



Log₂ fold changes of probes
[Taurine and phosphonate metabolism; n = 6]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K00681 ggt; gamma-glutamyltranspeptidase / glutathione hydrolase [EC:2.3.2.2 3.4.19.13]							SAR116_1348 gamma-glutamyltranspeptidase
K03119 tauD; taurine dioxygenase [EC:1.14.11.17]							SAR116_0893 putative dioxygenase
K00259 ald; alanine dehydrogenase [EC:1.4.1.1]							SAR116_2542 alanine dehydrogenase
K03852 xsc; sulfoacetaldehyde acetyltransferase [EC:2.3.3.15]							SAR116_1367 hypothetical protein
K00625 E2.3.1.8; phosphate acetyltransferase [EC:2.3.1.8]							SAR116_2025 Phosphate acetyltransferase
K06167 phnP; phosphoribosyl 1,2-cyclic phosphate phosphodiesterase [EC:3.1.4.55]							SAR116_0360 beta-lactamase-like protein