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| --- | --- | --- | --- |
| **Process** | **Gene** | **KO** | **notes** |
| Aerobic C fixation | ribulose-bisphosphate carboxylase large (*rbc*L) | K01601 | Calvin cycle |
| ribulose-bisphosphate carboxylase small (*rbc*S) | K01602 | Calvin cycle |
| phosphoribulose kinase (*prk*A) | K00855 | Calvin cycle |
| Respiration | cytochrome C oxidase subunit I (*cox*I) | K02256 | Eucaryotic |
| cytochrome C oxidase subunit III (*cox*III) | K02262 | Eucaryotic |
| cytochrome C oxidase subunit I (*cox*A) | K02274 | Bacterial |
| cytochrome C oxidase subunit III (*cox*C) | K02276 | Bacterial |
| Anaerobic C fixation | ATP-citrate lyase alpha subunit (*acl*A) | K15230 | Bacterial, rTCA |
| ATP-citrate lyase beta subunit (*acl*B) | K15231 | Bacterial, rTCA |
| citryl-CoA lyase (*ccl*) | K15234 | rTCA |
| citryl-CoA synthetase (*ccs*B) | K15233 | rTCA |
| fumerate reductase flavoprotein subunit (*frd*A) | K00244 | rTCA |
| carbon monoxide dehydrogenase/acetyl-CoA synthase alpha (CODH-ACSA) | K14138 | WL |
| carbon monoxide dehydrogenase/acetyl-CoA synthase beta (CODH-ACSA) | K00190 | WL |
| Fermentation | L-lactate dehydrogenase | K00016 |  |
| pyruvate:ferredoxin oxidoreductase alpha subunit | K00169 |  |
| pyruvate:ferredoxin oxidoreductase beta subunit | K00170 |  |
| CO oxidation | carbon-monoxide dehydrogenase large (*cox*L) | K03520 |  |
| carbon-monoxide dehydrogenase medium (*cox*M) | K03519 |  |
| carbon-monoxide dehydrogenase small (*cox*S) | K03518 |  |
| Methanogenesis | coenzyme M methyl reductase (*mcr*B)  methyl coenzyme M reductase system | K00401  K00400 |  |
| Methane oxidation | soluble methane monooxygenase | K08684 | KO deprecated |
| N fixation | nitrogenase | K00531 |  |
| nitrogenase molybdenum-iron protein alpha (*nif*D) | K02586 |  |
| nitrogenase iron protein (*nif*H) | K02588 |  |
| nitrogenase molybdenum-iron protein beta (*nif*K) | K02591 |  |
| Denitrification | nitric oxide reductase (*nor*B) | K02305 |  |
| nitric oxide reductase (*nor*C) | K04561 |  |
| nitrous oxide reductase (*nos*Z) | K00376 |  |
| DNRA | NrfA periplasmic cytochrome c552 subunit | K03385 |  |
| Anammox | hydroxylamine oxidase (hao) | K10535 |  |
| N mineralization | glutamate dehydrogenase (*gdh*A) | K00260 |  |
| glutamate dehydrogenase (*gdh*A) | K00261 |  |
| glutamate dehydrogenase (*gdh*A) | K00262 |  |
| N assimilation | assimilatory nitrate reductase | K00360 |  |
| assimilatory nitrate reductase | K00361 |  |
| glutamine synthetase (*gln*A) | K01915 |  |
| glutamate synthetase (NADPH/NADH)(*glt*B) | K00265 |  |
| glutamate synthetase (ferredoxin) (*glt*S) | K00284 |  |
| Nitrification | ammonia monooxygenase subunit A (*amo*A) | K10944 |  |
| ammonia monooxygenase subunit B (*amo*B) | K10945 |  |
| ammonia monooxygenase subunit C (*amo*C) | K10946 |  |
| ASR | Adenyl sulfate kinase (*cys*C)  Sulfate adenylyltransferase (*cys*N, *cys*D) | K00860  K00956  K00957 |  |
| DSR | APS reductase (*apr*A) | K00394 |  |
| S mineralization | Cysteine diogenase  3-Mercaptopyruvate sulfurtransferase | K00456  K01011 |  |

Table 2. Full list of KEGG Orthologs (KO) searched for in Organic Lake profile metagenome.