



Log₂ fold changes of probes
[Polynucleotides; n = 12]

	LL ₃	LD ₃	DD ₃	LL ₆	LD ₆	DD ₆	
	DD ₃	LL ₃	LD ₃	DD ₆	LL ₆	LD ₆	
							SAR116_0091 DNA recombination–mediator protein A (TIGR00730 family Rossman fold protein)
K03978 engB; GTP–binding protein							SAR116_1501 ribosome biogenesis GTP–binding protein YihA/YsxC
K03574 mutT; 8–oxo–dGTP diphosphatase [EC:3.6.1.55]							SAR116_1921 NUDIX hydrolase
							SAR116_2183 probable MutT/nudix family protein
K02316 dnaG; DNA primase [EC:2.7.7.101]							SAR116_2232 DNA primase
K06204 dksA; DnaK suppressor protein							SAR116_0086 dksA
K03060 rpoZ; DNA–directed RNA polymerase subunit omega [EC:2.7.7.6]							SAR116_0537 RpoZ/omega
K08311 nudH; putative (di)nucleoside polyphosphate hydrolase [EC:3.6.1.–]							SAR116_1846 RNA pyrophosphohydrolase
K03086 rpoD; RNA polymerase primary sigma factor							SAR116_2233 RpoD/sigma 70
K03043 rpoB; DNA–directed RNA polymerase subunit beta [EC:2.7.7.6]							SAR116_2384 RpoB
K03046 rpoC; DNA–directed RNA polymerase subunit beta' [EC:2.7.7.6]							SAR116_2385 RpoC
K03040 rpoA; DNA–directed RNA polymerase subunit alpha [EC:2.7.7.6]							SAR116_2415 RpoA