Log₂ probe abundance

-20 -15 -10 -5

[Oxidative phosphorylation; n = 39]							
	DD ₃	LL ₃	LD ₃	DD ₆	LL ₆	LD ₆	7
K00330 nuoA; NADH-quinone oxidoreductase subunit A [EC:7.1.1.2]							SAR116_0141 NADH:ubiquinone oxidoreductase subunit 3
K00331 nuoB; NADH-quinone oxidoreductase subunit B [EC:7.1.1.2]							SAR116_0142 NADH:ubiquinone oxidoreductase 20 kD subunit and related Fe-S oxidoreductase
K00332 nuoC; NADH-quinone oxidoreductase subunit C [EC:7.1.1.2]							SAR116_0143 NADH (or F420H2) dehydrogenase
K00333 nuoD; NADH-quinone oxidoreductase subunit D [EC:7.1.1.2]							SAR116_0144 NADH:ubiquinone oxidoreductase 49 kD subunit 7
K00334 nuoE; NADH-quinone oxidoreductase subunit E [EC:7.1.1.2]							SAR116_0145 NADH dehydrogenase
K00335 nuoF; NADH-quinone oxidoreductase subunit F [EC:7.1.1.2]							SAR116_0146 NADH-quinone oxidoreductase
K00336 nuoG; NADH-quinone oxidoreductase subunit G [EC:7.1.1.2]							SAR116_0147 NADH-quinone oxidoreductase
K00337 nuoH; NADH-quinone oxidoreductase subunit H [EC:7.1.1.2]							SAR116_0148 Respiratory–chain NADH dehydrogenase
K00338 nuol; NADH-quinone oxidoreductase subunit I [EC:7.1.1.2]							SAR116_0149 NADH-quinone oxidoreductase
K00339 nuoJ; NADH–quinone oxidoreductase subunit J [EC:7.1.1.2]							SAR116_0150 NADH–ubiquinone/plastoquinone oxidoreductase chain 6
K00340 nuoK; NADH-quinone oxidoreductase subunit K [EC:7.1.1.2]							SAR116_0151 NADH-ubiquinone oxidoreductase chain 4L
K00341 nuoL; NADH-quinone oxidoreductase subunit L [EC:7.1.1.2]							SAR116_0152 NADH:ubiquinone oxidoreductase subunit 5
K00342 nuoM; NADH-quinone oxidoreductase subunit M [EC:7.1.1.2]							SAR116_0153 Proton-translocating NADH-quinone oxidoreductase
K00343 nuoN; NADH-quinone oxidoreductase subunit N [EC:7.1.1.2]							SAR116_0154 Proton-translocating NADH-quinone oxidoreductase
K00239 sdhA; succinate dehydrogenase / fumarate reductase, flavoprotein subunit [EC:1.3.5.1 1.3.5.4]							SAR116_1828 succinate dehydrogenase flavoprotein subunit
K00240 sdhB; succinate dehydrogenase / fumarate reductase, iron-sulfur subunit [EC:1.3.5.1 1.3.5.4]							SAR116_1829 succinate dehydrogenase and fumarate reductase iron-sulfur protein
K00241 sdhC; succinate dehydrogenase / fumarate reductase, cytochrome b subunit							SAR116_1826 succinate dehydrogenase subunit C
K00242 sdhD; succinate dehydrogenase / fumarate reductase, membrane anchor subunit							SAR116_1827 succinate dehydrogenase
K00411 UQCRFS1; ubiquinol-cytochrome c reductase iron-sulfur subunit [EC:7.1.1.8]							SAR116_2187 ubiquinol–cytochrome c reductase
K00412 CYTB; ubiquinol-cytochrome c reductase cytochrome b subunit							SAR116_2188 Cytochrome b
K00413 CYC1; ubiquinol-cytochrome c reductase cytochrome c1 subunit							SAR116_2189 cytochrome c1
K02257 COX10; heme o synthase [EC:2.5.1.141]							SAR116_1118 putative protoheme IX farnesyltransferase (heme o synthase)
K02276 coxC; cytochrome c oxidase subunit III [EC:7.1.1.9]							SAR116_1115 cytochrome–c oxidase
K02274 coxA; cytochrome c oxidase subunit I [EC:7.1.1.9]							SAR116_1119 cytochrome–c oxidase
K02275 coxB; cytochrome c oxidase subunit II [EC:7.1.1.9]							SAR116_1120 cytochrome C oxidase subunit II
K02258 COX11; cytochrome c oxidase assembly protein subunit 11							SAR116_1116 cytochrome C oxidase assembly protein
K02259 COX15; heme a synthase [EC:1.17.99.9]							SAR116_1891 Uncharacterized protein required for cytochrome oxidase assembly
K08738 CYC; cytochrome c							SAR116_1486 cytochrome C
K08738 CYC; cytochrome c							SAR116_0742 cytochrome c family protein
K02111 ATPF1A; F-type H+/Na+-transporting ATPase subunit alpha [EC:7.1.2.2 7.2.2.1]							SAR116_1842 ATP synthase F1
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
K02113 ATPF1D; F-type H+-transporting ATPase subunit delta							SAR116_1841 H+-transporting two-sector ATPase
K02114 ATPF1E; F-type H+-transporting ATPase subunit epsilon							SAR116_1845 H+-transporting two-sector ATPase
K02108 ATPF0A; F-type H+-transporting ATPase subunit a							SAR116_1101 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
K22468 ppk2; polyphosphate kinase [EC:2.7.4.34]							SAR116_0942 polyphosphate kinase