Log$_2$ fold changes of probes
[Arginine and proline metabolism; n = 25]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K08688 E3.5.3.3; creatinase [EC:3.5.3.3]							SAR116_0762 creatinase
K01480 speB; agmatinase [EC:3.5.3.11]							SAR116_0766 agmatinase
K01480 speB; agmatinase [EC:3.5.3.11]							SAR116_2035 agmatinase
K12256 spuC; putrescine–––pyruvate transaminase [EC:2.6.1.113]							SAR116_2037 probable aminotransferase
K12256 spuC; putrescine–––pyruvate transaminase [EC:2.6.1.113]							SAR116_1762 hypothetical protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_0754 aldehyde dehydrogenase family protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_1715 Betaine–aldehyde dehydrogenase
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_2169 Betaine–aldehyde dehydrogenase
K12255 gbuA; guanidinobutyrase [EC:3.5.3.7]							SAR116_2014 agmatinase
K01426 E3.5.1.4; amidase [EC:3.5.1.4]							SAR116_1728 putative Acetamidase/Formamidase
K09471 puuB; gamma–glutamylputrescine oxidase [EC:1.4.3.–]							SAR116_2036 FAD dependent oxidoreductase
K09472 puuC; 4–(gamma–glutamylamino)butanal dehydrogenase [EC:1.2.1.99]							SAR116_0312 hypothetical protein
K09472 puuC; 4–(gamma–glutamylamino)butanal dehydrogenase [EC:1.2.1.99]							SAR116_0055 aldehyde dehydrogenase family protein
K09472 puuC; 4–(gamma–glutamylamino)butanal dehydrogenase [EC:1.2.1.99]							SAR116_2168 aldehyde dehydrogenase protein
K00286 proC; pyrroline–5–carboxylate reductase [EC:1.5.1.2]							SAR116_2200 Delta 1–pyrroline–5–carboxylate reductase
K13821 putA; RHH–type transcriptional regulator, proline utilization regulon repressor / proline dehydrogenase / delta 1–pyrroline–5–carboxylate dehydrogenase [EC:1.5.5.2 1.2.1.88]							SAR116_0963 proline dehydrogenase
K00931 proB; glutamate 5–kinase [EC:2.7.2.11]							SAR116_1853 Glutamate 5–kinase
K00147 proA; glutamate–5–semialdehyde dehydrogenase [EC:1.2.1.41]							SAR116_1852 gamma–glutamyl phosphate reductase
K01259 pip; proline iminopeptidase [EC:3.4.11.5]							SAR116_1069 proline iminopeptidase
K00812 aspB; aspartate aminotransferase [EC:2.6.1.1]							SAR116_2214 aminotransferase
K00812 aspB; aspartate aminotransferase [EC:2.6.1.1]							SAR116_0974 aspartate aminotransferase
K00813 aspC; aspartate aminotransferase [EC:2.6.1.1]							SAR116_0778 aspartate aminotransferase
K00824 dat; D–alanine transaminase [EC:2.6.1.21]							SAR116_0792 aminotransferase
K22548 lhpL; trans–L–3–hydroxyproline dehydratase [EC:4.2.1.77]							SAR116_0313 proline racemase
K21624 c3lhypd; cis–L–3–hydroxyproline dehydratase [EC:4.2.1.171]							SAR116_0305 muconate lactonizing enzyme

Log$_2$ fold change

−6 −4 −2 0 2 4 6