Log$_2$ fold changes of probes
[TADB2.0; n = 27]

$\frac{LL_3}{DD_3}$	$\frac{Diel_3}{LL_3}$	$\frac{DD_3}{Diel_3}$	$\frac{LL_6}{DD_6}$	$\frac{Diel_6}{LL_6}$	$\frac{DD_6}{Diel_6}$

Log$_2$ fold change

−6 −4 −2 0 2 4 6

SAR116_0006 hypothetical protein SAR116_0006

SAR116_0013 transcriptional regulator, BadM/Rrf2 family

SAR116_0101 Superfamily II DNA and RNA helicase

SAR116_0455 GCN5−related N−acetyltransferase

K00764 purF; amidophosphoribosyltransferase [EC:2.4.2.14] — SAR116_0651 Glutamine phosphoribosylpyrophosphate amidotransferase

SAR116_0697 Spermidine N1−acetyltransferase

K03824 yhbS; putative acetyltransferase [EC:2.3.1.−] — SAR116_0701 hypothetical protein SAR116_0701

SAR116_0880 GCN5−related N−acetyltransferase

K00640 cysE; serine O−acetyltransferase [EC:2.3.1.30] — SAR116_1024 serine O−acetyltransferase

SAR116_1185 probable transcriptional regulator

K07657 phoB; two−component system, OmpR family, phosphate regulon response regulator PhoB — SAR116_1201 two component transcriptional regulator, winged helix family

K07316 mod; adenine−specific DNA−methyltransferase [EC:2.1.1.72] — SAR116_1283 DNA methylase N−4/N−6 domain protein

SAR116_1325 CBS domain containing protein

K01520 dut; dUTP pyrophosphatase [EC:3.6.1.23] — SAR116_1566 dUTPase

SAR116_1630 GCN5−related N−acetyltransferase

SAR116_1631 Transcriptional Regulator, XRE family

SAR116_1706 DNA−binding protein

SAR116_1748 phage integrase family protein

K09768 K09768; uncharacterized protein — SAR116_1997 YaiI/YqxD family protein, putative

K07552 bcr; MFS transporter, DHA1 family, multidrug resistance protein — SAR116_2000 drug resistance transporter, Bcr/CflA subfamily

SAR116_2155 Aha1 domain protein

SAR116_2220 Transcriptional Regulator, AsnC family

SAR116_2221 DNA binding domain, excisionase family

K10778 ada; AraC family transcriptional regulator, regulatory protein of adaptative response / methylated−DNA−[protein]−cysteine methyltransferase [EC:2.1.1.63] — SAR116_2236 transcriptional regulator, AraC family protein

K07171 mazF; mRNA interferase MazF [EC:3.1.−.−] — SAR116_2309 hypothetical protein SAR116_2309

K03657 uvrD; ATP−dependent DNA helicase UvrD/PcrA [EC:5.6.2.4] — SAR116_2321 DNA helicase II

K00799 GST; glutathione S−transferase [EC:2.5.1.18] — SAR116_2377 glutathione S−transferase family protein