## Log<sub>2</sub> normalized probes [ABC transporters (3); n = 31]

	DD <sub>3</sub>	LL <sub>3</sub>	Diel <sub>3</sub>	DD <sub>6</sub>	LL <sub>6</sub>	Diel <sub>6</sub>	
K07335 bmpA; basic membrane protein A and related proteins							SAR116_1606 membrane lipoprotein
K15554 ssuC; sulfonate transport system permease protein							SAR116_2098 nitrate ABC transporter
K15555 ssuB; sulfonate transport system ATP-binding protein [EC:7.6.2.14]	_						SAR116_2097 ABC-type nitrate/sulfonate/bicarbonate transport system ATPase component-like protein
K15553 ssuA; sulfonate transport system substrate-binding protein							SAR116_2101 ABC-type nitrate/sulfonate/bicarbonate transport systems periplasmic component-like protein
K11074 potl; putrescine transport system permease protein							SAR116_2034 Ornithine carbamoyltransferase
K09815 znuA; zinc transport system substrate-binding protein							SAR116_2227 periplasmic solute binding protein
K10227 smoE; polyol transport system substrate-binding protein							SAR116_0970 periplasmic sorbitol–binding protein
K11720 lptG; lipopolysaccharide export system permease protein							SAR116_0632 permease YjgP/YjgQ
K02037 pstC; phosphate transport system permease protein	_						SAR116_1205 phosphate ABC transporter
K02038 pstA; phosphate transport system permease protein							SAR116_1204 phosphate ABC transporter permease protein
K02040 pstS; phosphate transport system substrate-binding protein							SAR116_1206 phosphate binding protein
K02036 pstB; phosphate transport system ATP-binding protein [EC:7.3.2.1]							SAR116_1203 ABC-type phosphate transporter ATP-binding protein
K02066 mlaE; phospholipid/cholesterol/gamma–HCH transport system permease protein							SAR116_2535 Protein of unknown function DUF140
K01999 livK; branched-chain amino acid transport system substrate-binding protein							SAR116_1786 putative ABC transport system
K23061 lhpP; hydroxyproline transporter system substrate-binding protein							SAR116_0309 putative ABC transporter periplasmic amino acid–binding protein
K02001 proW; glycine betaine/proline transport system permease protein							SAR116_1390 putative permease
K23062 lhpM; hydroxyproline transport system permease protein							SAR116_0308 putative permease of ABC transporter
K11076 potG; putrescine transport system ATP-binding protein [EC:7.6.2.16]							SAR116_2032 spermidine/putrescine ABC transporter ATPase subunit
K02002 proX; glycine betaine/proline transport system substrate-binding protein	_						SAR116_1391 Substrate-binding region of ABC-type glycine betaine transport system
K10112 msmX; multiple sugar transport system ATP-binding protein [EC:7.5.2]							SAR116_0044 sugar ABC transporter
K17325 glpT; glycerol transport system ATP-binding protein							SAR116_0888 sugar ABC transporter
K17324 glpS; glycerol transport system ATP-binding protein							SAR116_0889 sugar ABC transporter
K17321 glpV; glycerol transport system substrate-binding protein	-						SAR116_0890 sugar ABC transporter
K10228 smoF; polyol transport system permease protein							SAR116_0969 sugar ABC transporter
K05773 tupB; tungstate transport system permease protein							SAR116_0590 sulfate/tungstate uptake family ABC transporter
K05772 tupA; tungstate transport system substrate-binding protein							SAR116_0592 sulfate/tungstate uptake family ABC transporter
K10831 tauB; taurine transport system ATP-binding protein [EC:7.6.2.7]							SAR116_1440 probable taurin uptake ABC transporter ATP-binding protein
K15551 tauA; taurine transport system substrate-binding protein							SAR116_1439 taurine transport system periplasmic protein
K15552 tauC; taurine transport system permease protein							SAR116_1441 probable taurine uptake ABC transporter permease protein
K07323 mlaC; phospholipid transport system substrate-binding protein							SAR116_2319 toluene tolerance family protein
K23536 nupC; general nucleoside transport system permease protein							SAR116_1609 uncharacterized ABC-type transport system

Log<sub>2</sub> probe abundance

-20 -15 -10 -5