



Log<sub>2</sub> fold changes of probes  
[Taurine and phosphonate metabolism; n = 6]

	<u>LL<sub>3</sub></u> DD <sub>3</sub>	<u>Diel<sub>3</sub></u> LL <sub>3</sub>	<u>DD<sub>3</sub></u> Diel <sub>3</sub>	<u>LL<sub>6</sub></u> DD <sub>6</sub>	<u>Diel<sub>6</sub></u> LL <sub>6</sub>	<u>DD<sub>6</sub></u> Diel <sub>6</sub>	
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