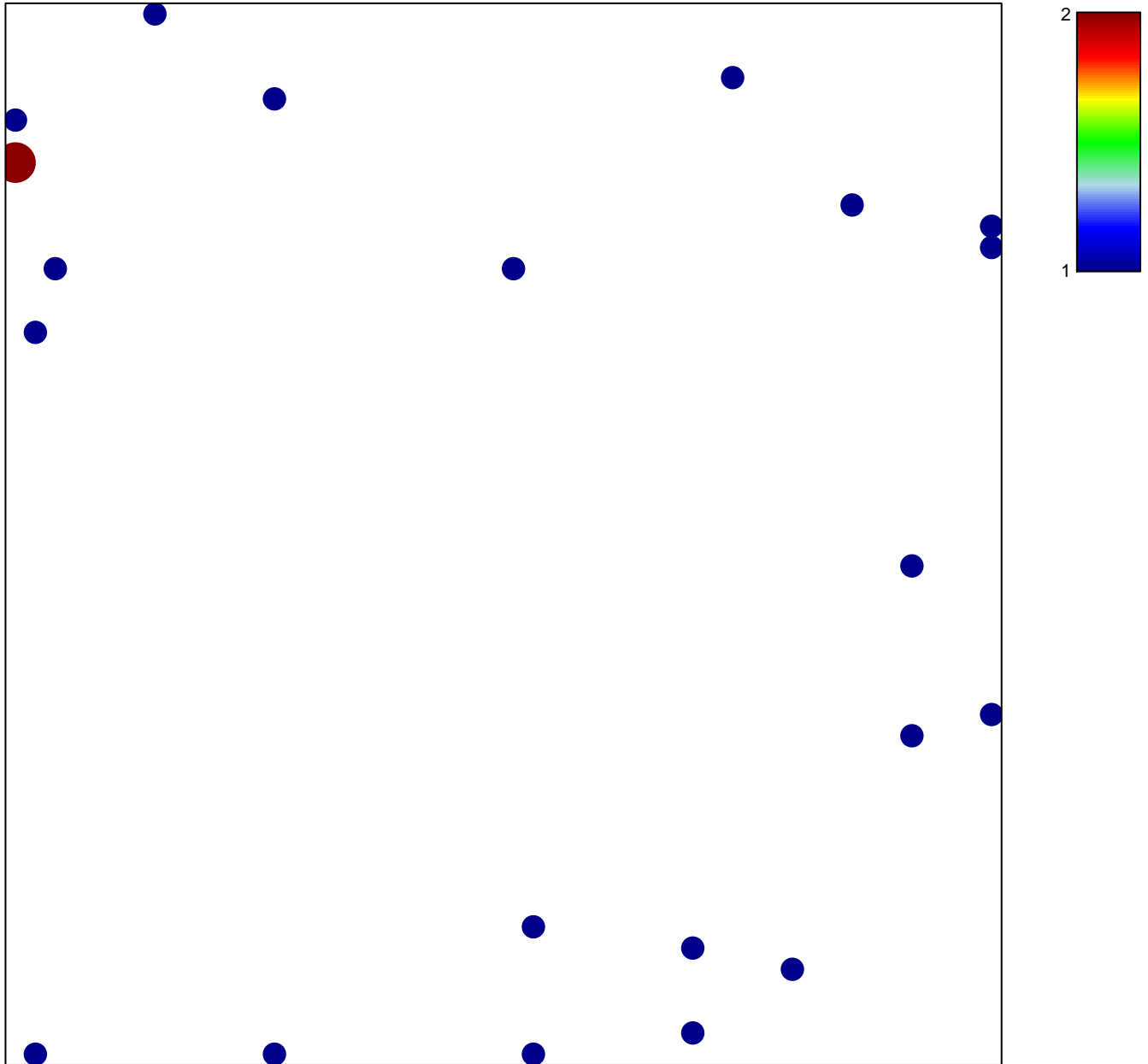
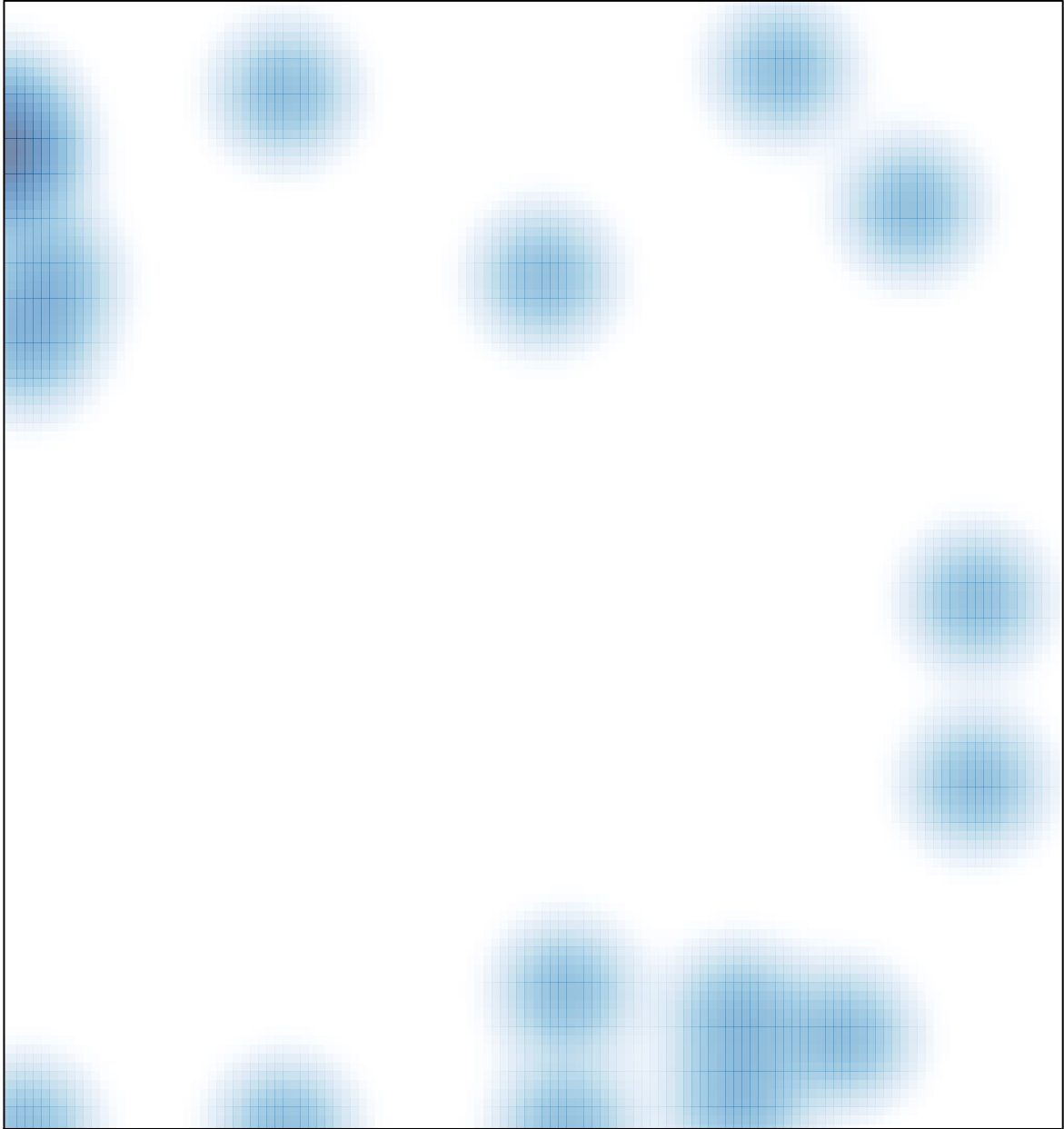


# positive regulation of proteolysis



# features = 22  
chi-square p = 0.82

# positive regulation of proteolysis



# features = 22 , max = 2