



Log₂ fold changes of probes
[Glycolysis / Gluconeogenesis; n = 24]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K00845 glk; glucokinase [EC:2.7.1.2]							SAR116_1985 glucokinase
K01810 GPI; glucose-6-phosphate isomerase [EC:5.3.1.9]							SAR116_2454 Glucose-6-phosphate isomerase
K03841 FBP; fructose-1,6-bisphosphatase I [EC:3.1.3.11]							SAR116_1413 fructose-1,6-bisphosphatase I
K01624 FBA; fructose-bisphosphate aldolase, class II [EC:4.1.2.13]							SAR116_1412 Fructose-bisphosphate aldolase
K01624 FBA; fructose-bisphosphate aldolase, class II [EC:4.1.2.13]							SAR116_0281 fructose-bisphosphate aldolase
K01803 TPI; triosephosphate isomerase (TIM) [EC:5.3.1.1]							SAR116_0198 Triosephosphate isomerase
K00134 GAPDH; glyceraldehyde 3-phosphate dehydrogenase (phosphorylating) [EC:1.2.1.12]							SAR116_0883 glyceraldehyde-3-phosphate dehydrogenase
K00134 GAPDH; glyceraldehyde 3-phosphate dehydrogenase (phosphorylating) [EC:1.2.1.12]							SAR116_2285 glyceraldehyde-3-phosphate dehydrogenase
K00927 PGK; phosphoglycerate kinase [EC:2.7.2.3]							SAR116_0729 Phosphoglycerate kinase
K15633 gpml; 2,3-bisphosphoglycerate-independent phosphoglycerate mutase [EC:5.4.2.12]							SAR116_1848 Phosphoglycerate mutase
K01689 ENO; enolase [EC:4.2.1.11]							SAR116_0203 Phosphopyruvate hydratase
K00873 PK; pyruvate kinase [EC:2.7.1.40]							SAR116_1437 pyruvate kinase
K01006 ppdK; pyruvate, orthophosphate dikinase [EC:2.7.9.1]							SAR116_2277 pyruvate, water dikinase
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
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
K01895 ACSS1_2; acetyl-CoA synthetase [EC:6.2.1.1]							SAR116_1878 Acetate--CoA ligase
K01785 galM; aldose 1-epimerase [EC:5.1.3.3]							SAR116_0583 aldose 1-epimerase