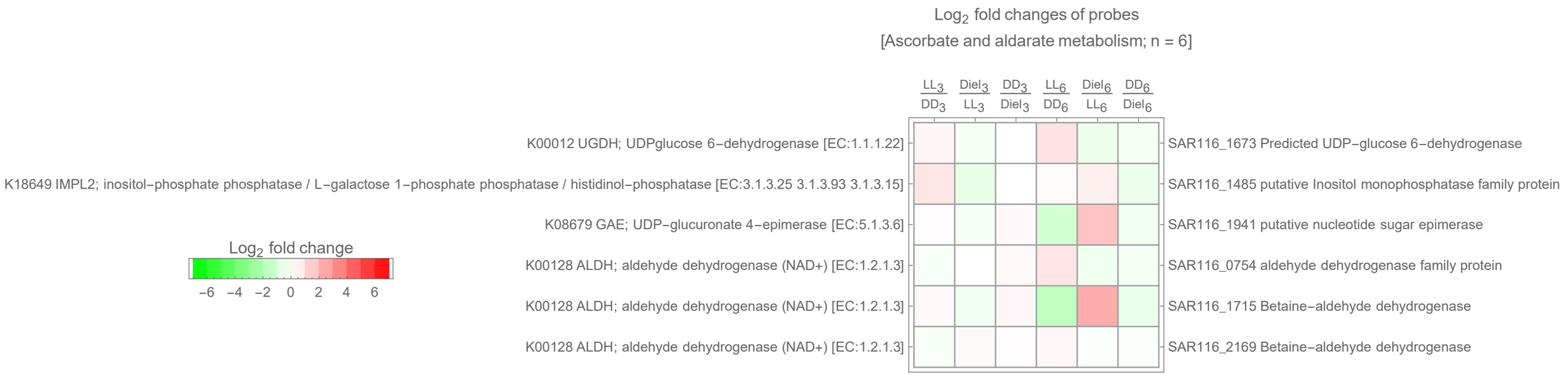
Log2 fold changes of probes
[Ascorbate and aldarate metabolism; n = 6]
LL3 / DD3
Diel3 / LL3
DD3 / Diel3
LL6 / DD6
Diel6 / LL6
DD6 / Diel6
K00012 UGDH; UDPglucose 6−dehydrogenase [EC:1.1.1.22]
K18649 IMPL2; inositol−phosphate phosphatase / L−galactose 1−phosphate phosphatase / histidinol−phosphatase [EC:3.1.3.25 3.1.3.93 3.1.3.15]
K08679 GAE; UDP−glucuronate 4−epimerase [EC:5.1.3.6]
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
SAR116_1673 Predicted UDP−glucose 6−dehydrogenase
SAR116_1485 putative Inositol monophosphatase family protein
SAR116_1941 putative nucleotide sugar epimerase
SAR116_0754 aldehyde dehydrogenase family protein
SAR116_1715 Betaine−aldehyde dehydrogenase
SAR116_2169 Betaine−aldehyde dehydrogenase
Log2 fold change
−6 −4 −2 0 2 4 6