Log₂ normalized probes [Anderson, et al. 2021; n = 27]

	DD ₃	LL ₃	Diel ₃	DD ₆	LL ₆ [Diel ₆
K01939 purA; adenylosuccinate synthase [EC:6.3.4.4]					SAR116_2308 adenylosuccinate synthase
K03979 obgE; GTPase [EC:3.6.5]					- SAR116_1854 ObgE
K01139 spoT; GTP diphosphokinase / guanosine-3',5'-bis(diphosphate) 3'-diphosphatase [EC:2.7.6.5 3.1.7.2]					SAR116_0536 SpoT/RelA
K01524 ppx-gppA; exopolyphosphatase / guanosine-5'-triphosphate,3'-diphosphate pyrophosphatase [EC:3.6.1.11 3.6.1.40]					SAR116_1993 Ppx/GppA phosphatase
K02316 dnaG; DNA primase [EC:2.7.7.101 Log ₂ probe abundance] -					SAR116_2232 DNA primase
K02355 fusA; elongation factor 0	à -					SAR116_2388 translation elongation factor G
−20 −15 −10 −5 K00942 gmk; guanylate kinase [EC:2.7.4.8] -					SAR116_0636 Guanylate kinase
K06207 typA; GTP-binding protein	1					SAR116_2002 TypA
K00769 gpt; xanthine phosphoribosyltransferase [EC:2.4.2.22]					- SAR116_1505 Xgpt
K06204 dksA; DnaK suppressor protein	1					- SAR116_0086 dksA
K03086 rpoD; RNA polymerase primary sigma facto	r -					SAR116_2233 RpoD/sigma 70
K03060 rpoZ; DNA-directed RNA polymerase subunit omega [EC:2.7.7.6]					SAR116_0537 RpoZ/omega
K03046 rpoC; DNA-directed RNA polymerase subunit beta' [EC:2.7.7.6] -					- SAR116_2385 RpoC
K03043 rpoB; DNA-directed RNA polymerase subunit beta [EC:2.7.7.6] -					SAR116_2384 RpoB
K03040 rpoA; DNA-directed RNA polymerase subunit alpha [EC:2.7.7.6] -					SAR116_2415 RpoA
K00764 purF; amidophosphoribosyltransferase [EC:2.4.2.14] -					SAR116_0651 purF; Glutamine phosphoribosylpyrophosphate amidotransferase
K03665 hflX; GTPase	> -					SAR116_0793 GTP-binding protein HflX
K06942 ychF; ribosome-binding ATPase	3 -					SAR116_2191 redox-regulated ATPase YchF
	_					SAR116_0091 DNA recombination-mediator protein A (TIGR00730 family Rossman fold protein)
K00760 hprT; hypoxanthine phosphoribosyltransferase [EC:2.4.2.8]					- SAR116_1426 HGPRT
K08311 nudH; putative (di)nucleoside polyphosphate hydrolase [EC:3.6.1] -					SAR116_1846 RNA pyrophosphohydrolase
						SAR116_0584 AKR_Tas-like
K19265 gpr; L-glyceraldehyde 3-phosphate reductase [EC:1.1.1]					SAR116_2446 GPR/YghZ/AKR14A1
	_					SAR116_0604 Aldo-keto reductase (AKR) superfamily
	-					SAR116_0247 AKR_BsYcsN_EcYdhF-like
						SAR116_1601 Adenosine kinase
K00759 APRT; adenine phosphoribosyltransferase [EC:2.4.2.7] -					SAR116_1668 Adenine phosphoribosyl transferase