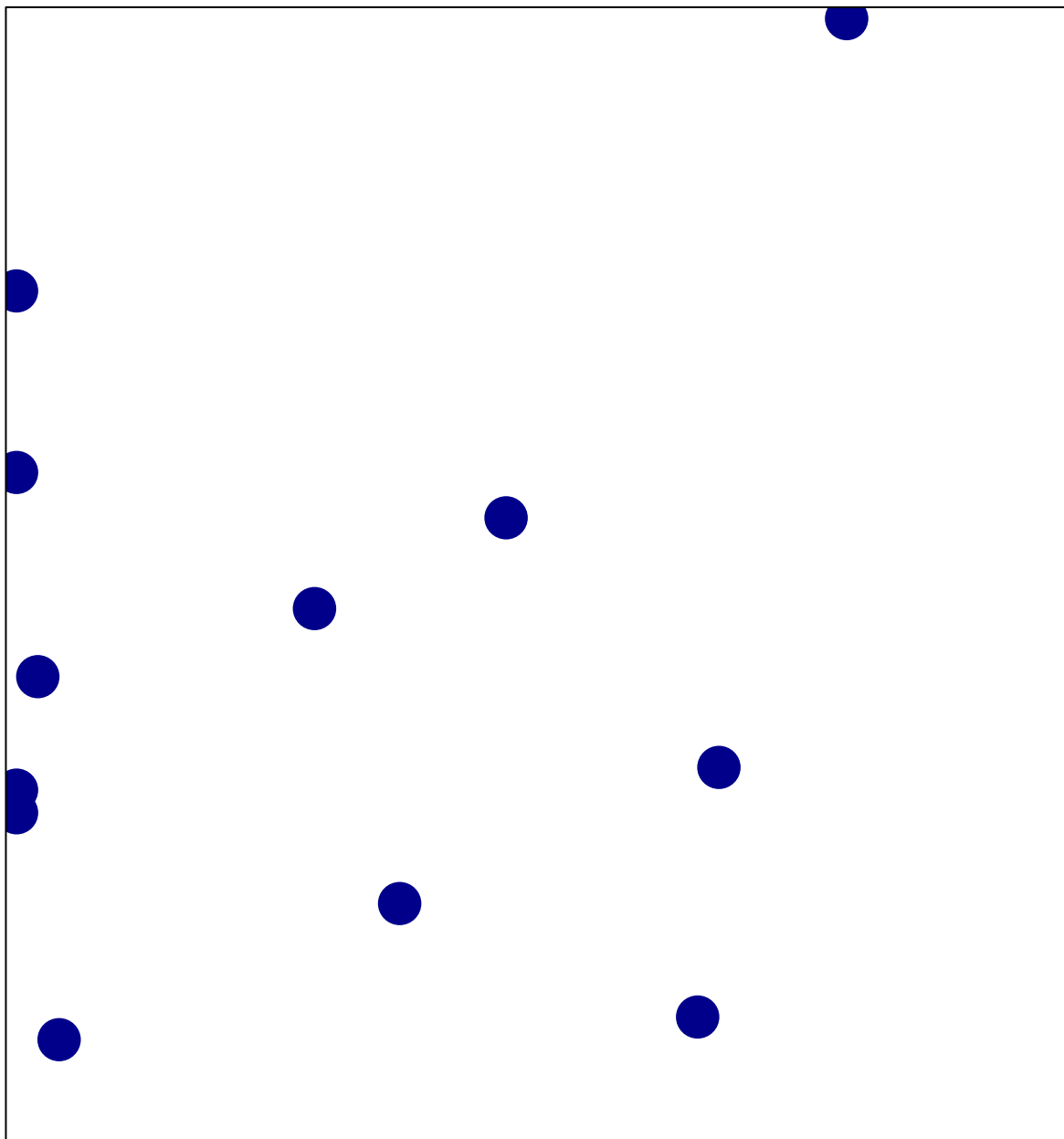
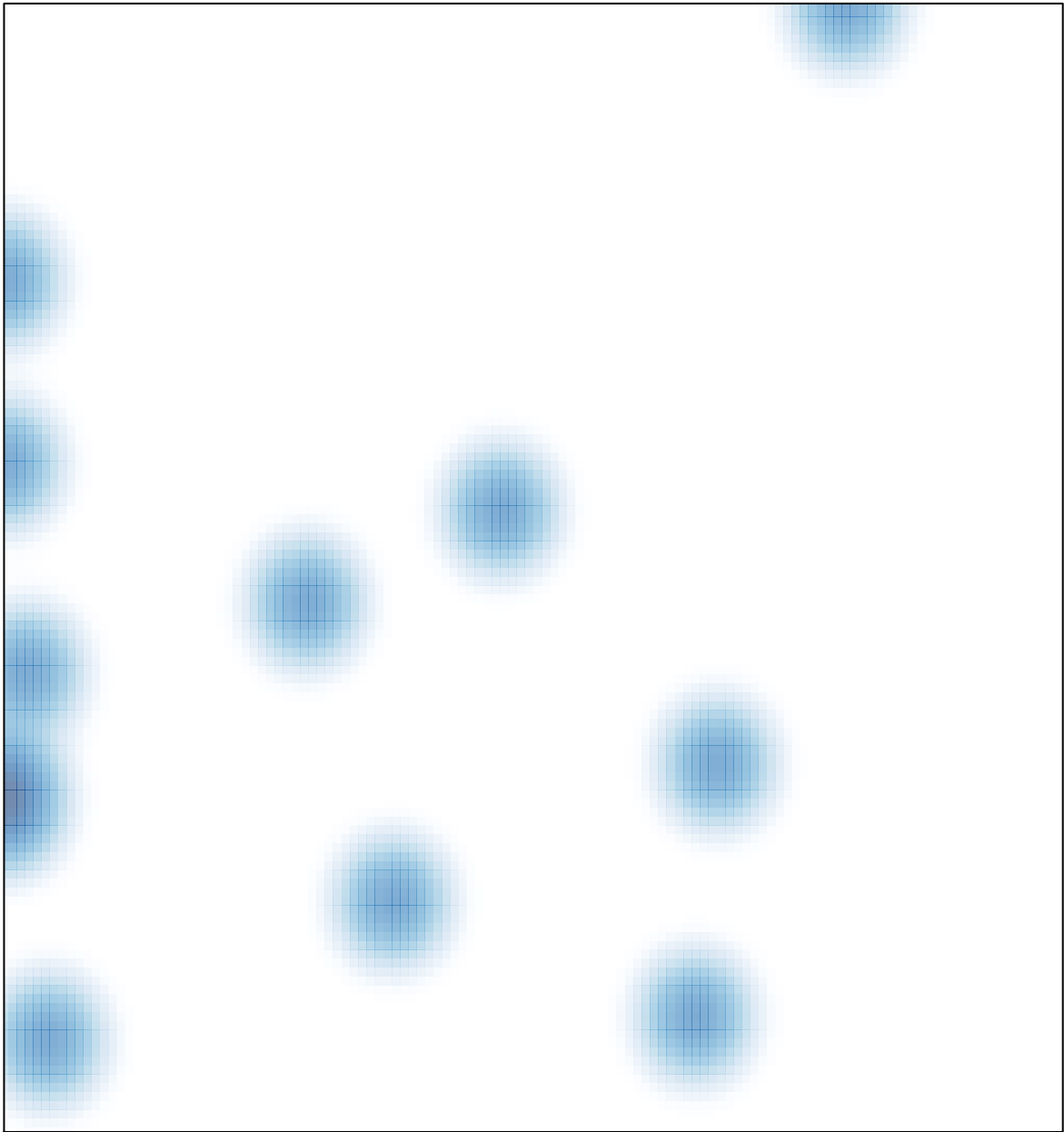


REACTOME_CHAPERONIN_MEDIATED_PROTEIN_FOLDING



features = 12
chi-square p = 0.82

ACTOME_CHAPERONIN_MEDIATED_PROTEIN_FOLD



features = 12 , max = 1