Glycerol uptake facilitator or related Ribosomal protein Cytoskeletal protein RodZ, Lyzozyme M1 (1,4-beta-N-acetyln DNA-directed RNA poly Anthranilate phosphol 5-methylthioribulose/5-deoxyribulose/Fuculose 1-phosphate ald L-asparaginase/archa Alkyl sulfatase BDS1 and related hydi Uncharacterized conserved protein MJ1445, calcingur Fluoride ion exporter CrcB/FE Uncharacterized conserved protein YqeY may ha NADH:ubiquinone oxide Predicted nuc Pyruvate/o Creatinine amidohydrolase/Fe(II)-dependent FAPy formamide hydrolase/ Midasin, AAA ATPase With WWA do ABC-type enterobactin trans Uncharacterized conserved protein YpbQ, isoprenylcysteine carb Spore coat polysaccharide biosynthesis protein 16S rRNA G527 N7-me Uncharacter UDP-N-acetylmuramat Endo alpha-1,4 polygalactosaminidase, GH114 family (was erroneously anno ABC-type sugar transport sys Thioredoxin/glutathione peroxidase Btu Predicted esterase of the alpha/Phosphopanic CRISPR/Cas system-associated protein Cas10, large subunit of type III CRISPR/Cas system-associated protein Cas10, large subunit of type III CRISPR/Cas mRNA-degrading endonuclease YafQ (mRNA interferase), tox Flavin-dependent oxidoreductase, luciferase family (includes alkanesulfonate monooxygenase SSUD an CRISPR-Cas system type III CSM-effector complete the complete complete the complete c-di-AMP phosphodieste Phenylpropionate dioxygenase or re Bifunctional protein GlmU, N-acetylglucosamine-1-phosphate-ur Fermentation-respiration sw Type IV Uncharacterized phage-associate Spore dipicolinate transporter SpoVV/YjiH/YlbJ, contains nucled Predicted GntP-related membrane per

GEM: Campylobacterota

