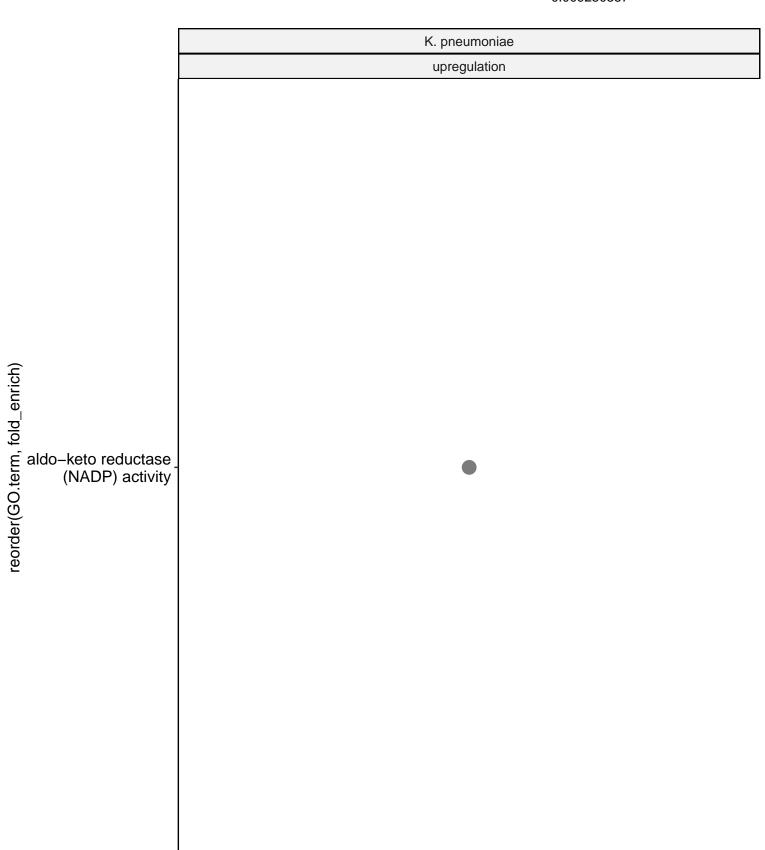


'eorder(GO.term, fold_enrich)



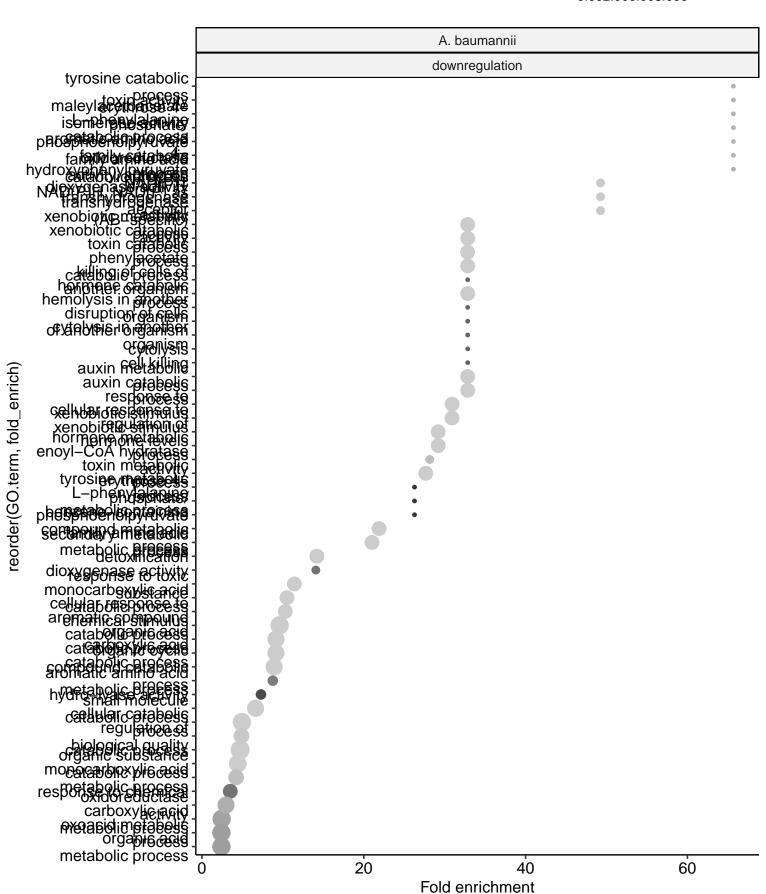
8.725

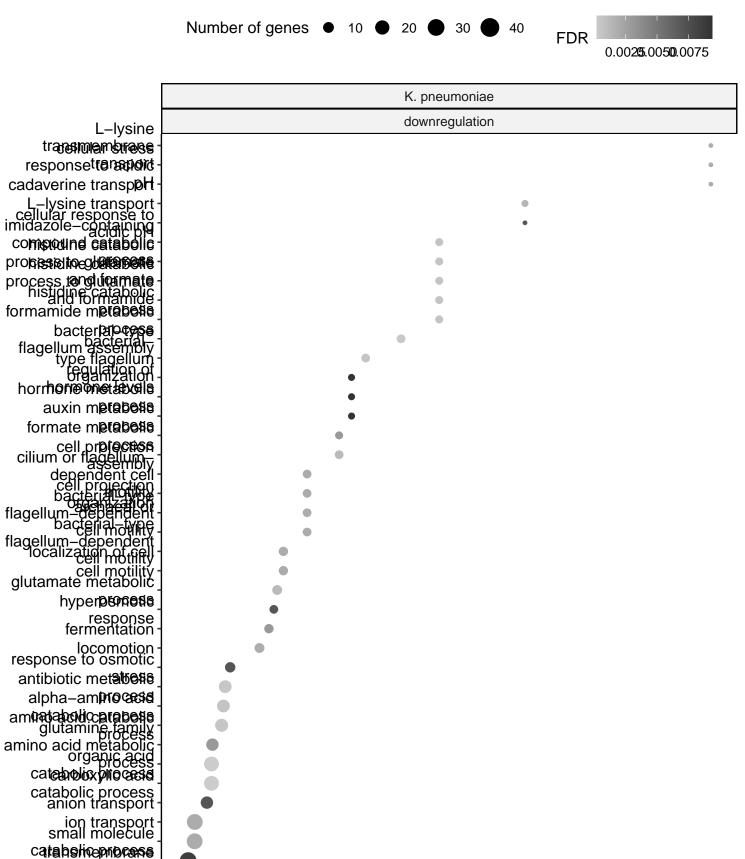
Fold enrichment

8.750

8.675

8.700





reorder(GO.term, fold_enrich)

flagellum

cellular transport

process localization