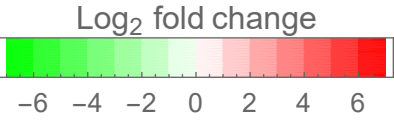


Log₂ fold changes of probes
[Fructose and mannose metabolism; n = 16]



		$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$		
	K00847 E2.7.1.4; fructokinase [EC:2.7.1.4]							SAR116_0620 fructokinase, ROK family protein	
	K00847 E2.7.1.4; fructokinase [EC:2.7.1.4]							SAR116_0973 fructokinase	
K16011	algA; mannose-1-phosphate guanylyltransferase / mannose-6-phosphate isomerase [EC:2.7.7.13 5.3.1.8]							SAR116_1966 Mannose-1-phosphate guanylyltransferase/mannose-6-phosphate isomerase	
	K01840 manB; phosphomannomutase [EC:5.4.2.8]							SAR116_1670 Phosphomannomutase	
	K01711 gmd; GDPmannose 4,6-dehydratase [EC:4.2.1.47]							SAR116_1961 GDPmannose 4,6-dehydratase	
	K02377 TSTA3; GDP-L-fucose synthase [EC:1.1.1.271]							SAR116_1960 NAD-dependent epimerase/dehydratase	
	K02377 TSTA3; GDP-L-fucose synthase [EC:1.1.1.271]							SAR116_0989 NAD-dependent epimerase/dehydratase	
	K18335 K18335; 2-keto-3-deoxy-L-fuconate dehydrogenase [EC:1.1.1.-]							SAR116_0303 short-chain dehydrogenase/reductase SDR	
K21883	lra5; 2-dehydro-3-deoxy-L-rhamnonate dehydrogenase (NAD+) [EC:1.1.1.401]							SAR116_0926 short-chain dehydrogenase/reductase SDR	
	K03841 FBP; fructose-1,6-bisphosphatase I [EC:3.1.3.11]							SAR116_1413 fructose-1,6-bisphosphatase I	
	K01805 xylA; xylose isomerase [EC:5.3.1.5]							SAR116_0264 putative xylose isomerase	
	K00045 E1.1.1.67; mannitol 2-dehydrogenase [EC:1.1.1.67]							SAR116_0965 Mannitol dehydrogenase-like protein	
	K01624 FBA; fructose-bisphosphate aldolase, class II [EC:4.1.2.13]							SAR116_1412 Fructose-bisphosphate aldolase	
	K01624 FBA; fructose-bisphosphate aldolase, class II [EC:4.1.2.13]							SAR116_0281 fructose-bisphosphate aldolase	
	K01803 TPI; triosephosphate isomerase (TIM) [EC:5.3.1.1]							SAR116_0198 Triosephosphate isomerase	
	K01808 rpiB; ribose 5-phosphate isomerase B [EC:5.3.1.6]							SAR116_0160 sugar-phosphate isomerase	