



Log₂ fold changes of probes
[Citrate cycle (TCA cycle); n = 19]

	<div><div>LL₃</div><div>DD₃</div></div>	<div><div>LD₃</div><div>LL₃</div></div>	<div><div>DD₃</div><div>LD₃</div></div>	<div><div>LL₆</div><div>DD₆</div></div>	<div><div>LD₆</div><div>LL₆</div></div>	<div><div>DD₆</div><div>LD₆</div></div>	
K01647 CS; citrate synthase [EC:2.3.3.1]							SAR116_0504 citrate synthase I
K01681 ACO; aconitate hydratase [EC:4.2.1.3]							SAR116_1075 Aconitate hydratase 1
K00031 IDH1; isocitrate dehydrogenase [EC:1.1.1.42]							SAR116_0090 Isocitrate dehydrogenase
K00030 IDH3; isocitrate dehydrogenase (NAD+) [EC:1.1.1.41]							SAR116_1739 probable 3-isopropylmalate dehydrogenase oxidoreductase protein
K00164 OGDH; 2-oxoglutarate dehydrogenase E1 component [EC:1.2.4.2]							SAR116_1834 2-oxoglutarate dehydrogenase, E1
K00658 DLST; 2-oxoglutarate dehydrogenase E2 component (dihydrolipoamide succinyltransferase) [EC:2.3.1.61]							SAR116_1835 2-oxoglutarate dehydrogenase, E2
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
K01902 sucD; succinyl-CoA synthetase alpha subunit [EC:6.2.1.5]							SAR116_1833 succinyl-CoA synthetase, alpha
K01903 sucC; succinyl-CoA synthetase beta subunit [EC:6.2.1.5]							SAR116_1832 succinyl-CoA synthetase, beta
K00239 sdhA; succinate dehydrogenase / fumarate reductase, flavoprotein subunit [EC:1.3.5.1 1.3.5.4]							SAR116_1828 succinate dehydrogenase, flavoprotein subunit
K00240 sdhB; succinate dehydrogenase / fumarate reductase, iron-sulfur subunit [EC:1.3.5.1 1.3.5.4]							SAR116_1829 succinate dehydrogenase and fumarate reductase, iron-sulfur protein
K00241 sdhC; succinate dehydrogenase / fumarate reductase, cytochrome b subunit							SAR116_1826 succinate dehydrogenase, cytochrome b subunit
K00242 sdhD; succinate dehydrogenase / fumarate reductase, membrane anchor subunit							SAR116_1827 succinate dehydrogenase, membrane anchor subunit
K01676 E4.2.1.2A; fumarate hydratase, class I [EC:4.2.1.2]							SAR116_0723 hydro-lyase;fumarate hydratase, class I
K00024 mdh; malate dehydrogenase [EC:1.1.1.37]							SAR116_1831 Malate dehydrogenase
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