Log$_2$ fold changes of probes

[Glycerolipid metabolism; n = 9]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K11529 gck; glycerate 2−kinase [EC:2.7.1.165]							SAR116_2472 Hydroxypyruvate reductase
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
K00864 glpK; glycerol kinase [EC:2.7.1.30]							SAR116_0884 glycerol kinase
K03621 plsX; phosphate acyltransferase [EC:2.3.1.274]							SAR116_0171 Fatty acid synthesis plsX protein
K08591 plsY; acyl phosphate:glycerol−3−phosphate acyltransferase [EC:2.3.1.275]							SAR116_0615 hypothetical protein
K00655 plsC; 1−acyl−sn−glycerol−3−phosphate acyltransferase [EC:2.3.1.51]							SAR116_2351 phospholipid/glycerol acyltransferase
K07406 melA; alpha−galactosidase [EC:3.2.1.22]							SAR116_1196 glycoside hydrolase family 4

Log$_2$ fold change

−6 −4 −2 0 2 4 6