Log$_2$ normalized probes

[Carbohydrates/Lipids/Amino Acids; n = 17]

	DD$_3$	LL$_3$	LD$_3$	DD$_6$	LL$_6$	LD$_6$	
K00031 IDH1; isocitrate dehydrogenase [EC:1.1.1.42]							SAR116_0090 Isocitrate dehydrogenase
							SAR116_0247 AKR_BsYcsN_EcYdhF−like
							SAR116_0304 6−phosphogluconate dehydrogenase, NAD−binding protein
							SAR116_0584 AKR_Tas−like
							SAR116_0604 Aldo−keto reductase (AKR) superfamily
K00927 PGK; phosphoglycerate kinase [EC:2.7.2.3]							SAR116_0729 Phosphoglycerate kinase
K01595 ppc; phosphoenolpyruvate carboxylase [EC:4.1.1.31]							SAR116_0912 Putative phosphoenolpyruvate carboxylase
K03841 FBP; fructose−1,6−bisphosphatase I [EC:3.1.3.11]							SAR116_1413 fructose−1,6−bisphosphatase I
K15633 gpmI; 2,3−bisphosphoglycerate−independent phosphoglycerate mutase [EC:5.4.2.12]							SAR116_1848 Phosphoglycerate mutase
K01929 murF; UDP−N−acetylmuramoyl−tripeptide−−D−alanyl−D−alanine ligase [EC:6.3.2.10]							SAR116_2338 UDP−N−acetylmuramoylalanyl−D−glutamyl−2
							SAR116_2436 Phosphoglycerate mutase
K01963 accD; acetyl−CoA carboxylase carboxyl transferase subunit beta [EC:6.4.1.2 2.1.3.15]							SAR116_1258 Acetyl−CoA carboxylase carboxyl transferase
K01716 fabA; 3−hydroxyacyl−[acyl−carrier protein] dehydratase / trans−2−decenoyl−[acyl−carrier protein] isomerase [EC:4.2.1.59 5.3.3.14]							SAR116_1572 3−hydroxydecanoyl−ACP dehydratase
K00765 hisG; ATP phosphoribosyltransferase [EC:2.4.2.17]							SAR116_0443 ATP phosphoribosyltransferase HisG
K00261 GLUD1_2; glutamate dehydrogenase (NAD(P)+) [EC:1.4.1.3]							SAR116_0453 Glu/Leu/Phe/Val dehydrogenase
K11754 folC; dihydrofolate synthase / folylpolyglutamate synthase [EC:6.3.2.12 6.3.2.17]							SAR116_1259 FolC bifunctional protein
K00020 HIBADH; 3−hydroxyisobutyrate dehydrogenase [EC:1.1.1.31]							SAR116_1425 3−hydroxyisobutyate dehydrogenase [EC:1.1.1.31]

Log$_2$ probe abundance

−20   −15   −10   −5