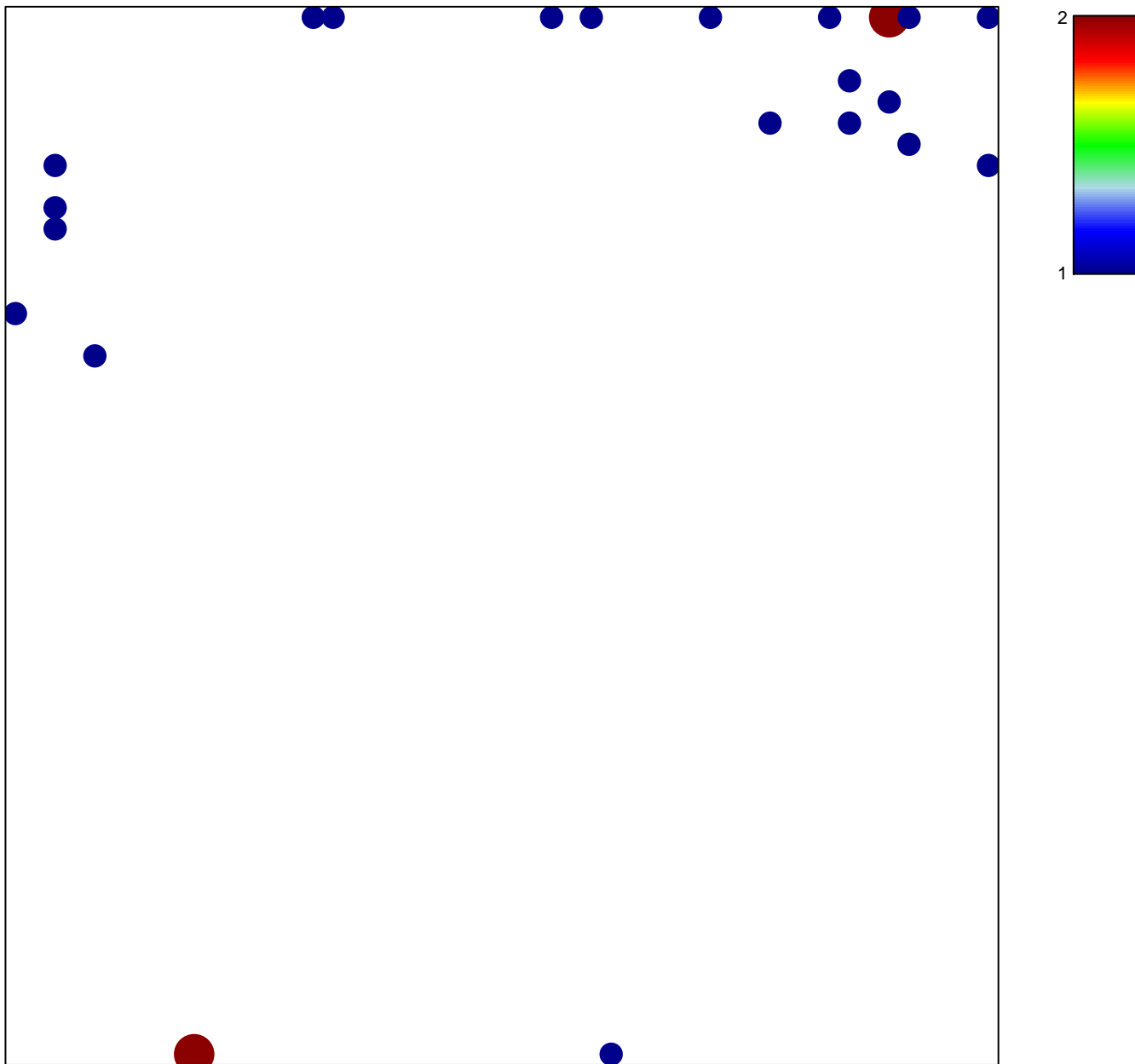
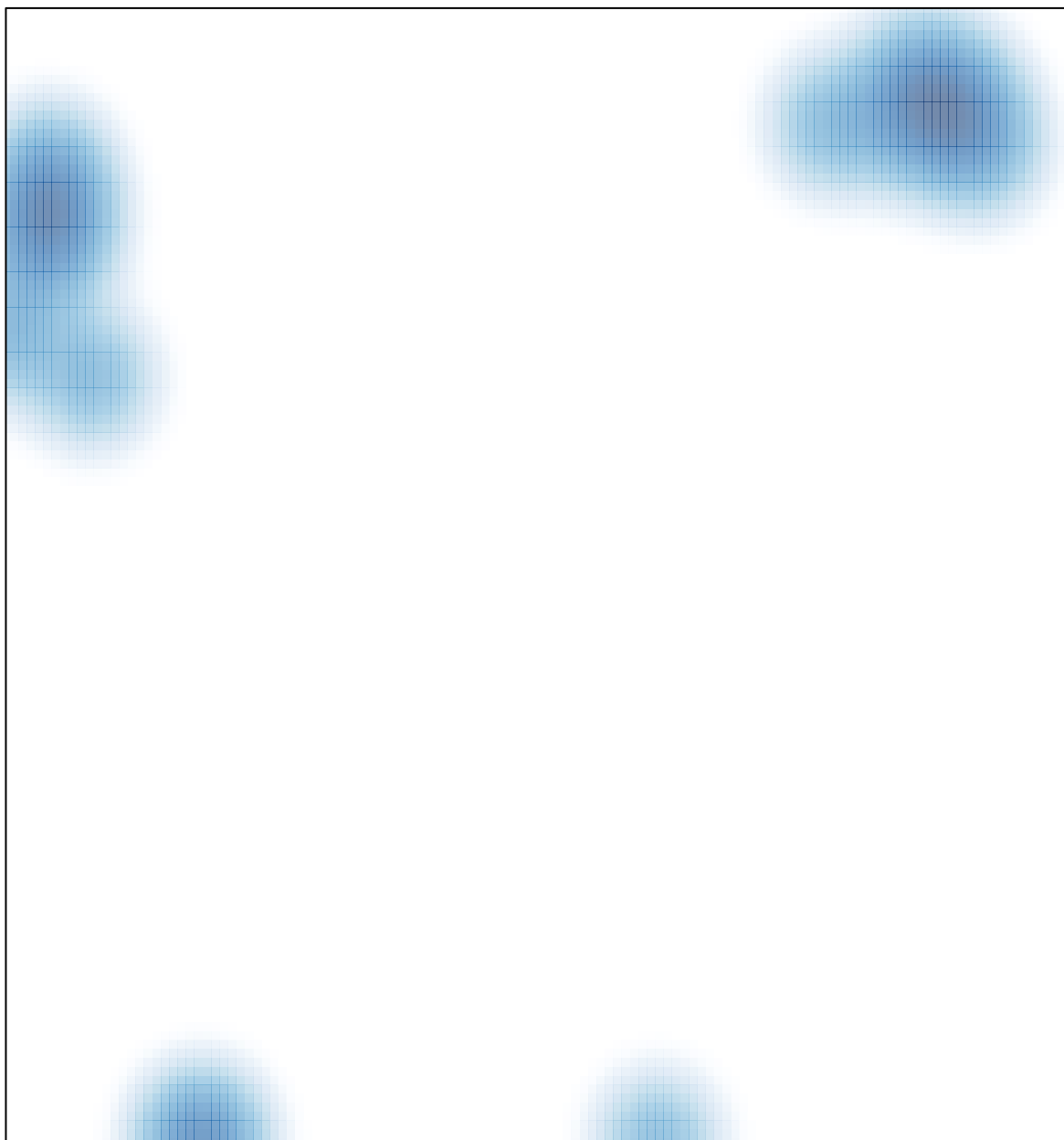


REACTOME_CYTOSOLIC_TRNA_AMINOACYLATION



features = 24
chi-square p = 0.73

REACTOME_CYTOSOLIC_TRNA_AMINOACYLATION



features = 24 , max = 2