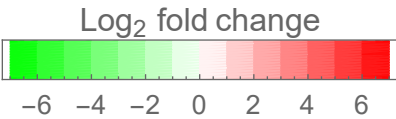


Log₂ fold changes of probes
[Nitrogen metabolism; n = 10]



	<u>LL₃</u> DD ₃	<u>LD₃</u> LL ₃	<u>DD₃</u> LD ₃	<u>LL₆</u> DD ₆	<u>LD₆</u> LL ₆	<u>DD₆</u> LD ₆
K01455 E3.5.1.49; formamidase [EC:3.5.1.49]						
K00459 ncd2; nitronate monooxygenase [EC:1.13.12.16]						
K00261 GLUD1_2; glutamate dehydrogenase (NAD(P)+) [EC:1.4.1.3]						
K01915 glnA; glutamine synthetase [EC:6.3.1.2]						
K01915 glnA; glutamine synthetase [EC:6.3.1.2]						
K01915 glnA; glutamine synthetase [EC:6.3.1.2]						
K01915 glnA; glutamine synthetase [EC:6.3.1.2]						
K00265 gltB; glutamate synthase (NADPH) large chain [EC:1.4.1.13]						
K00266 gltD; glutamate synthase (NADPH) small chain [EC:1.4.1.13]						
K01673 cynT; carbonic anhydrase [EC:4.2.1.1]						
SAR116_1711 Acetamidase/Formamidase						
SAR116_2516 hypothetical protein						
SAR116_0453 Glu/Leu/Phe/Val dehydrogenase						
SAR116_0106 Glutamine synthetase type I						
SAR116_0113 glutamine synthetase						
SAR116_2038 Glutamate--ammonia ligase						
SAR116_2030 glutamine synthetase						
SAR116_1240 Glutamate synthase (ferredoxin)						
SAR116_1241 glutamate synthase [NADPH] small chain						
SAR116_2462 Carbonate dehydratase						