REACTOME_PROTEIN_FOLDING
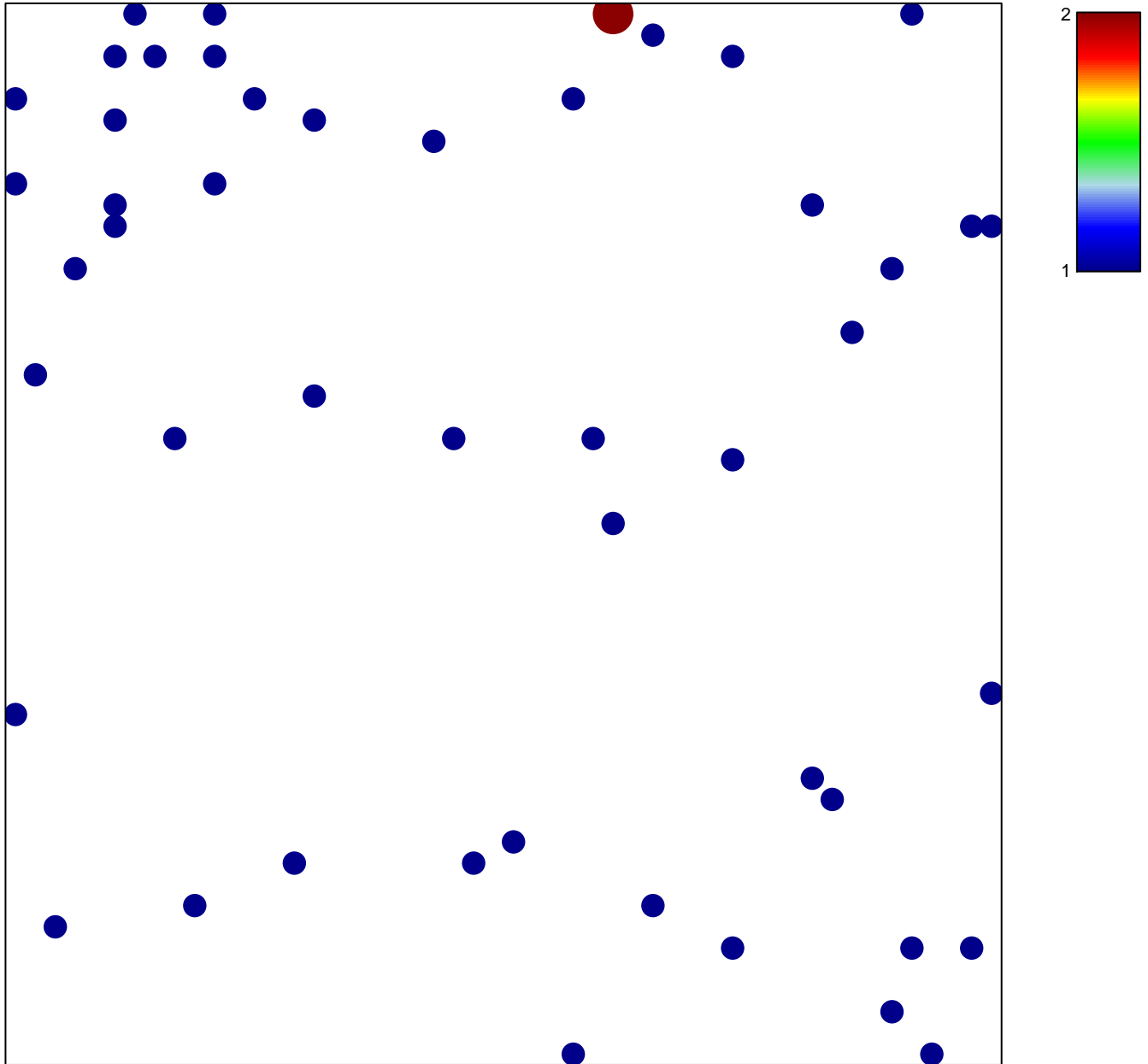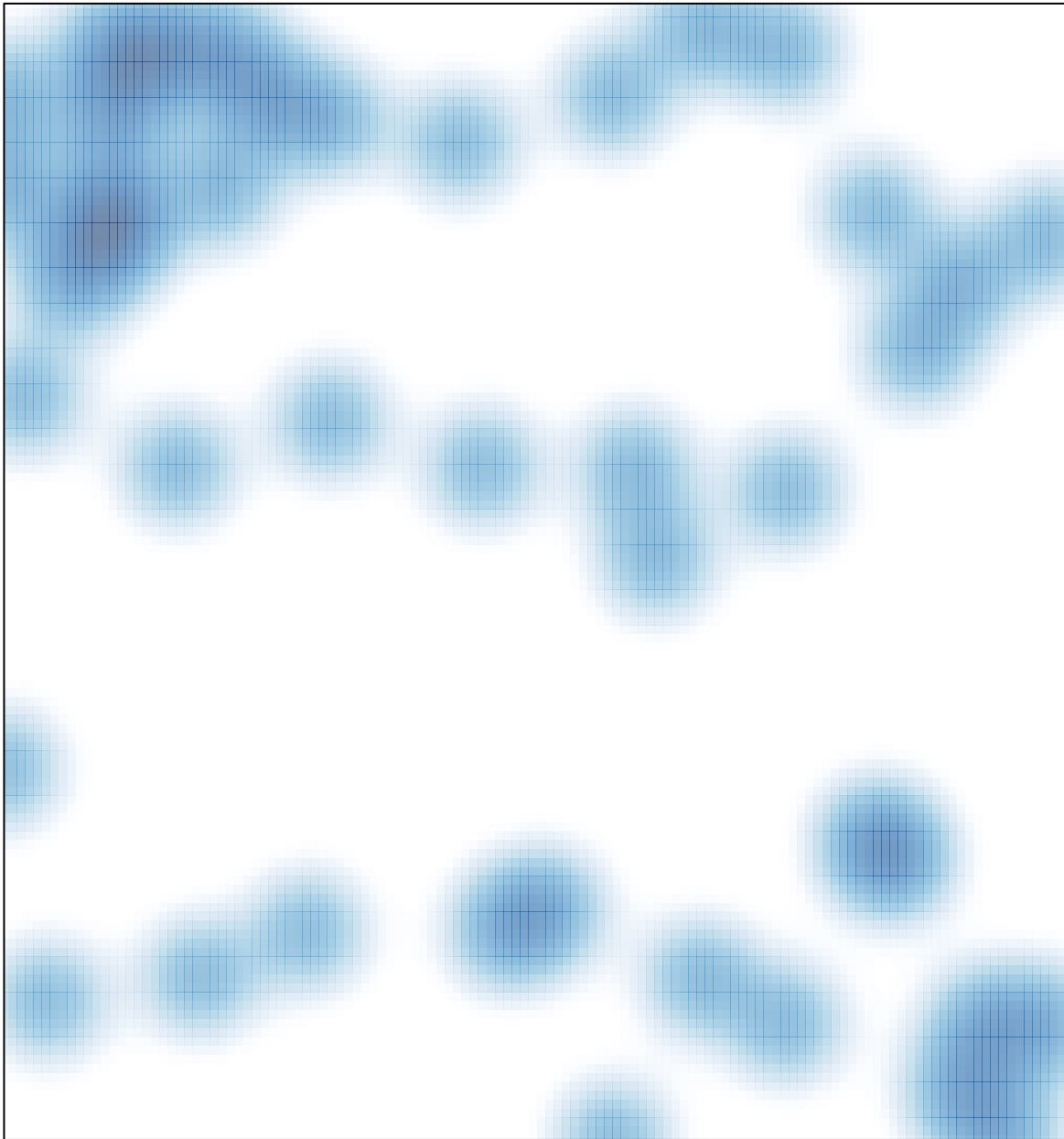
features = 49
chi−square p = 0.82

REACTOME_PROTEIN_FOLDING

features = 49 , max = 2