GO enrichment xanthophyll biosynthetic process vacuole triterpenoid biosynthetic process sucrose:proton symporter activity sucrose transport --1 \* log10(Qvalue) 1.55 sucrose transmembrane transporter activity scopolin beta-glucosidase activity -1.50 pentacyclic triterpenoid biosynthetic process -1.45 multivesicular body -1.40 GO term lanosterol synthase activity -1.35 glucosinolate catabolic process -Genes cycloartenol synthase activity cellular zinc ion homeostasis carotenoid biosynthetic process carotene beta-ring hydroxylase activity -20 carbohydrate metabolic process brassinosteroid metabolic process beta-glucosidase activity beta-amyrin synthase activity -ADP binding -3.6 3.0 3.9 3.3 -log10(Pvalue)