Top 30 of GO Enrichment propionyl-CoA carboxylase activity -NAD+ ADP-ribosyltransferase activity linoleoyl-CoA desaturase activity hydrogen:proline symporter activity guanylate cyclase activity glycine betaine:proton symporter activity coenzyme binding -Gene\_number ATP-dependent DNA helicase activity -• 2 acetyl-CoA carboxylase activity protein complex bacterial nucleoid acetyl-CoA carboxylase complex regulation of lipid biosynthetic process q\_value quaternary ammonium group transport -0.20 proline transmembrane transport positive regulation of DNA binding -0.15 polysaccharide biosynthetic process -0.10 peptidyl-threonine phosphorylation organic cation transport -GO domain negative regulation of fatty acid biosynthetic process biological\_process negative regulation by symbiont of host innate immune response cellular\_component mycolic acid biosynthetic process molecular\_function modified amino acid transport long-chain fatty acid biosynthetic process hyperosmotic salinity response cGMP biosynthetic process cellular metabolic process carbon fixation amino-acid betaine transport -3-phenylpropionate catabolic process -7.5 10.0 5.0 12.5 15.0 Rich Factor