Log$_2$ probe abundance

-20 -15 -10 -5

Log$_2$ normalized probes
[Chlorohydrocarbon degradation; n = 7]

	DD$_3$	LL$_3$	Diel$_3$	DD$_6$	LL$_6$	Diel$_6$	
K00121 frmA; S−(hydroxymethyl)glutathione dehydrogenase / alcohol dehydrogenase [EC:1.1.1.284 1.1.1.1]							SAR116_2012 Alcohol dehydrogenase GroES−like protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_0754 aldehyde dehydrogenase family protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_1715 Betaine−aldehyde dehydrogenase
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_2169 Betaine−aldehyde dehydrogenase
K01561 dehH; haloacetate dehalogenase [EC:3.8.1.3]							SAR116_0318 putative hydrolase or acyltransferase
K01856 catB; muconate cycloisomerase [EC:5.5.1.1]							SAR116_1527 mandelate racemase/muconate lactonizing enzyme
K01561 dehH; haloacetate dehalogenase [EC:3.8.1.3]							SAR116_0318 putative hydrolase or acyltransferase