:2.8.3.1]							SAR116_0923 coenzyme A transferase
K00101 lldD; L–lactate dehydrogenase (cytochrome) [EC:1.1.2.3]							SAR116_0741 L–lactate dehydrogenase
K00101 lldD; L–lactate dehydrogenase (cytochrome) [EC:1.1.2.3]							SAR116_0771 L–lactate dehydrogenase
K01759 GLO1; lactoylglutathione lyase [EC:4.4.1.5]							SAR116_2143 Glyoxalase/bleomycin resistance protein/dioxygenase
K01069 gloB; hydroxyacylglutathione hydrolase [EC:3.1.2.6]							SAR116_2142 Hydroxyacylglutathione hydrolase
K12972 ghrA; glyoxylate/hydroxypyruvate reductase [EC:1.1.1.79 1.1.1.81]							SAR116_1484 putative oxidoreductase
K00029 E1.1.1.40; malate dehydrogenase (oxaloacetate–decarboxylating)(NADP+) [EC:1.1.1.40]							SAR116_1600 malate dehydrogenase
K00024 mdh; malate dehydrogenase [EC:1.1.1.37]							SAR116_1831 Malate dehydrogenase
K01676 E4.2.1.2A; fumarate hydratase, class I [EC:4.2.1.2]							SAR116_0723 hydro–lyase
K01595 ppc; phosphoenolpyruvate carboxylase [EC:4.1.1.31]							SAR116_0912 Putative phosphoenolpyruvate carboxylase
K01006 ppdK; pyruvate, orthophosphate dikinase [EC:2.7.9.1]							SAR116_2277 pyruvate, water dikinase
K01638 aceB; malate synthase [EC:2.3.3.9]							SAR116_1669 malate synthase G
K00626 ACAT; acetyl–CoA C–acetyltransferase [EC:2.3.1.9]							SAR116_0063 beta–ketothiolase
K01649 leuA; 2–isopropylmalate synthase [EC:2.3.3.13]							SAR116_1223 2–isopropylmalate synthase
K01649 leuA; 2–isopropylmalate synthase [EC:2.3.3.13]							SAR116_0677 2–isopropylmalate synthase/homocitrate synthase

Log$_2$ fold change

−6   −4   −2   0   2   4   6