Log$_2$ normalized probes
[ATP generation; n = 12]

	DD$_3$	LL$_3$	Diel$_3$	DD$_6$	LL$_6$	Diel$_6$	
K21034 virH2; cytochrome P450 family 104							SAR116_2093 Cytochrome P450
K02639 petF; ferredoxin							SAR116_0299 ferredoxin
K02114 ATPF1E; F−type H+−transporting ATPase subunit epsilon							SAR116_1845 H+−transporting two−sector ATPase
K02112 ATPF1B; F−type H+/Na+−transporting ATPase subunit beta [EC:7.1.2.2 7.2.2.1]							SAR116_1844 F1−ATP Synthase Beta Chain
K02115 ATPF1G; F−type H+−transporting ATPase subunit gamma							SAR116_1843 ATP synthase F1
K02111 ATPF1A; F−type H+/Na+−transporting ATPase subunit alpha [EC:7.1.2.2 7.2.2.1]							SAR116_1842 ATP synthase F1
K02113 ATPF1D; F−type H+−transporting ATPase subunit delta							SAR116_1841 H+−transporting two−sector ATPase
K02109 ATPF0B; F−type H+−transporting ATPase subunit b							SAR116_1103 ATP synthase subunit B'
K02109 ATPF0B; F−type H+−transporting ATPase subunit b							SAR116_1104 ATP synthase B chain precursor
K02110 ATPF0C; F−type H+−transporting ATPase subunit c							SAR116_1102 ATP synthase C chain
K02108 ATPF0A; F−type H+−transporting ATPase subunit a							SAR116_1101 H+−transporting two−sector ATPase
K02116 atpI; ATP synthase protein I							SAR116_1100 ATP synthase protein I

Log$_2$ probe abundance

−20 −15 −10 −5