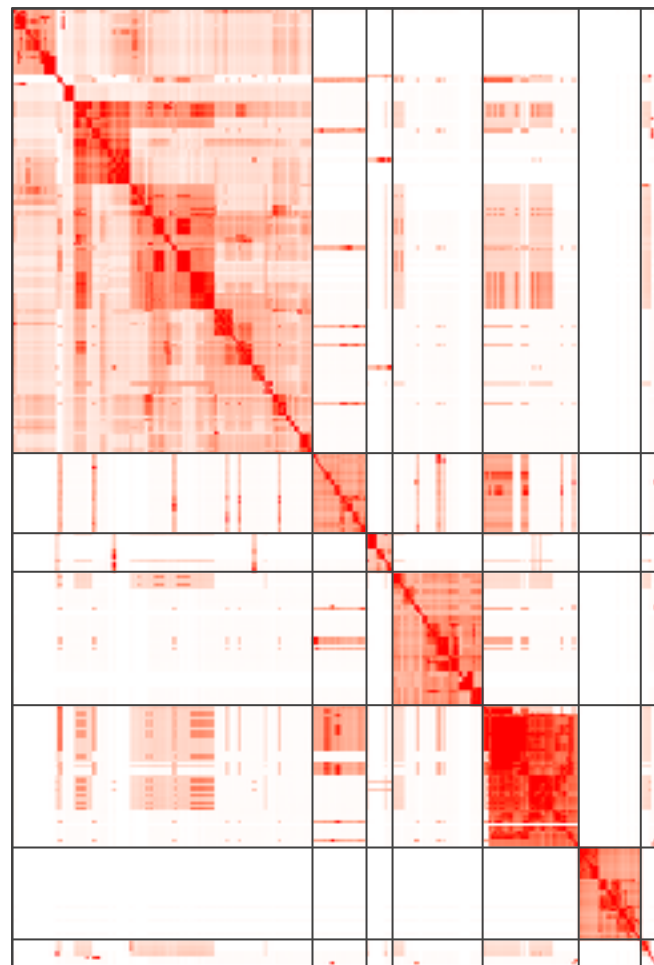
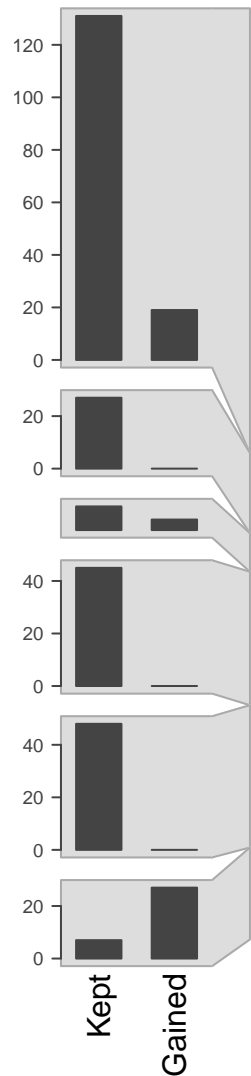


323 GO terms clustered by 'binary_cut'

Number of significant GO terms in each cluster (padj < 0.01)



metabolic expression
macromolecule rna splicing dnatemplated
process modification mrna

ribonucleoprotein fission assembly projection chromatin organelle
organization change biogenesis subunit

response bmp cellular stimulus compound stress growth

er
vesiclemediated membrane cotranslational nucleus insertion
transport localization targeting

chromatid metaphaseanaphase chromosome cell separation
cycle mitotic transition segregation sister

anatomical
specification formation determination embryo neural
development symmetry pattern mesoderm

not available
available

Similarity

