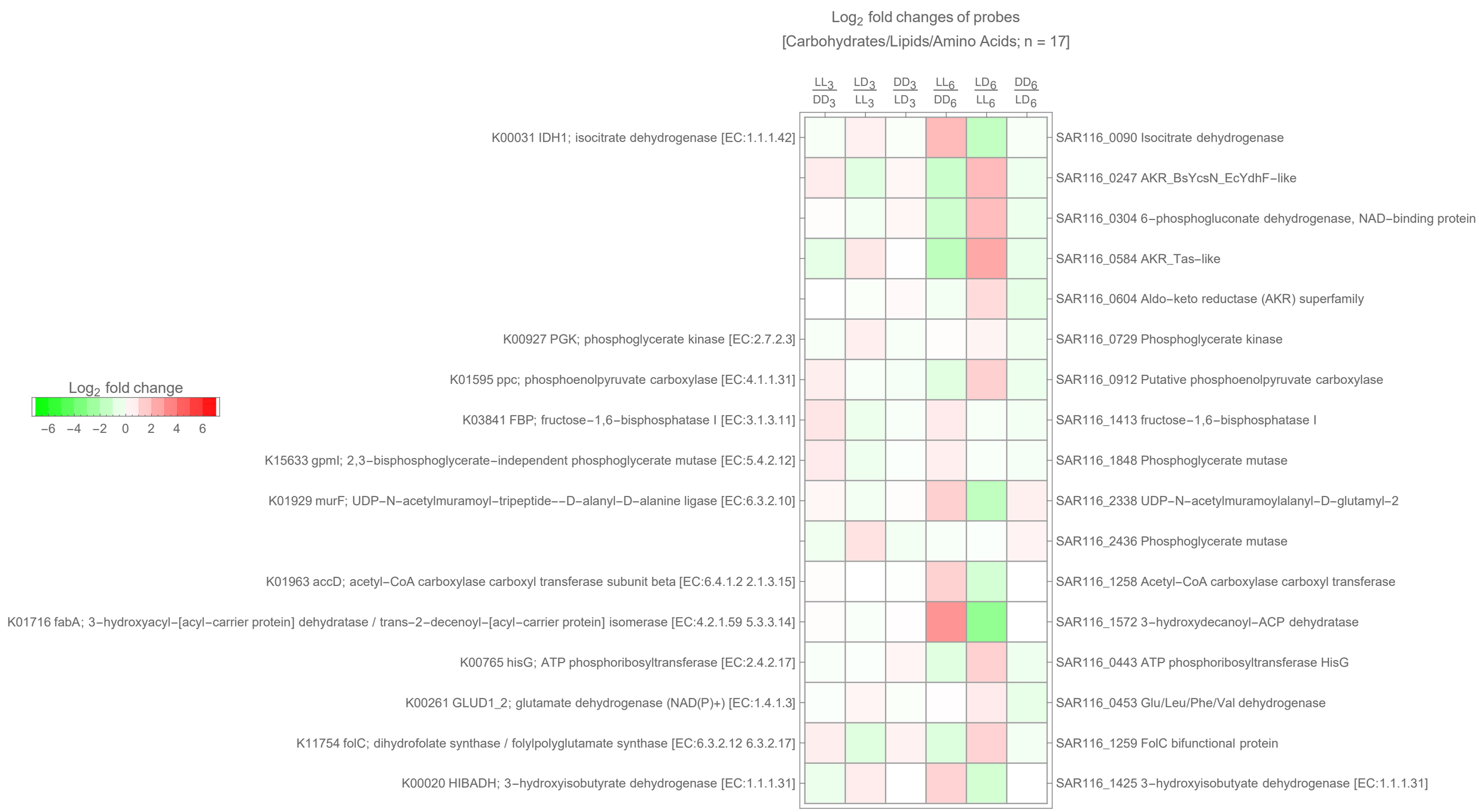
Log2 fold changes of probes
[Carbohydrates/Lipids/Amino Acids; n = 17]
LL3/DD3
LD3/LL3
DD3/LD3
LL6/DD6
LD6/LL6
DD6/LD6
K00031 IDH1; isocitrate dehydrogenase [EC:1.1.1.42]
K00927 PGK; phosphoglycerate kinase [EC:2.7.2.3]
K01595 ppc; phosphoenolpyruvate carboxylase [EC:4.1.1.31]
K03841 FBP; fructose−1,6−bisphosphatase I [EC:3.1.3.11]
K15633 gpmI; 2,3−bisphosphoglycerate−independent phosphoglycerate mutase [EC:5.4.2.12]
K01929 murF; UDP−N−acetylmuramoyl−tripeptide−−D−alanyl−D−alanine ligase [EC:6.3.2.10]
K01963 accD; acetyl−CoA carboxylase carboxyl transferase subunit beta [EC:6.4.1.2 2.1.3.15]
K01716 fabA; 3−hydroxyacyl−[acyl−carrier protein] dehydratase / trans−2−decenoyl−[acyl−carrier protein] isomerase [EC:4.2.1.59 5.3.3.14]
K00765 hisG; ATP phosphoribosyltransferase [EC:2.4.2.17]
K00261 GLUD1_2; glutamate dehydrogenase (NAD(P)+) [EC:1.4.1.3]
K11754 folC; dihydrofolate synthase / folylpolyglutamate synthase [EC:6.3.2.12 6.3.2.17]
K00020 HIBADH; 3−hydroxyisobutyrate dehydrogenase [EC:1.1.1.31]
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
SAR116_0729 Phosphoglycerate kinase
SAR116_0912 Putative phosphoenolpyruvate carboxylase
SAR116_1413 fructose−1,6−bisphosphatase I
SAR116_1848 Phosphoglycerate mutase
SAR116_2338 UDP−N−acetylmuramoylalanyl−D−glutamyl−2
SAR116_2436 Phosphoglycerate mutase
SAR116_1258 Acetyl−CoA carboxylase carboxyl transferase
SAR116_1572 3−hydroxydecanoyl−ACP dehydratase
SAR116_0443 ATP phosphoribosyltransferase HisG
SAR116_0453 Glu/Leu/Phe/Val dehydrogenase
SAR116_1259 FolC bifunctional protein
SAR116_1425 3−hydroxyisobutyate dehydrogenase [EC:1.1.1.31]
Log2 fold change
−6 −4 −2 0 2 4 6