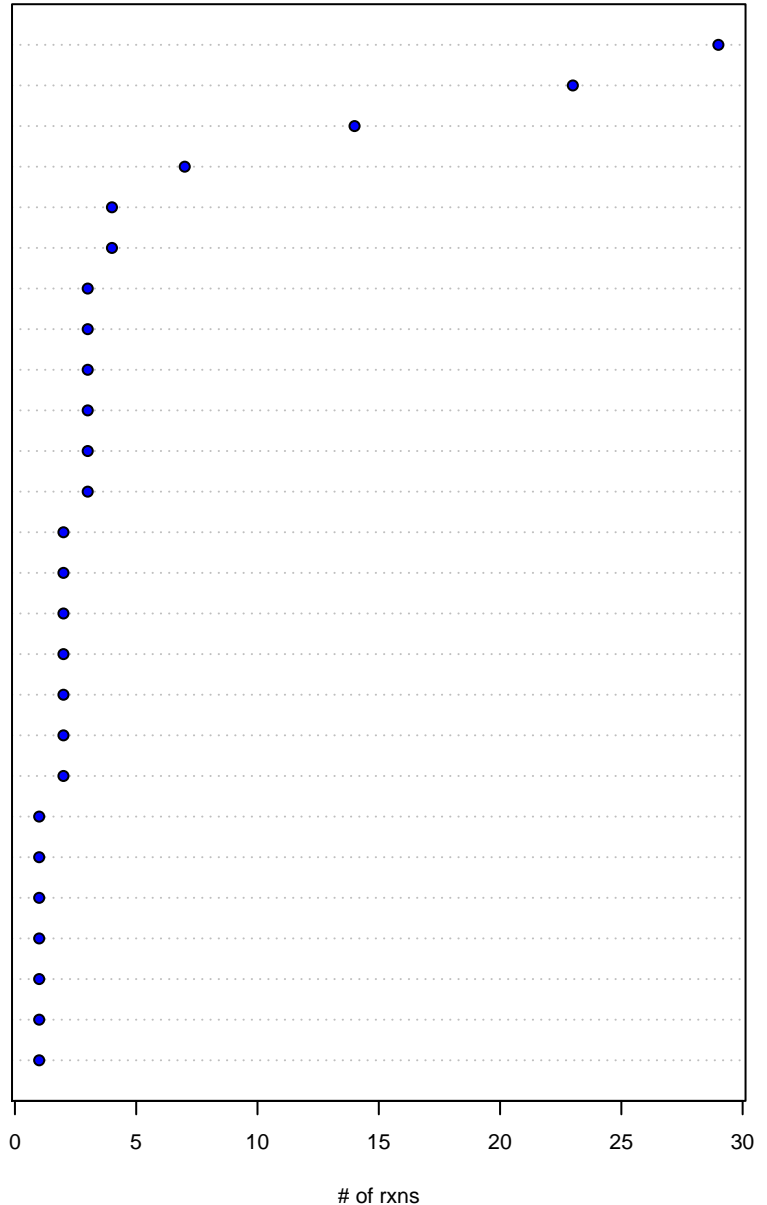
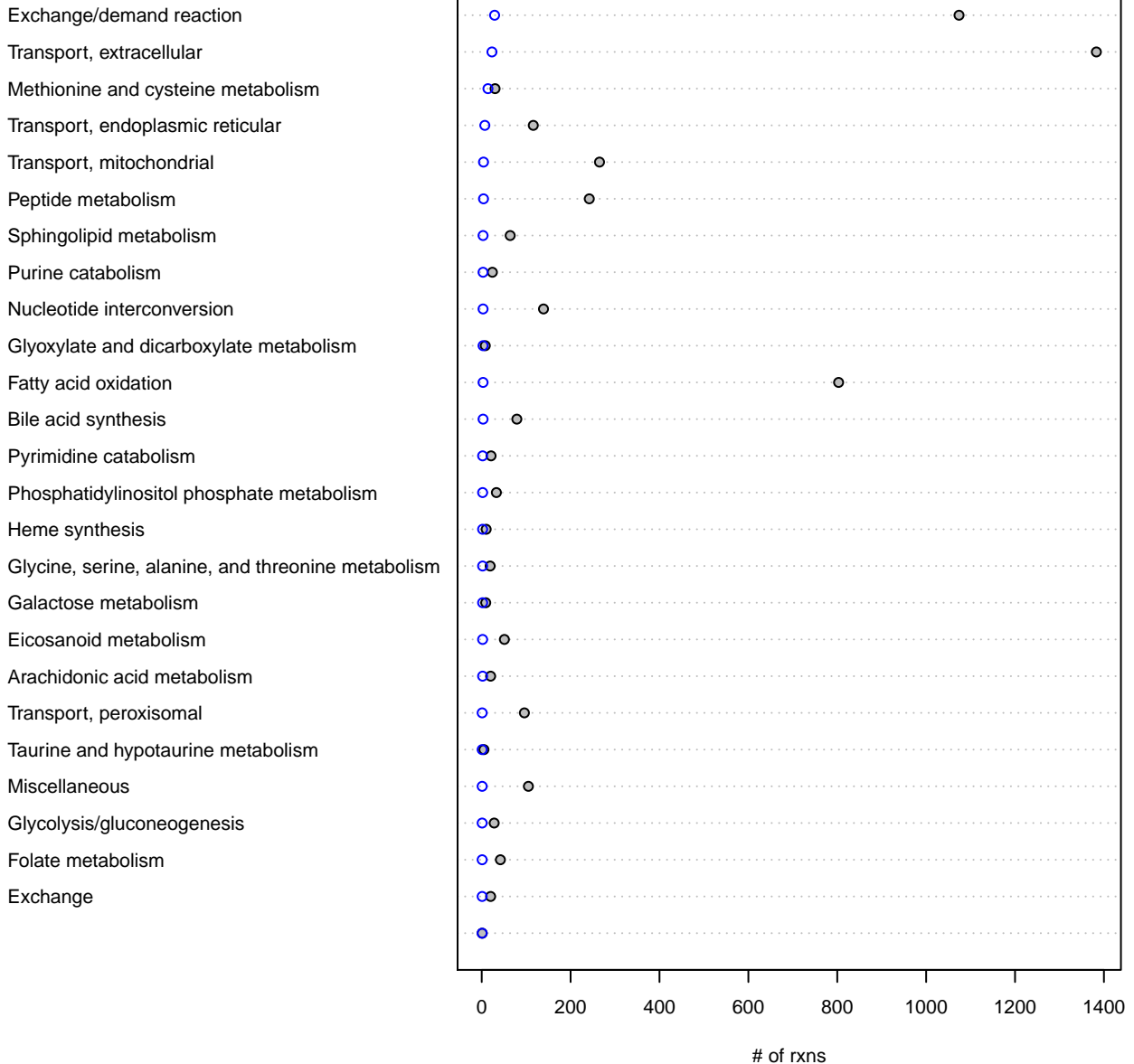


# of disrupted rxns (n=120, ST)

Exchange/demand reaction  
Transport, extracellular  
Methionine and cysteine metabolism  
Transport, endoplasmic reticular  
Transport, mitochondrial  
Peptide metabolism  
Sphingolipid metabolism  
Purine catabolism  
Nucleotide interconversion  
Glyoxylate and dicarboxylate metabolism  
Fatty acid oxidation  
Bile acid synthesis  
Pyrimidine catabolism  
Phosphatidylinositol phosphate metabolism  
Heme synthesis  
Glycine, serine, alanine, and threonine metabolism  
Galactose metabolism  
Eicosanoid metabolism  
Arachidonic acid metabolism  
Transport, peroxisomal  
Taurine and hypotaurine metabolism  
Miscellaneous  
Glycolysis/gluconeogenesis  
Folate metabolism  
Exchange

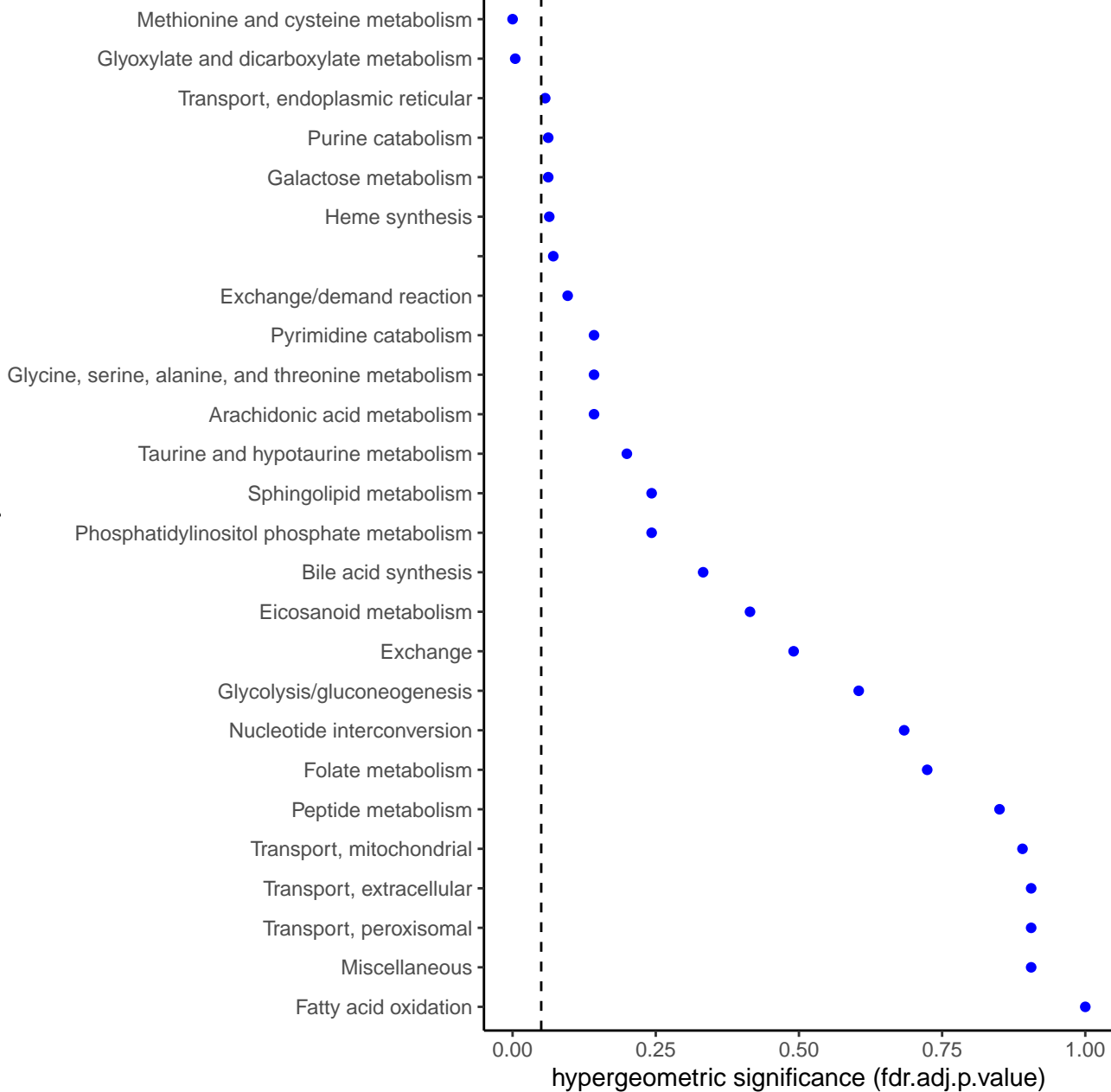


# of disrupted rxns (n=120, ST) vs all rxns

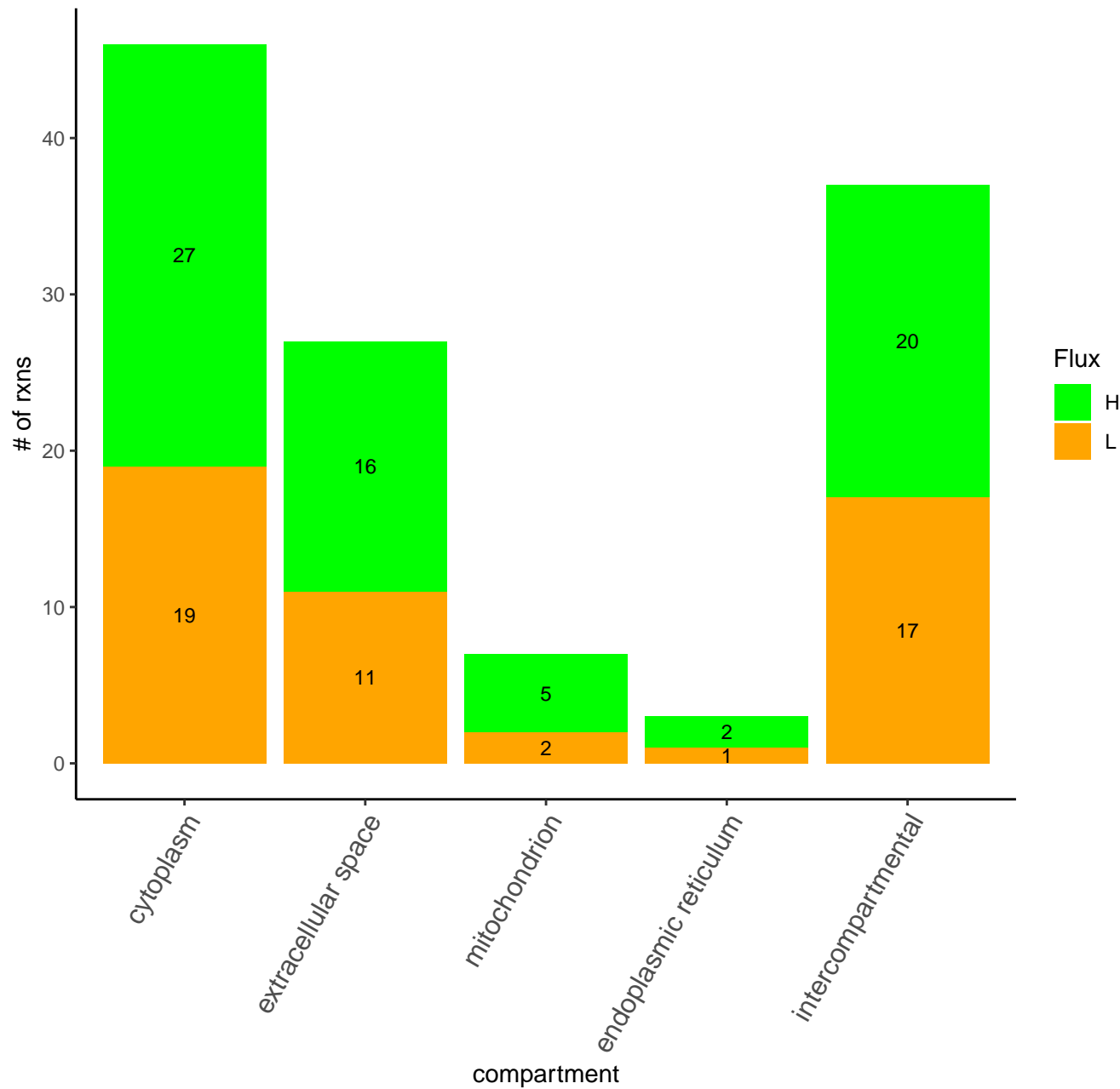


# over-representation analysis, ST

subSystem



Disrupted rxns (all, n=120) per compartment (ST)



Disrupted rxns (fdr.significant, n=17) per compartment (ST)

