Additional file 7

Putative methylases/methyltransferases in the genome of Microcystis aeruginosa PCC 7806

Gene ID	COG number		Gene ID	COG number	Function
mic0260	COG0270	Site-specific DNA methylase (putative <i>Eco47II</i> , <i>Sau96I</i> modification methylase)	mic3149	COG0863	DNA modification methylase (putative <i>CfrBI</i> modification methylase)
mic0262	COG0863	DNA modification methylase	mic3329	COG1092	Predicted SAM-dependent methyltransferase
mic0319		SAM-dependent methyltransferase	mic3380	COG1743	Adenine-specific DNA methylase containing a Zn-ribbon
mic0367		SAM-dependent methyltransferase	mic3417	COG2189	Adenine specific DNA methylase (Putative type II DNA modification enzyme)
mic0434	COG4889	Adenine specific DNA methyltransferase	mic3467	COG1002	Type II restriction enzyme, methylase subunit
mic0510		SAM-dependent methyltransferase	mic3571	COG0116	Predicted N6-adenine-specific DNA methylase
mic0553		SAM-dependent methyltransferase	mic3707	COG0286	Type I restriction-modification system methyltransferase subunit
mic0555		SAM-dependent methyltransferase	mic3729	COG0220	Predicted SAM-dependent methyltransferase
mic0714	COG0286	Type I restriction-modification system methyltransferase subunit	mic3737		SAM-dependent methyltransferase
mic0757		SAM-dependent methyltransferase	mic3776		Methyltransferase (putative <i>SsoI</i> methylase)
mic0919		Modification methylase (putative <i>EcoRV</i> methylase)	mic3919	COG0270	Site-specific DNA methylase (putative type II <i>HaeII</i> modification methylase)
mic1132		SAM-dependent methyltransferase	mic4082	COG0863	DNA modification methylase (putative <i>BstYI</i> methyltransferase)
mic1135		Predicted O-methyltransferase	mic4091	COG1002	Type II restriction enzyme, methylase subunit
mic1145	COG0286	Type I restriction-modification system methyltransferase subunit	mic4194		SAM-dependent methyltransferase
mic1231	COG0270	Site-specific DNA methylase	mic4261	COG0338	Site-specific DNA methylase (putative <i>EcoRV</i> modification methylase)
mic1247		SAM-dependent methyltransferase	mic4360		SAM-dependent methyltransferase
mic1485	COG0270	Site-specific DNA methylase	mic4390	COG0270	Site-specific DNA methylase
mic1536	COG0270	Site-specific DNA methylase	mic4632	COG0270	Site-specific DNA methylase
mic1611	COG0338	Site-specific DNA methylase (MspA11 methyltransferase)	mic4878		SAM-dependent methyltransferase
mic1716	COG0286	Type I restriction-modification system methyltransferase subunit	mic4888	COG1002	Type II restriction enzyme, methylase subunit
mic1905		predicted N-6 adenine-specific DNA methylase	mic4913	COG0286	Type I restriction-modification system methyltransferase subunit
mic1963	COG0313	Predicted methyltransferase	mic4976		SAM-dependent methyltransferase
mic2077		SAM-dependent methyltransferase	mic5066	COG0270	Site-specific DNA methylase (putative <i>HhaI</i> modification methylase)
mic2080		SAM-dependent methyltransferase	mic5070	COG1002	Type II restriction enzyme, methylase subunit
mic2088	COG1002	Type II restriction enzyme, methylase subunit	mic5255		SAM-dependent methyltransferase
mic2201	COG0270	Site-specific DNA methylase	mic5298		Modification methylase (putative NspV methylase)
mic2315		Site-specific DNA methylase	mic5314		SAM-dependent methyltransferase
mic2318	COG0863	DNA modification methylase (putative PvuII modification methylase)	mic5330		Type II restriction enzyme, methylase subunit
mic2460		SAM-dependent methyltransferase	mic5528		N6-adenine-specific methylase
mic2464	COG0863	DNA modification methylase	mic5612		SAM-dependent methyltransferase
mic2569		SAM-dependent methyltransferase	mic5613	COG0338	Site-specific DNA methylase
mic2593	COG1002	Type II restriction enzyme, methylase subunit	mic5683	COG0338	Site-specific DNA methylase
mic2808	COG0863	DNA modification methylase (putative AvaI modification methylase)	mic6590	COG1002	Type II restriction enzyme, methylase subunit
mic3127		SAM-dependent methyltransferase	mic6600	COG0270	Site-specific DNA methylase (putative <i>HhaI</i> modification methylase)
mic3146	COG0863	DNA modification methylase (cytosine methylase)	mic7256	COG0286	Type I restriction-modification system methyltransferase subunit

Enzymes shown in bold are co-localized with putative methylases shown in the table of additional file 6.