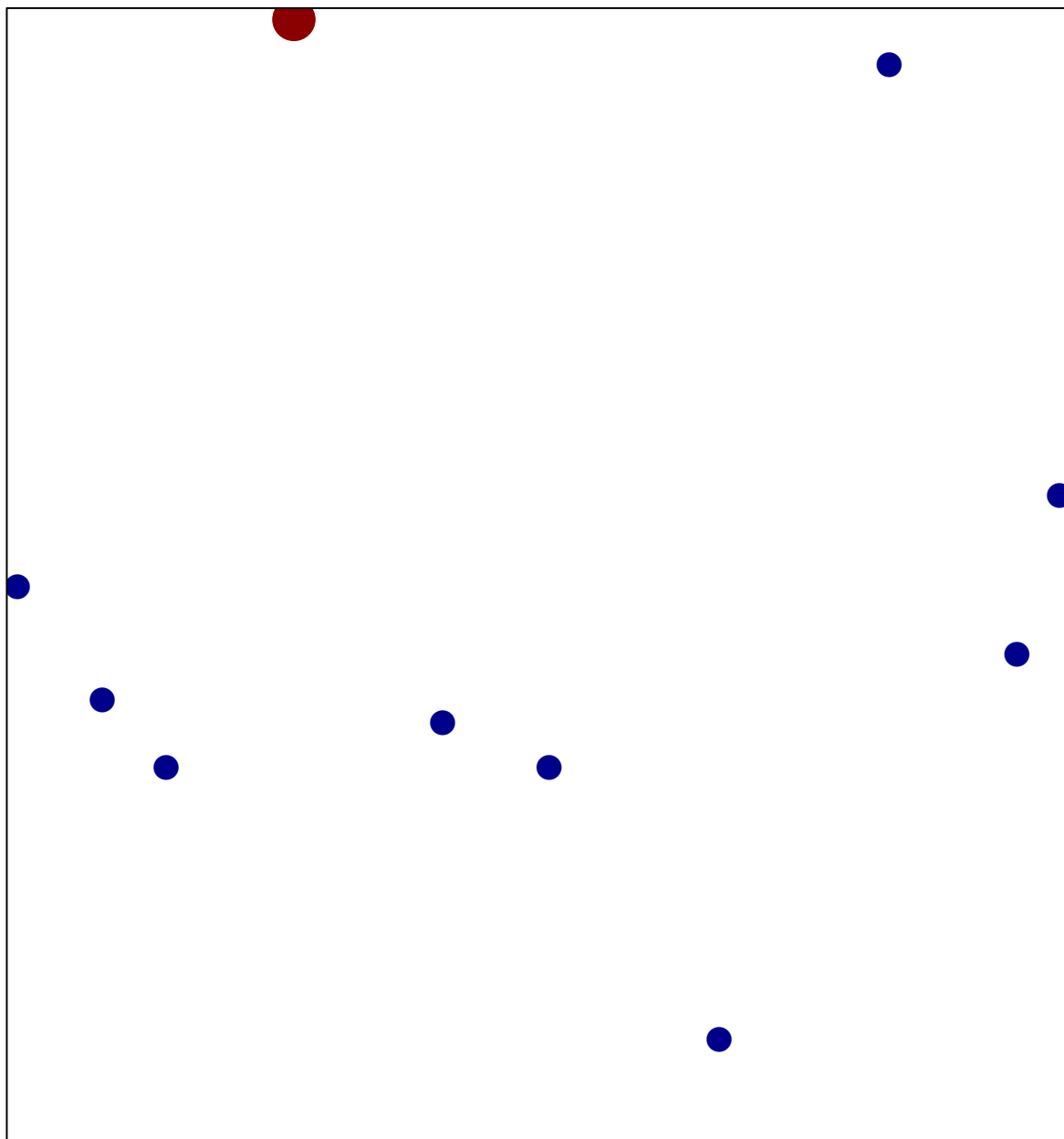


REACTOME_APOPTOSIS_INDUCED_DNA_FRAGMENTATION



features = 11
chi-square p = 0.84

ACTOME_APOPTOSIS_INDUCED_DNA_FRAGMENTA



features = 11 , max = 2