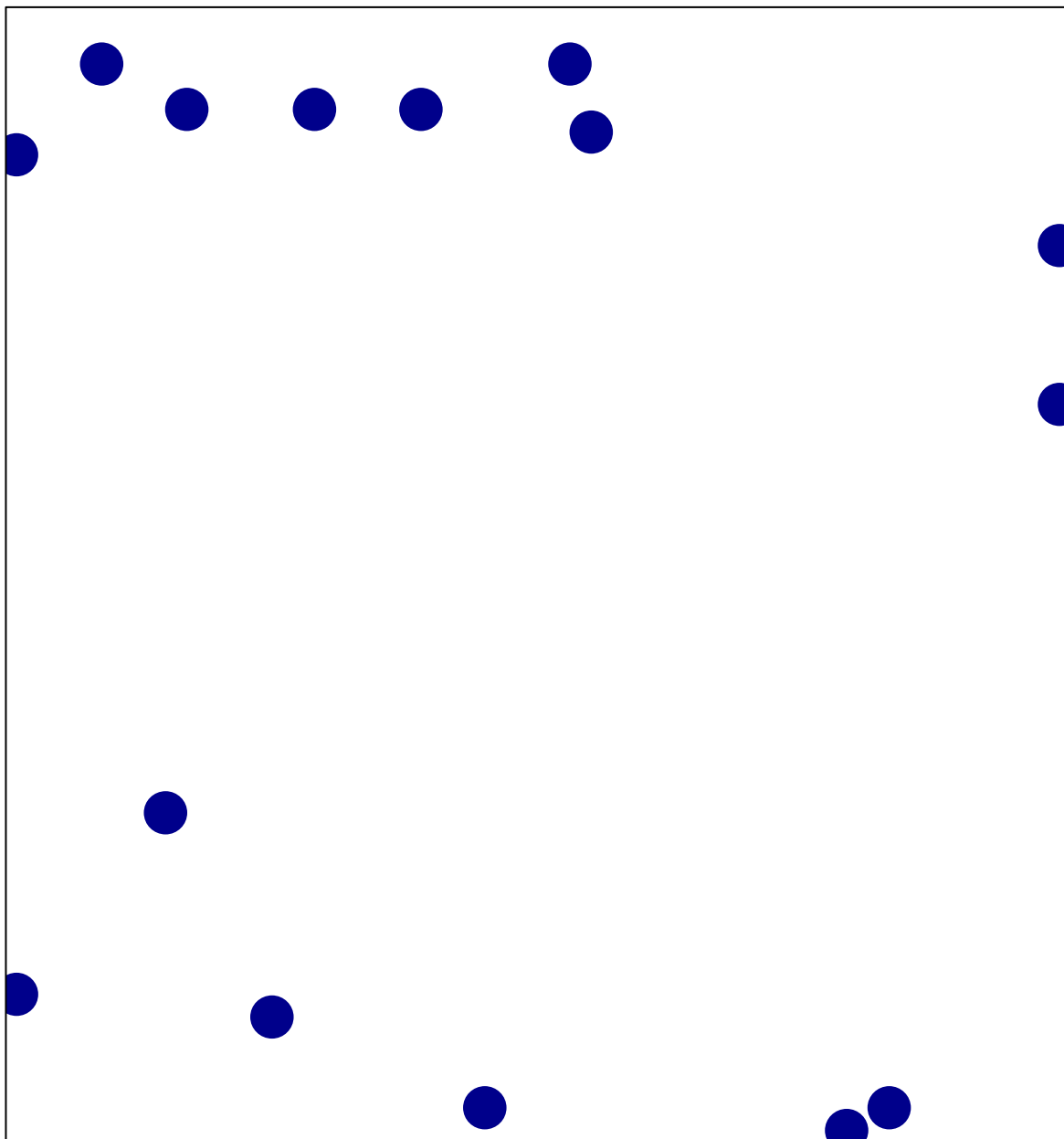


REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS



features = 15
chi-square p = 0.83

ACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOL



features = 15 , max = 1