## $Log_2$ normalized probes [Quorum sensing (1); n = 39]

		[Quorum sensing (1), n = 39]					
		DD <sub>3</sub>	LL <sub>3</sub> Die	el <sub>3</sub> DE	D <sub>6</sub> L	L <sub>6</sub> Diel <sub>6</sub>	
	K02031 ddpD; peptide/nickel transport system ATP-binding protein						SAR116_0064 ABC transporter ATP-binding protein
	K02034 ABC.PE.P1; peptide/nickel transport system permease protein						SAR116_0065 hypothetical ABC transporter permease
	K02033 ABC.PE.P; peptide/nickel transport system permease protein						SAR116_0066 peptide ABC transporter permease protein-like protein
	K02035 ABC.PE.S; peptide/nickel transport system substrate-binding protein						SAR116_0067 hypothetical ABC transporter
	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
Log <sub>2</sub> probe	K01897 ACSL; long-chain acyl-CoA synthetase [EC:6.2.1.3]						SAR116_0193 Putative long-chain-fatty-acidCoA ligase (Long-chain acyl-CoA synthetase)
	K01657 trpE; anthranilate synthase component I [EC:4.1.3.27]						SAR116_0195 Anthranilate/para-aminobenzoate synthase component I
	K03075 secG; preprotein translocase subunit SecG						SAR116_0200 preprotein translocase SecG subunit
	K03666 hfq; host factor–I protein						SAR116_0220 RNA-binding protein
	abundance K01658 trpG; anthranilate synthase component II [EC:4.1.3.27]						SAR116_0232 Glutamine amidotransferase of anthranilate synthase
	K01999 livK; branched-chain amino acid transport system substrate-binding protein						SAR116_0346 high affinity branched–chain amino acid ABC transporter
	K01997 livH; branched–chain amino acid transport system permease protein						SAR116_0347 branched–chain amino acid permease protein
	K01998 livM; branched–chain amino acid transport system permease protein						- SAR116_0348 inner–membrane translocator
	K01995 livG; branched–chain amino acid transport system ATP–binding protein						SAR116_0349 ABC transporter related protein
	K01996 livF; branched–chain amino acid transport system ATP–binding protein –						SAR116_0350 branched–chain amino acid ABC transporter
	K03210 yajC; preprotein translocase subunit YajC –						- SAR116_0407 YajC
	K02031 ddpD; peptide/nickel transport system ATP-binding protein						SAR116_0600 putative ABC transporter
	K02034 ABC.PE.P1; peptide/nickel transport system permease protein						SAR116_0601 Binding-protein-dependent transport systems inner membrane component
	K02033 ABC.PE.P; peptide/nickel transport system permease protein						SAR116_0602 binding-protein-dependent transport systems inner membrane component
	K02035 ABC.PE.S; peptide/nickel transport system substrate-binding protein						SAR116_0603 extracellular solute–binding protein family 5
	K01626 E2.5.1.54; 3-deoxy-7-phosphoheptulonate synthase [EC:2.5.1.54]						SAR116_0682 3-deoxy-7-phosphoheptulonate synthase
	K02053 ABC.SP.P; putative spermidine/putrescine transport system permease protein						SAR116_0758 binding–protein–dependent transport systems inner membrane component
	K02054 ABC.SP.P1; putative spermidine/putrescine transport system permease protein						SAR116_0759 polyamine ABC transporter
	K02055 ABC.SP.S; putative spermidine/putrescine transport system substrate-binding protein						SAR116_0760 polyamine ABC trasnporter
	K02052 ABC.SP.A; putative spermidine/putrescine transport system ATP-binding protein						SAR116_0761 spermidine/putrescine ABC transporter ATPase subunit
	K01998 livM; branched–chain amino acid transport system permease protein						SAR116_0779 inner–membrane translocator
	K01997 livH; branched–chain amino acid transport system permease protein						SAR116_0780 inner–membrane translocator
	K01996 livF; branched–chain amino acid transport system ATP–binding protein						SAR116_0781 ABC transporter related protein
	K01995 livG; branched–chain amino acid transport system ATP–binding protein						SAR116_0782 ABC transporter related protein
	K01999 livK; branched–chain amino acid transport system substrate–binding protein –						SAR116_0783 branched–chain amino acid ABC transporter
	K01996 livF; branched–chain amino acid transport system ATP–binding protein –						SAR116_0928 ABC transporter related protein
	K01997 livH; branched–chain amino acid transport system permease protein						SAR116_0929 branched–chain amino acid ABC transporter
	K01999 livK; branched–chain amino acid transport system substrate–binding protein						SAR116_0930 Extracellular ligand–binding receptor
	K01998 livM; branched–chain amino acid transport system permease protein						SAR116_0931 inner–membrane translocator
	K01995 livG; branched–chain amino acid transport system ATP–binding protein						SAR116_0932 high affinity branched–chain amino acid ABC transporter
	K02031 ddpD; peptide/nickel transport system ATP-binding protein						SAR116_0949 Oligopeptide/dipeptide ABC transporter
	K02034 ABC.PE.P1; peptide/nickel transport system permease protein						SAR116_0950 peptide ABC transporter
	K02033 ABC.PE.P; peptide/nickel transport system permease protein						SAR116_0951 binding-protein-dependent transport systems inner membrane component
	K02035 ABC.PE.S; peptide/nickel transport system substrate-binding protein						SAR116_0952 extracellular solute-binding protein
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