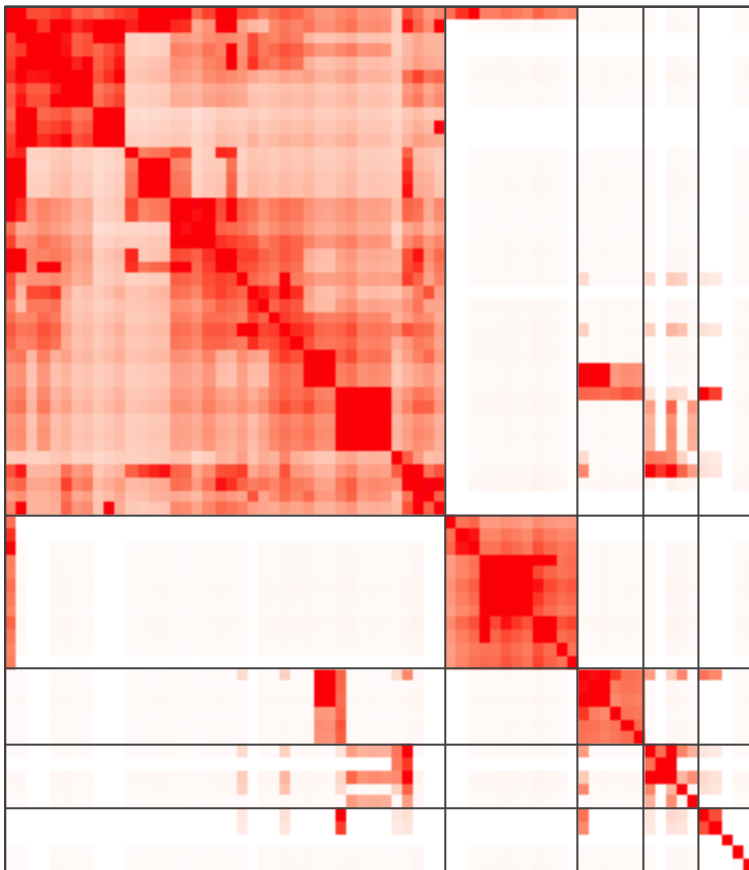




# 68 GO terms clustered by 'binary\_cut'



acid  
compound synthesis coupled atp transcription electron  
catabolic rna polymerase negative cellular mitochondrial  
metabolic repair regulation transport glucose  
**process** biosynthetic

**Similarity**  
0.6  
0.4  
0.2  
0

**transport** ion intracellular

regulation positive signaling