Log$_2$ normalized probes
[DMSP metabolism; n = 10]

	DD$_3$	LL$_3$	Diel$_3$	DD$_6$	LL$_6$	Diel$_6$	
dddP							SAR116_1427 peptidase M24 (EC:3.4.13.9)
dddP homolog							SAR116_0769 metallopeptidase, family M24
dddQ							SAR116_0023 cupin 2 barrel domain−containing protein (EC:2.7.7.22)
dddQ homolog							SAR116_0050 Cupin 2 conserved barrel domain protein
dddW							SAR116_1790 potential Cupin domain−containing protein (EC:5.3.1.9)
dmdA							SAR116_0074 aminomethyl transferase (EC:2.1.2.10)
dmdA homolog							SAR116_0916 glycine cleavage T protein (aminomethyl transferase) (EC:2.1.2.10)
dmdA homolog							SAR116_0235 aminomethyltransferase (EC:2.1.2.10)
dmdA homolog							SAR116_1344 aminomethyltransferase (glycine cleavage system T protein) (EC:2.1.2.10)
dmdA homolog							SAR116_2172 aminomethyl transferase (EC:2.1.2.10)

Log$_2$ probe abundance

−20   −15   −10   −5