Cluster 6 BP Top 30 GO enrichment result organonitrogen compound metabolic process phosphorus metabolic process small molecule metabolic process oxoacid metabolic process organic acid metabolic process carboxylic acid metabolic process phosphorylation organic substance catabolic process small molecule biosynthetic process macromolecule catabolic process organonitrogen compound catabolic process Gene_Number organic acid biosynthetic process 100 carboxylic acid biosynthetic process 200 300 protein catabolic process alpha-amino acid biosynthetic process Qvalue dicarboxylic acid metabolic process malate metabolic process 0.05 aspartate family amino acid metabolic process 0.10 potassium ion transmembrane transport-0.15 branched-chain amino acid biosynthetic process protein refolding isoleucine metabolic process isoleucine biosynthetic process diaminopimelate metabolic process lysine biosynthetic process via diaminopimelate lysine metabolic process lysine biosynthetic process pollen tube developmentregulation of ubiquitin-dependent protein catabolic process regulation of oxidoreductase activity 0.0 0.2 0.3 0.1 0.4 Gene_Ratio