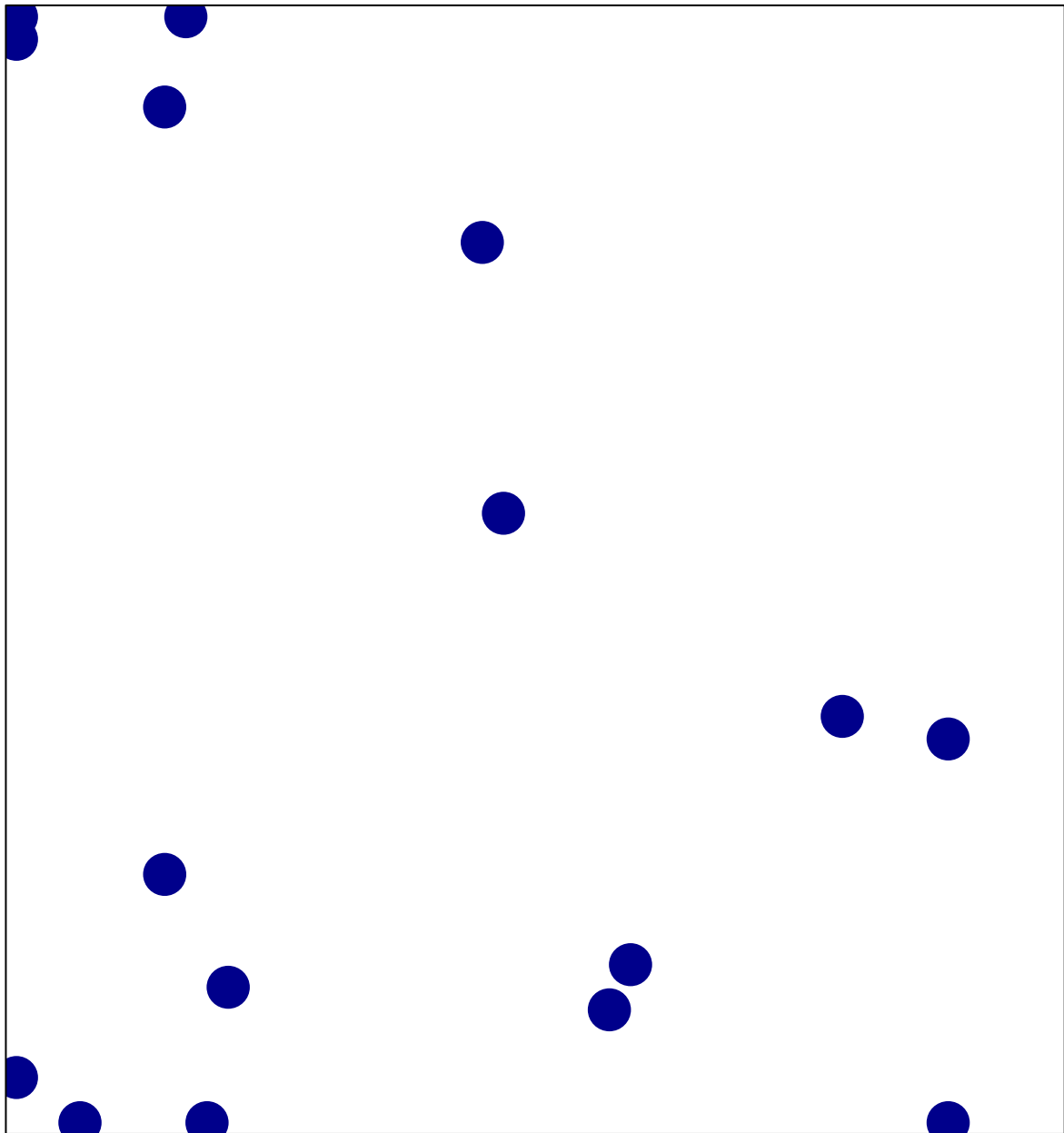
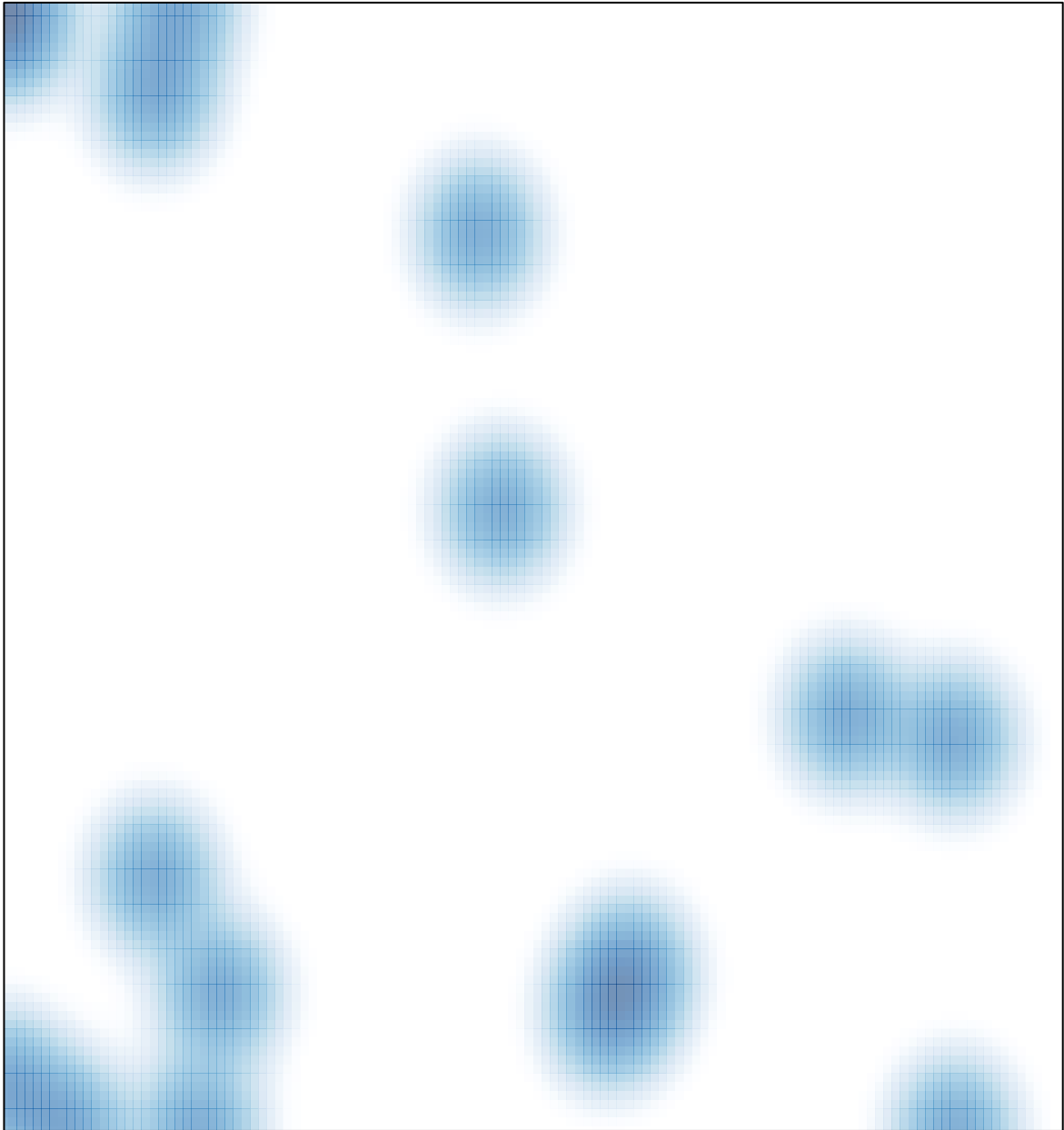


# negative regulation of proteolysis



# features = 16  
chi-square p = 0.82

# negative regulation of proteolysis



# features = 16 , max = 1