LL<sub>3</sub> LD<sub>3</sub> DD<sub>3</sub> LL<sub>6</sub> LD<sub>6</sub> DD<sub>6</sub> DD<sub>3</sub> LL<sub>3</sub> LD<sub>3</sub> DD<sub>6</sub> LL<sub>6</sub> LD<sub>6</sub> K24821 atm1; ATP-binding cassette, subfamily B, heavy metal transporter SAR116\_0020 ABC transporter ATP-binding protein/permease K10112 msmX; multiple sugar transport system ATP-binding protein [EC:7.5.2.-] SAR116\_0044 ABC transporter ATP-binding protein K02031 ddpD; peptide/nickel transport system ATP-binding protein SAR116\_0064 ABC transporter ATP-binding protein SAR116\_0271 ATP-binding cassette domain-containing protein K02056 ABC.SS.A; simple sugar transport system ATP-binding protein [EC:7.5.2.-] K23064 IhpO; hydroxyproline transport system ATP-binding protein [EC:7.4.2.1] SAR116\_0306 amino acid ABC transporter ATP-binding protein K09972 aapP; general L-amino acid transport system ATP-binding protein [EC:7.4.2.1] SAR116\_0332 amino acid ABC transporter ATP-binding protein SAR116\_0349 ATP-binding cassette domain-containing protein K01995 livG; branched-chain amino acid transport system ATP-binding protein K01996 livF; branched-chain amino acid transport system ATP-binding protein SAR116\_0350 ABC transporter ATP-binding protein K02010 afuC; iron(III) transport system ATP-binding protein [EC:7.2.2.7] SAR116\_0416 ABC transporter ATP-binding protein K09972 aapP; general L-amino acid transport system ATP-binding protein [EC:7.4.2.1] SAR116\_0448 amino acid ABC transporter ATP-binding protein K09810 lolD; lipoprotein-releasing system ATP-binding protein [EC:7.6.2.-] SAR116\_0521 ABC transporter ATP-binding protein K03595 era; GTPase SAR116\_0530 GTPase Era SAR116\_0591 ATP-binding cassette domain-containing protein K06857 tupC; tungstate transport system ATP-binding protein [EC:7.3.2.6] K02031 ddpD; peptide/nickel transport system ATP-binding protein SAR116\_0600 ABC transporter ATP-binding protein SAR116\_0608 ATP-binding cassette domain-containing protein K01990 ABC-2.A; ABC-2 type transport system ATP-binding protein SAR116\_0648 ATP-binding cassette domain-containing protein K02065 mlaF; phospholipid/cholesterol/gamma-HCH transport system ATP-binding protein K02052 ABC.SP.A; putative spermidine/putrescine transport system ATP-binding protein SAR116\_0761 spermidine/putrescine ABC transporter ATPase subunit K01996 livF; branched-chain amino acid transport system ATP-binding protein SAR116\_0781 ABC transporter ATP-binding protein K01995 livG; branched-chain amino acid transport system ATP-binding protein SAR116\_0782 ABC transporter ATP-binding protein SAR116\_0793 GTPase HflX K03665 hflX; GTPase SAR116\_0888 ABC transporter ATP-binding protein K17325 glpT; glycerol transport system ATP-binding protein K17324 glpS; glycerol transport system ATP-binding protein SAR116\_0889 ABC transporter ATP-binding protein K01990 ABC-2.A; ABC-2 type transport system ATP-binding protein SAR116\_0894 ABC transporter ATP-binding protein K01996 livF; branched-chain amino acid transport system ATP-binding protein SAR116\_0928 ABC transporter ATP-binding protein K01995 livG; branched-chain amino acid transport system ATP-binding protein SAR116\_0932 ABC transporter ATP-binding protein SAR116\_0948 ATP-binding cassette domain-containing protein K02031 ddpD; peptide/nickel transport system ATP-binding protein SAR116\_0949 ABC transporter ATP-binding protein K02028 ABC.PA.A; polar amino acid transport system ATP-binding protein [EC:7.4.2.1] SAR116\_0959 amino acid ABC transporter ATP-binding protein K10111 malK; multiple sugar transport system ATP-binding protein [EC:7.5.2.-] SAR116\_0967 ABC transporter ATP-binding protein K02193 ccmA; heme exporter protein A [EC:7.6.2.5] SAR116\_1076 heme ABC exporter ATP-binding protein CcmA SAR116\_1130 ABC transporter ATP-binding protein/permease K11085 msbA; ATP-binding cassette, subfamily B, bacterial MsbA [EC:7.5.2.6] K12541 lapB; ATP-binding cassette, subfamily C, bacterial LapB SAR116\_1168 ATP-binding cassette domain-containing protein K10111 malK; multiple sugar transport system ATP-binding protein [EC:7.5.2.-] SAR116\_1192 sn-glycerol-3-phosphate ABC transporter ATP-binding protein UgpC K02036 pstB; phosphate transport system ATP-binding protein [EC:7.3.2.1] SAR116\_1203 phosphate ABC transporter ATP-binding protein PstB K06861 lptB; lipopolysaccharide export system ATP-binding protein [EC:7.5.2.5] SAR116\_1247 LPS export ABC transporter ATP-binding protein K06925 tsaE; tRNA threonylcarbamoyladenosine biosynthesis protein TsaE SAR116\_1263 tRNA (adenosine(37)–N6)–threonylcarbamoyltransferase complex ATPase subunit type 1 TsaE K01990 ABC-2.A; ABC-2 type transport system ATP-binding protein SAR116\_1270 ABC transporter ATP-binding protein K02003 ABC.CD.A; putative ABC transport system ATP-binding protein SAR116\_1418 ATP-binding cassette domain-containing protein K10831 tauB; taurine transport system ATP-binding protein [EC:7.6.2.7] SAR116\_1440 probable taurin uptake ABC transporter ATP-binding protein K02028 ABC.PA.A; polar amino acid transport system ATP-binding protein [EC:7.4.2.1] SAR116\_1446 glutamine ABC transporter, ATP-binding protein K01995 livG; branched-chain amino acid transport system ATP-binding protein SAR116\_1461 ABC transporter ATP-binding protein K01996 livF; branched-chain amino acid transport system ATP-binding protein SAR116\_1462 ABC transporter ATP-binding protein K03978 engB; GTP-binding protein SAR116\_1501 ribosome biogenesis GTP-binding protein YihA/YsxC K10112 msmX; multiple sugar transport system ATP-binding protein [EC:7.5.2.-] SAR116\_1512 sn-glycerol-3-phosphate ABC transporter ATP-binding protein UgpC K23537 nupA; general nucleoside transport system ATP-binding protein SAR116\_1607 ABC transporter ATP-binding protein K03650 mnmE; tRNA modification GTPase [EC:3.6.-.-] SAR116\_1645 tRNA uridine-5-carboxymethylaminomethyl(34) synthesis GTPase MnmE K02052 ABC.SP.A; putative spermidine/putrescine transport system ATP-binding protein SAR116\_1724 ABC transporter ATP-binding protein K01996 livF; branched-chain amino acid transport system ATP-binding protein SAR116\_1758 ABC transporter ATP-binding protein K01995 livG; branched-chain amino acid transport system ATP-binding protein SAR116\_1759 ATP-binding cassette domain-containing protein SAR116\_1784 ABC transporter ATP-binding protein SAR116\_1785 ABC transporter ATP-binding protein K01995 livG; branched-chain amino acid transport system ATP-binding protein K02052 ABC.SP.A; putative spermidine/putrescine transport system ATP-binding protein SAR116\_1793 ABC transporter ATP-binding protein SAR116\_1854 GTPase ObgE K03979 obgE; GTPase [EC:3.6.5.-] K24846 pglK; ATP-binding cassette, subfamily B, bacterial PglK [EC:7.5.2.-] SAR116\_1954 ABC transporter ATP-binding protein/permease SAR116\_1968 ABC transporter ATP-binding protein K15600 thiZ; putative hydroxymethylpyrimidine transport system ATP-binding protein K11076 potG; putrescine transport system ATP-binding protein [EC:7.6.2.16] SAR116\_2032 ABC transporter ATP-binding protein K06158 ABCF3; ATP-binding cassette, subfamily F, member 3 SAR116\_2053 ABC-F family ATP-binding cassette domain-containing protein K25152 gldA; gliding motility–associated transport system ATP–binding protein SAR116\_2095 ABC transporter ATP-binding protein K15555 ssuB; sulfonate transport system ATP-binding protein [EC:7.6.2.14] SAR116\_2097 ABC transporter ATP-binding protein K15555 ssuB; sulfonate transport system ATP-binding protein [EC:7.6.2.14] SAR116\_2099 ABC transporter ATP-binding protein K06147 ABCB-BAC; ATP-binding cassette, subfamily B, bacterial SAR116\_2279 ABC transporter transmembrane domain-containing protein SAR116\_2318 energy-dependent translational throttle protein EttA K06020 ettA; energy-dependent translational throttle protein EttA K15738 uup; ABC transport system ATP-binding/permease protein SAR116\_2347 ABC-F family ATP-binding cassette domain-containing protein K23537 nupA; general nucleoside transport system ATP-binding protein SAR116\_2466 ABC transporter ATP-binding protein SAR116\_2501 ribosome biogenesis GTPase Der

SAR116\_0406 protein translocase subunit SecD [Candidatus Puniceispirillum marinum]

SAR116\_2191 redox-regulated ATPase YchF [Candidatus Puniceispirillum marinum]

K03072 secD; preprotein translocase subunit SecD

K06942 ychF; ribosome-binding ATPase

Log<sub>2</sub> fold changes of probes

[obgE; n = 67]

