Log$_2$ probe abundance

−20 −15 −10 −5

Log$_2$ normalized probes

[Riboflavin metabolism; n = 6]

	DD$_3$	LL$_3$	Diel$_3$	DD$_6$	LL$_6$	Diel$_6$	
K01497 ribA; GTP cyclohydrolase II [EC:3.5.4.25]							SAR116_1657 GTP cyclohydrolase II
K14652 ribBA; 3,4−dihydroxy 2−butanone 4−phosphate synthase / GTP cyclohydrolase II [EC:4.1.99.12 3.5.4.25]							SAR116_0165 3,4−dihydroxy 2−butanone 4−phosphate synthase / GTP cyclohydrolase II
K11752 ribD; diaminohydroxyphosphoribosylaminopyrimidine deaminase / 5−amino−6−(5−phosphoribosylamino)uracil reductase [EC:3.5.4.26 1.1.1.193]							SAR116_0163 bifunctional deaminase−reductase
K00794 ribH; 6,7−dimethyl−8−ribityllumazine synthase [EC:2.5.1.78]							SAR116_0166 riboflavin synthase
K00793 ribE; riboflavin synthase [EC:2.5.1.9]							SAR116_0164 Lumazine−binding protein
K11753 ribF; riboflavin kinase / FMN adenylyltransferase [EC:2.7.1.26 2.7.7.2]							SAR116_1145 FMN adenylyltransferase