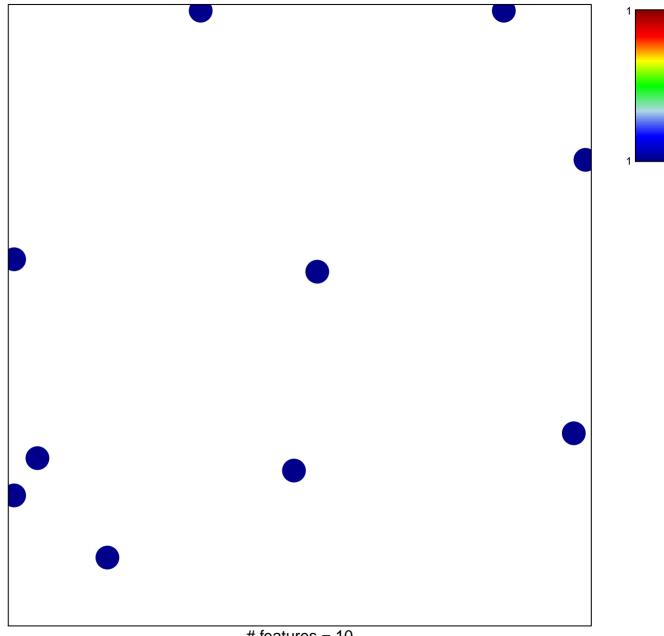
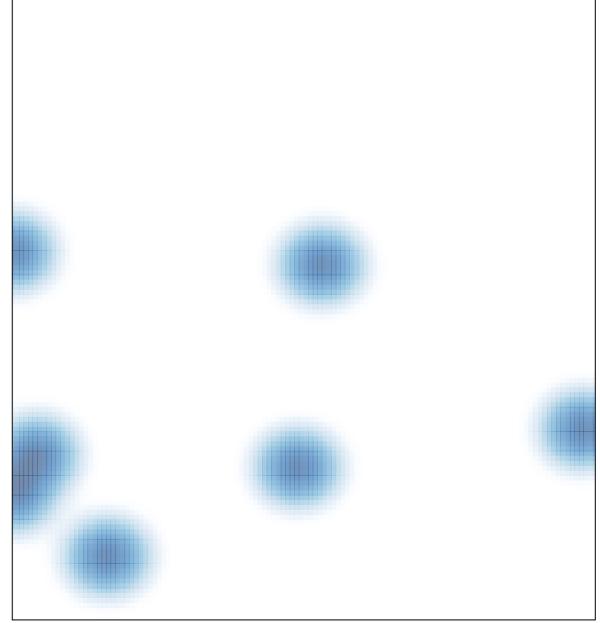
REACTOME_REGULATED_PROTEOLYSIS_OF_P75NTR



features = 10chi-square p = 0.84

REACTOME_REGULATED_PROTEOLYSIS_OF_P75N1



features = 10, max = 1