Top 30 of GO Enrichment transferase activity, transferring glycosyl groups protein serine/threonine kinase activity protein kinase activity nucleoside-triphosphatase activity -NADP binding methyltransferase activity -Gene_number linoleoyl-CoA desaturase activity -• 2 electron carrier activity -ATPase activity -3-oxoacyl-[acyl-carrier-protein] reductase (NADPH) activity -2-hydroxy-3-oxopropionate reductase activity -ATP-binding cassette (ABC) transporter complex response to stress response to organic substance q_value response to nitrosative stress -0.125 protein secretion by the type VII secretion system -0.100 positive regulation of catalytic activity -0.075 negative regulation of fatty acid biosynthetic process -0.050 negative regulation of catalytic activity -0.025 negative regulation by symbiont of host cell-mediated immune response -GO domain mycolic acid biosynthetic process biological_process mannosylation cellular_component long-chain fatty acid biosynthetic process molecular_function growth involved in symbiotic interaction glycolipid biosynthetic process fatty acid elongation carnitine catabolic process -ATP catabolic process alkanesulfonate transport active evasion of host immune response -7.5 5.0 12.5 2.5 10.0 Rich Factor