NC_019396 Gluconobacter oxydans H24
NC_010125 Gluconacetobacter diazotrophicus PA1 5
NC_013355 Zymomonas mobilis subsp. mobilis NCIMB 11163
NC_018485 Methylocystis sp. SC2
NC_016594 Azospirillum brasilense Sp245
NC_011894 Methylobacterium nodulans ORS 2060
NC_017249 Bradyrhizobium japonicum USDA 6
NC_015717 Hyphomicrobium sp. MC1
NC_011386 Oligotropha carboxidovorans OM5
NC_005296 Rhodopseudomonas palustris CGA009
NC_020453 Bradyrhizobium oligotrophicum S58
NC_017249 Bradyrhizobium japonicum USDA 6
NC_007964 Nitrobacter hamburgensis X14
NC_011666 Methylocella silvestris BL2
NC_010581 Beijerinckia indica subsp. indica ATCC 9039
NC_014664 Rhodomicrobium vannielii ATCC 17100
NC_015723 Cupriavidus necator N-1
NC_007952 Burkholderia xenovorans LB400
NC_022904 Pandoraea pnomenusa 3kgm
NC_012997 Teredinibacter turnerae T7901
NC_016112 Methylomicrobium alcaliphilum 20Z
NC_015572 Methylomonas methanica MC09
NC_017033 Frateuria aurantia DSM 6220
NC_002977 Methylococcus capsulatus str. Bath
NC_013960 Nitrosococcus halophilus Nc 4
NC_015731 Nitrosomonas sp. Is79A3
NC_007614 Nitrosospira multiformis ATCC 25196
NC_012560 Azotobacter vinelandii DJ
NC_011761 Acidithiobacillus ferrooxidans ATCC 23270
COG0775 Nucleoside phosphorylase
COG1562 Phytoene/squalene synthetase
COG1657 Squalene cyclase
COG0535 Radical SAM superfamily enzyme, MoaA/NifB/PqqE/SkfB family