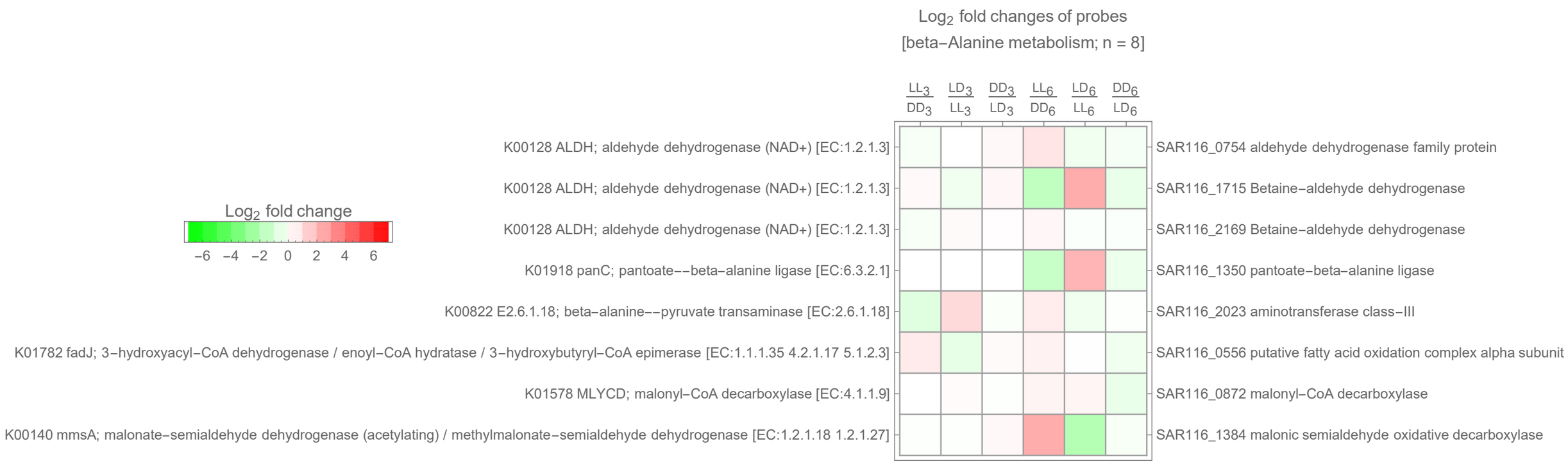
Log2 fold changes of probes
[beta−Alanine metabolism; n = 8]
LL3/DD3
LD3/LL3
DD3/LD3
LL6/DD6
LD6/LL6
DD6/LD6
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]
K01918 panC; pantoate−−beta−alanine ligase [EC:6.3.2.1]
K00822 E2.6.1.18; beta−alanine−−pyruvate transaminase [EC:2.6.1.18]
K01782 fadJ; 3−hydroxyacyl−CoA dehydrogenase / enoyl−CoA hydratase / 3−hydroxybutyryl−CoA epimerase [EC:1.1.1.35 4.2.1.17 5.1.2.3]
K01578 MLYCD; malonyl−CoA decarboxylase [EC:4.1.1.9]
K00140 mmsA; malonate−semialdehyde dehydrogenase (acetylating) / methylmalonate−semialdehyde dehydrogenase [EC:1.2.1.18 1.2.1.27]
SAR116_0754 aldehyde dehydrogenase family protein
SAR116_1715 Betaine−aldehyde dehydrogenase
SAR116_2169 Betaine−aldehyde dehydrogenase
SAR116_1350 pantoate−beta−alanine ligase
SAR116_2023 aminotransferase class−III
SAR116_0556 putative fatty acid oxidation complex alpha subunit
SAR116_0872 malonyl−CoA decarboxylase
SAR116_1384 malonic semialdehyde oxidative decarboxylase
Log2 fold change
−6 −4 −2 0 2 4 6