Cluster 7 CC Top 30 GO enrichment result intracellular anatomical structure intracellular non-membrane-bounded organelle non-membrane-bounded organelle ribosome endomembrane systemcytosolendoplasmic reticulum ribosomal subunitchloroplast stromaplastid stroma Qvalue nuclear outer membrane-endoplasmic reticulum membrane network-0.05 endoplasmic reticulum membrane 0.10 endoplasmic reticulum subcompartment-0.15 vacuolar membrane 0.20 cytosolic ribosome 0.25 chloroplast envelope Gene\_Number large ribosomal subunit-100 extracellular region -200 cytosolic large ribosomal subunit-300 400 cell wallproton-transporting V-type ATPase complexapoplastproton-transporting V-type ATPase, V0 domainlate endosome vacuolar proton-transporting V-type ATPase, V0 domainchloroplast inner membrane plastid inner membrane tricarboxylic acid cycle enzyme complexglyoxysome mitochondrial nucleoid 0.0 0.2 0.4 0.6 Gene\_Ratio