



Log₂ fold changes of probes
[DMSP metabolism; n = 10]

	<u>LL₃</u> DD ₃	<u>Diel₃</u> LL ₃	<u>DD₃</u> Diel ₃	<u>LL₆</u> DD ₆	<u>Diel₆</u> LL ₆	<u>DD₆</u> Diel ₆	
							SAR116_1427 peptidase M24 (EC:3.4.13.9)
K01271 pepQ; Xaa-Pro dipeptidase [EC:3.4.13.9]							SAR116_0769 metallopeptidase, family M24
							SAR116_0023 cupin 2 barrel domain-containing protein (EC:2.7.7.22)
Cupin 2 conserved barrel domain protein							SAR116_0050 Cupin 2 conserved barrel domain protein
							SAR116_1790 potential Cupin domain-containing protein (EC:5.3.1.9)
K17486 dmdA; dimethylsulfoniopropionate demethylase [EC:2.1.1.269]							SAR116_0074 aminomethyl transferase (EC:2.1.2.10)
							SAR116_0916 glycine cleavage T protein (aminomethyl transferase) (EC:2.1.2.10)
K00605 gcvT; aminomethyltransferase [EC:2.1.2.10]							SAR116_0235 aminomethyltransferase (EC:2.1.2.10)
K00605 gcvT; aminomethyltransferase [EC:2.1.2.10]							SAR116_1344 aminomethyltransferase (glycine cleavage system T protein) (EC:2.1.2.10)
K01620 ltaE; threonine aldolase [EC:4.1.2.48]							SAR116_2172 aminomethyl transferase (EC:2.1.2.10)