Log$_2$ fold changes of probes
[Glutathione metabolism; n = 20]

	$\frac{LL_3}{DD_3}$	$\frac{Diel_3}{LL_3}$	$\frac{DD_3}{Diel_3}$	$\frac{LL_6}{DD_6}$	$\frac{Diel_6}{LL_6}$	$\frac{DD_6}{Diel_6}$	
K00681 ggt; gamma–glutamyltranspeptidase / glutathione hydrolase [EC:2.3.2.2 3.4.19.13]							SAR116_1348 gamma–glutamyltranspeptidase
K07232 CHAC; glutathione–specific gamma–glutamylcyclotransferase [EC:4.3.2.7]							SAR116_1147 ChaC family protein
K01469 OPLAH; 5–oxoprolinase (ATP–hydrolysing) [EC:3.5.2.9]							SAR116_0487 5–oxoprolinase (ATP–hydrolyzing)
K07160 pxpA; 5–oxoprolinase (ATP–hydrolysing) subunit A [EC:3.5.2.9]							SAR116_1772 hypothetical protein
K01919 gshA; glutamate––cysteine ligase [EC:6.3.2.2]							SAR116_1123 glutamate––cysteine ligase
K01920 gshB; glutathione synthase [EC:6.3.2.3]							SAR116_1628 glutathione synthetase
K01255 CARP; leucyl aminopeptidase [EC:3.4.11.1]							SAR116_0630 Leucyl aminopeptidase
K01255 CARP; leucyl aminopeptidase [EC:3.4.11.1]							SAR116_1483 Leucyl aminopeptidase
K01256 pepN; aminopeptidase N [EC:3.4.11.2]							SAR116_0100 aminopeptidase N
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_2368 Glutathione S–transferase
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_1238 Glutathione S–transferase
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_0032 glutathione S–transferase
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_2377 glutathione S–transferase family protein
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_0230 Glutathione S–transferase–like protein
K00799 GST; glutathione S–transferase [EC:2.5.1.18]							SAR116_2447 Glutathione S–transferase domain protein
K00383 GSR; glutathione reductase (NADPH) [EC:1.8.1.7]							SAR116_0683 glutathione–disulfide reductase
K00031 IDH1; isocitrate dehydrogenase [EC:1.1.1.42]							SAR116_0090 Isocitrate dehydrogenase
K00036 G6PD; glucose–6–phosphate 1–dehydrogenase [EC:1.1.1.49 1.1.1.363]							SAR116_1986 glucose–6–phosphate 1–dehydrogenase
K00432 gpx; glutathione peroxidase [EC:1.11.1.9]							SAR116_2363 Phospholipid hydroperoxide glutathione peroxidase
K24138 prx3; glutaredoxin/glutathione–dependent peroxiredoxin [EC:1.11.1.25 1.11.1.27]							SAR116_1109 Redoxin domain protein

Log$_2$ fold change

−6 −4 −2 0 2 4 6