Log$_2$ fold changes of probes

[Toxin; n = 26]

| | LL$_3$ | Diel$_3$ | DD$_3$ | LL$_6$ | Diel$_6$ | DD$_6$ | |
	DD$_3$	LL$_3$	Diel$_3$	DD$_6$	LL$_6$	Diel$_6$	
							SAR116_0006 hypothetical protein
K09013 sufC; Fe–S cluster assembly ATP–binding protein							SAR116_0015 FeS assembly ATPase SufC
K03499 trkA; trk/ktr system potassium uptake protein							SAR116_0219 K+ transport systems, NAD–binding component
K03116 tatA; sec–independent protein translocase protein TatA							SAR116_0415 twin–arginine translocation protein, TatA/E family subunit
K00817 hisC; histidinol–phosphate aminotransferase [EC:2.6.1.9]							SAR116_0442 histidinol–phosphate aminotransferase
							SAR116_0444 hypothetical protein
K05568 mnhD; multicomponent Na+:H+ antiporter subunit D							SAR116_0460 NADH dehydrogenase (quinone)
K02234 cobW; cobalamin biosynthesis protein CobW							SAR116_0572 cobalamin biosynthesis protein CobW
K03192 ureJ; urease accessory protein							SAR116_0573 hydrogenase accessory protein
K03321 TC.SULP; sulfate permease, SulP family							SAR116_0878 sulfate permease family protein
K01686 uxuA; mannonate dehydratase [EC:4.2.1.8]							SAR116_0904 mannonate hydrolase
							SAR116_0933 putative major facilitator family (MFS) transporter
K01963 accD; acetyl–CoA carboxylase carboxyl transferase subunit beta [EC:6.4.1.2 2.1.3.15]							SAR116_1258 Acetyl–CoA carboxylase carboxyl transferase, beta subunit
K01918 panC; pantoate––beta–alanine ligase [EC:6.3.2.1]							SAR116_1350 pantoate–beta–alanine ligase
							SAR116_1353 Integrase
K07402 xdhC; xanthine dehydrogenase accessory factor							SAR116_1378 protein of unknown function DUF182
K02029 ABC.PA.P; polar amino acid transport system permease protein							SAR116_1445 polar amino acid uptake family ABC transporter, permease protein
K06217 phoH; phosphate starvation–inducible protein PhoH and related proteins							SAR116_1589 PhoH family protein
K01995 livG; branched–chain amino acid transport system ATP–binding protein							SAR116_1785 ABC transporter permease protein
							SAR116_2079 hypothetical protein
							SAR116_2124 acyltransferase domain protein
K03640 pal; peptidoglycan–associated lipoprotein							SAR116_2300 Peptidoglycan–associated lipoprotein
K07171 mazF; mRNA interferase MazF [EC:3.1.–.–]							SAR116_2309 hypothetical protein
K03631 recN; DNA repair protein RecN (Recombination protein N)							SAR116_2325 ATPase involved in DNA repair
							SAR116_2360 Rhodanese–like protein
K23536 nupC; general nucleoside transport system permease protein							SAR116_2468 inner–membrane translocator

Log$_2$ fold change

−6 −4 −2 0 2 4 6