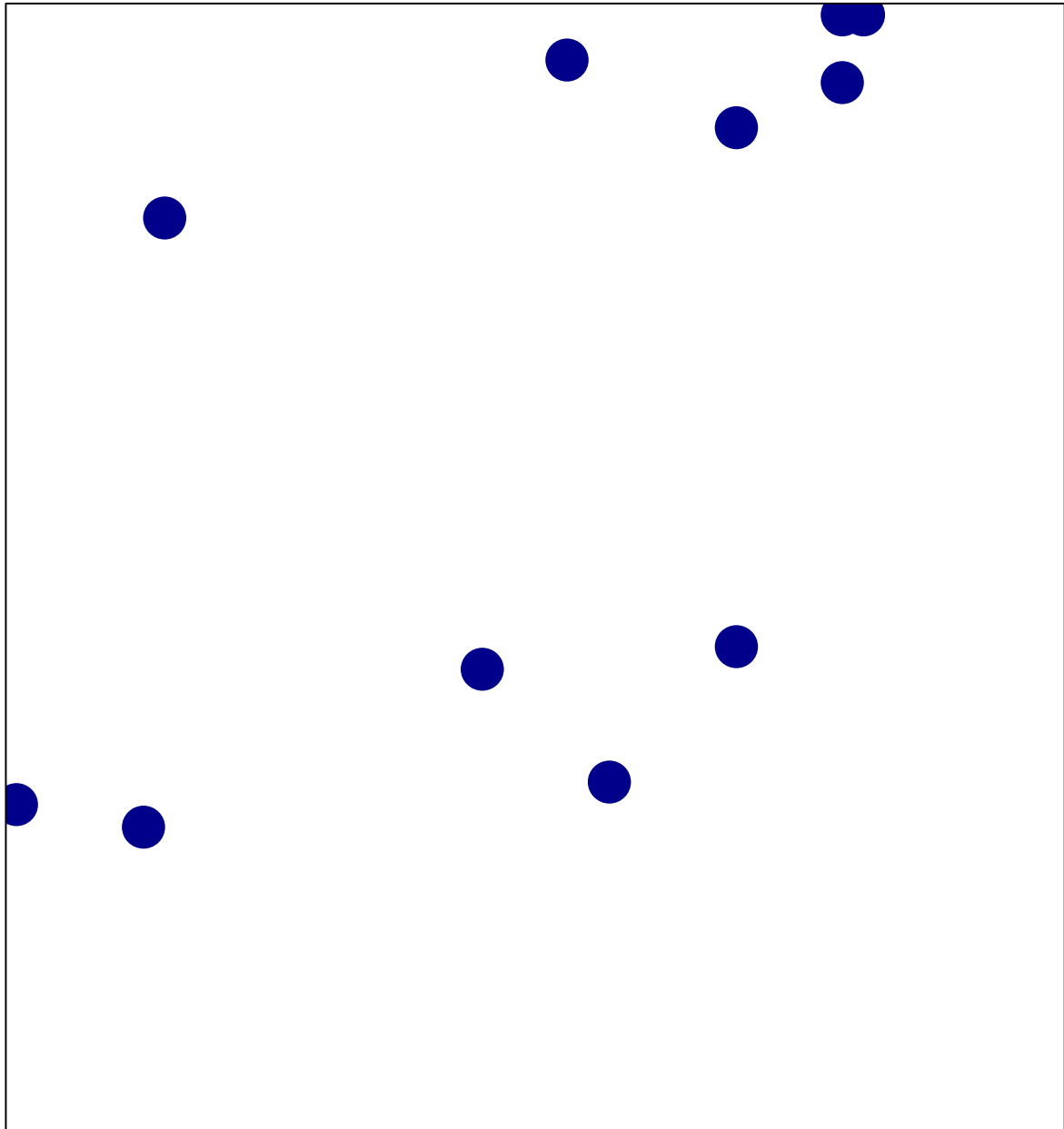


chaperone-mediated protein complex assembly



features = 11
chi-square p = 0.84

chaperone-mediated protein complex assembly



features = 11 , max = 1