

$Log_2 \ \, \text{fold changes of probes}$ [Protein families: genetic information processing; n = 10]

K03978 engB; GTP-binding protein SAR116_1501 ribosome biogenesis GTP-binding protein YihA/YsxC K03665 hflX; GTPase SAR116_0793 GTPase HflX K03979 obgE; GTPase [EC:3.6.5.-] SAR116_1854 GTPase ObgE K03595 era; GTPase SAR116_0530 GTPase Era K03977 engA; GTPase SAR116_2501 ribosome biogenesis GTPase Der K06942 ychF; ribosome-binding ATPase SAR116_2191 redox-regulated ATPase YchF [Candidatus Puniceispirillum marinum] K06925 tsaE; tRNA threonylcarbamoyladenosine biosynthesis protein TsaE SAR116_1263 tRNA (adenosine(37)–N6)-threonylcarbamoyltransferase complex ATPase subunit type 1 TsaE K03650 mnmE; tRNA modification GTPase [EC:3.6.-.-] SAR116_1645 tRNA uridine-5-carboxymethylaminomethyl(34) synthesis GTPase MnmE K06020 ettA; energy-dependent translational throttle protein EttA SAR116_2318 energy-dependent translational throttle protein EttA K06158 ABCF3; ATP-binding cassette, subfamily F, member 3 SAR116_2053 ABC-F family ATP-binding cassette domain-containing protein