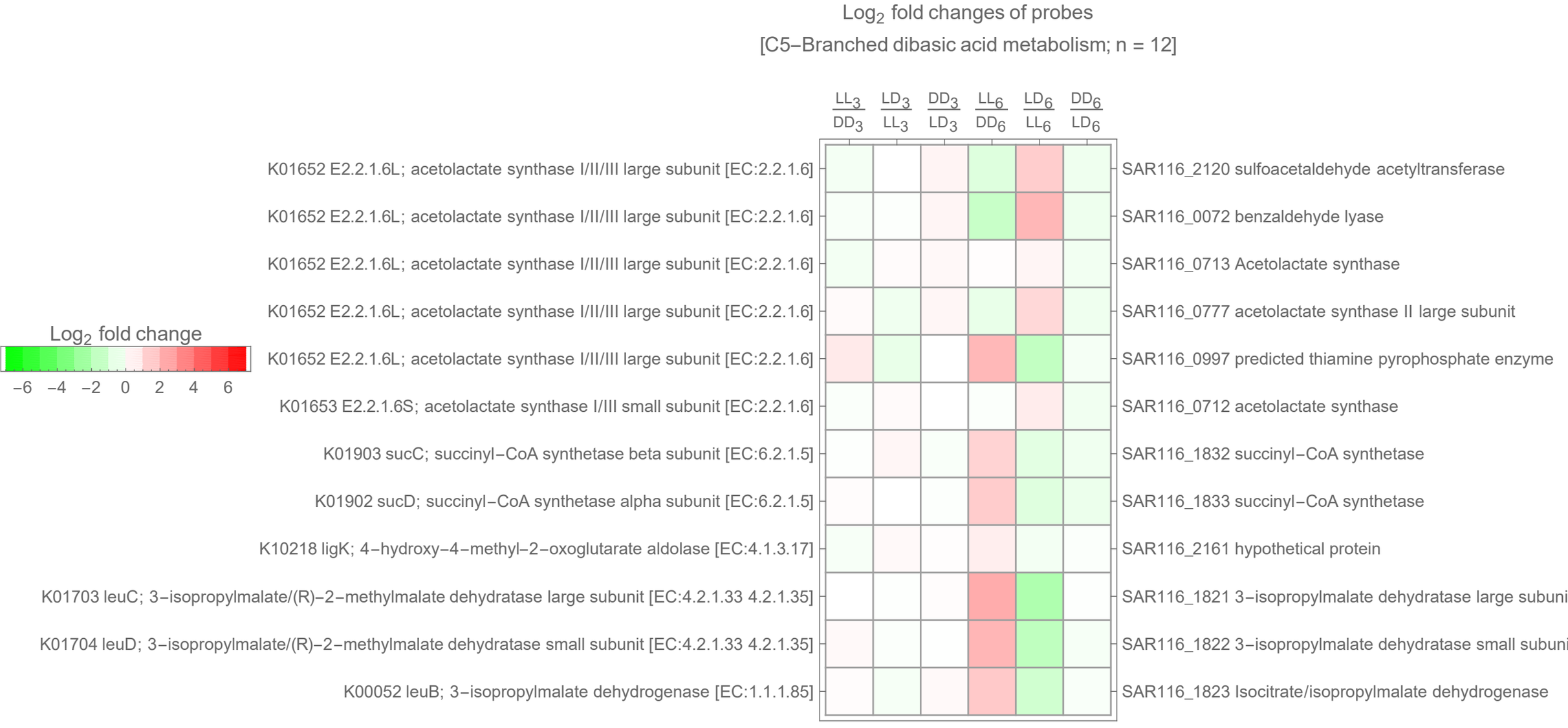
Log2 fold changes of probes
[C5−Branched dibasic acid metabolism; n = 12]
LL3/DD3
LD3/LL3
DD3/LD3
LL6/DD6
LD6/LL6
DD6/LD6
K01652 E2.2.1.6L; acetolactate synthase I/II/III large subunit [EC:2.2.1.6]
K01652 E2.2.1.6L; acetolactate synthase I/II/III large subunit [EC:2.2.1.6]
K01652 E2.2.1.6L; acetolactate synthase I/II/III large subunit [EC:2.2.1.6]
K01652 E2.2.1.6L; acetolactate synthase I/II/III large subunit [EC:2.2.1.6]
K01652 E2.2.1.6L; acetolactate synthase I/II/III large subunit [EC:2.2.1.6]
K01653 E2.2.1.6S; acetolactate synthase I/III small subunit [EC:2.2.1.6]
K01903 sucC; succinyl−CoA synthetase beta subunit [EC:6.2.1.5]
K01902 sucD; succinyl−CoA synthetase alpha subunit [EC:6.2.1.5]
K10218 ligK; 4−hydroxy−4−methyl−2−oxoglutarate aldolase [EC:4.1.3.17]
K01703 leuC; 3−isopropylmalate/(R)−2−methylmalate dehydratase large subunit [EC:4.2.1.33 4.2.1.35]
K01704 leuD; 3−isopropylmalate/(R)−2−methylmalate dehydratase small subunit [EC:4.2.1.33 4.2.1.35]
K00052 leuB; 3−isopropylmalate dehydrogenase [EC:1.1.1.85]
SAR116_2120 sulfoacetaldehyde acetyltransferase
SAR116_0072 benzaldehyde lyase
SAR116_0713 Acetolactate synthase
SAR116_0777 acetolactate synthase II large subunit
SAR116_0997 predicted thiamine pyrophosphate enzyme
SAR116_0712 acetolactate synthase
SAR116_1832 succinyl−CoA synthetase
SAR116_1833 succinyl−CoA synthetase
SAR116_2161 hypothetical protein
SAR116_1821 3−isopropylmalate dehydratase large subunit region
SAR116_1822 3−isopropylmalate dehydratase small subunit
SAR116_1823 Isocitrate/isopropylmalate dehydrogenase
Log2 fold change
−6 −4 −2 0 2 4 6