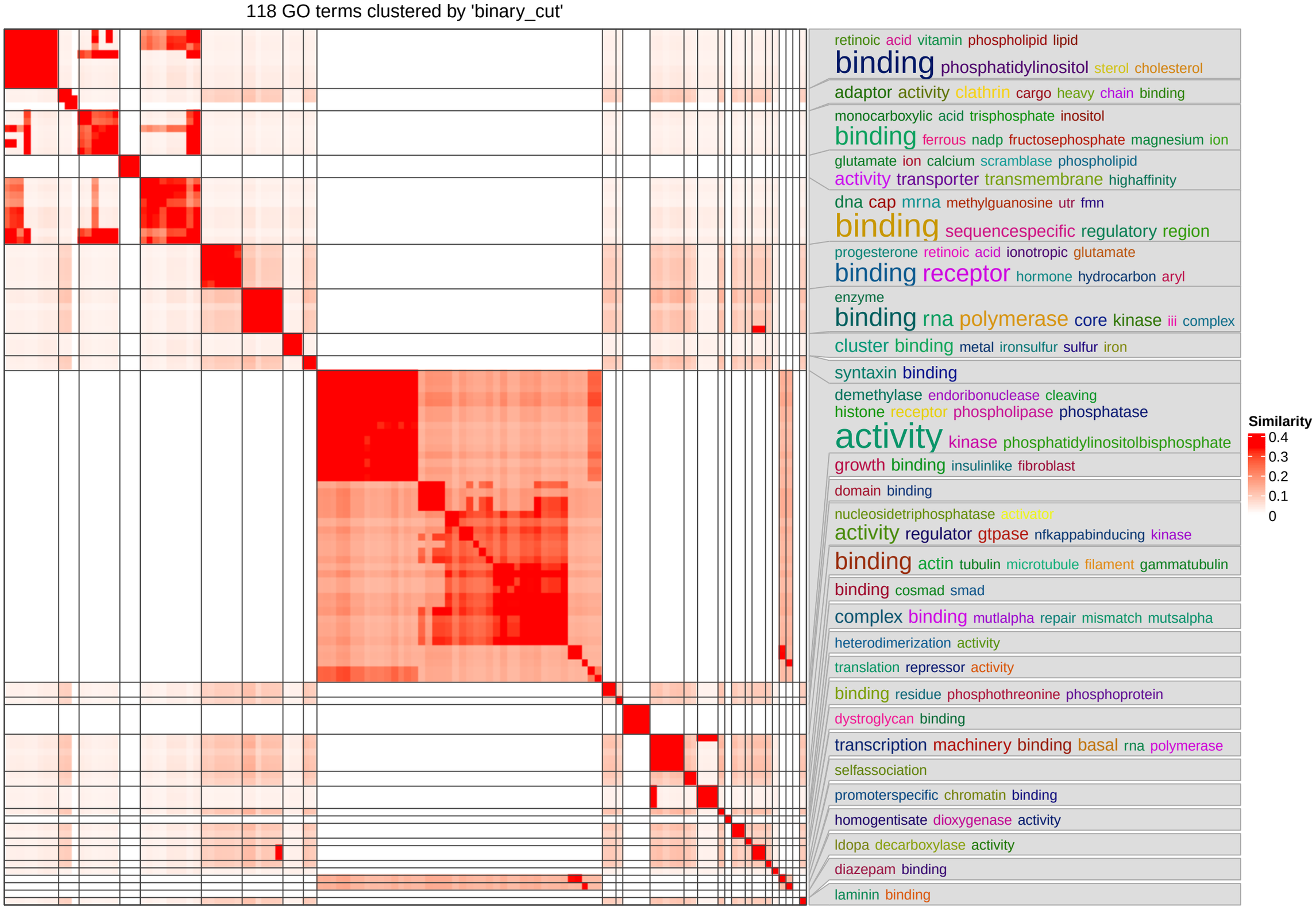
118 GO terms clustered by 'binary_cut'
retinoic acid vitamin phospholipid lipid
binding phosphatidylinositol sterol cholesterol
adaptor activity clathrin cargo heavy chain binding
monocarboxylic acid trisphosphate inositol
binding ferrous nadp fructosephosphate magnesium ion
glutamate ion calcium scramblase phospholipid
activity transporter transmembrane highaffinity
dna cap mrna methylguanosine utr fmn
binding sequencespecific regulatory region
progesterone retinoic acid ionotropic glutamate
binding receptor hormone hydrocarbon aryl
enzyme
binding rna polymerase core kinase iii complex
cluster binding metal ironsulfur sulfur iron
syntaxin binding
demethylase endoribonuclease cleaving
histone receptor phospholipase phosphatase
activity kinase phosphatidylinositolbisphosphate
growth binding insulinlike fibroblast
domain binding
nucleosidetriphosphatase activator
activity regulator gtpase nfkappabinducing kinase
binding actin tubulin microtubule filament gammatubulin
binding cosmad smad
complex binding mutlalpha repair mismatch mutsalpha
heterodimerization activity
translation repressor activity
binding residue phosphothreonine phosphoprotein
dystroglycan binding
transcription machinery binding basal rna polymerase
selfassociation
promoterspecific chromatin binding
homogentisate dioxygenase activity
ldopa decarboxylase activity
diazepam binding
laminin binding
Similarity
0.4
0.3
0.2
0.1
0