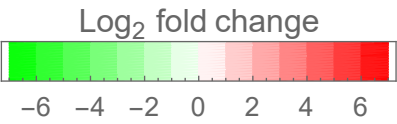


K18649 IMPL2; inositol-phosphate phosphatase / L-galactose 1-phosphate phosphatase / histidinol-phosphatase [EC:3.1.3.25 3.1.3.93 3.1.3.15]



Log₂ fold changes of probes
[Ascorbate and aldarate metabolism; n = 6]

	$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K00012 UGDH; UDPglucose 6-dehydrogenase [EC:1.1.1.22]							SAR116_1673 Predicted UDP-glucose 6-dehydrogenase
K08679 GAE; UDP-glucuronate 4-epimerase [EC:5.1.3.6]							SAR116_1485 putative Inositol monophosphatase family protein
K00128 ALDH; aldehyde dehydrogenase (NAD+) [EC:1.2.1.3]							SAR116_1941 putative nucleotide sugar epimerase
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