Log$_2$ normalized probes
[Nucleotide metabolism; n = 26]

Columns: DD$_3$, LL$_3$, LD$_3$, DD$_6$, LL$_6$, LD$_6$

Log$_2$ probe abundance
−20 −15 −10 −5

Left label	Right label
	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
K01139 spoT; GTP diphosphokinase / guanosine−3',5'−bis(diphosphate) 3'−diphosphatase [EC:2.7.6.5 3.1.7.2]	SAR116_0536 SpoT/RelA, (p)ppGpp synthetase I (GTP pyrophosphokinase)
K00942 gmk; guanylate kinase [EC:2.7.4.8]	SAR116_0636 Guanylate kinase
	SAR116_1601 Adenosine kinase
K00759 APRT; adenine phosphoribosyltransferase [EC:2.4.2.7]	SAR116_1668 Adenine phosphoribosyl transferase
K00764 purF; amidophosphoribosyltransferase [EC:2.4.2.14]	SAR116_0651 purF; Glutamine phosphoribosylpyrophosphate amidotransferase
K01756 purB; adenylosuccinate lyase [EC:4.3.2.2]	SAR116_0667 Adenylosuccinate lyase
K01923 purC; phosphoribosylaminoimidazole−succinocarboxamide synthase [EC:6.3.2.6]	SAR116_0668 Phosphoribosylaminoimidazole−succinocarboxamide synthase
K00945 cmk; CMP/dCMP kinase [EC:2.7.4.25]	SAR116_1251 Cytidylate kinase
K00760 hprT; hypoxanthine phosphoribosyltransferase [EC:2.4.2.8]	SAR116_1426 HGPRT
K00769 gpt; xanthine phosphoribosyltransferase [EC:2.4.2.22]	SAR116_1505 Xanthine−guanine phosphoribosyltransferase
K00761 upp; uracil phosphoribosyltransferase [EC:2.4.2.9]	SAR116_1611 Uracil phosphoribosyltransferase
K01524 ppx−gppA; exopolyphosphatase / guanosine−5'−triphosphate,3'−diphosphate pyrophosphatase [EC:3.6.1.11 3.6.1.40]	SAR116_1993 Ppx/GppA phosphatase
K01939 purA; adenylosuccinate synthase [EC:6.3.4.4]	SAR116_2308 Adenylosuccinate synthase
K00088 IMPDH; IMP dehydrogenase [EC:1.1.1.205]	SAR116_2483 Inosine−5'−monophosphate dehydrogenase
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
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
K08311 nudH; putative (di)nucleoside polyphosphate hydrolase [EC:3.6.1.−]	SAR116_1846 RNA pyrophosphohydrolase