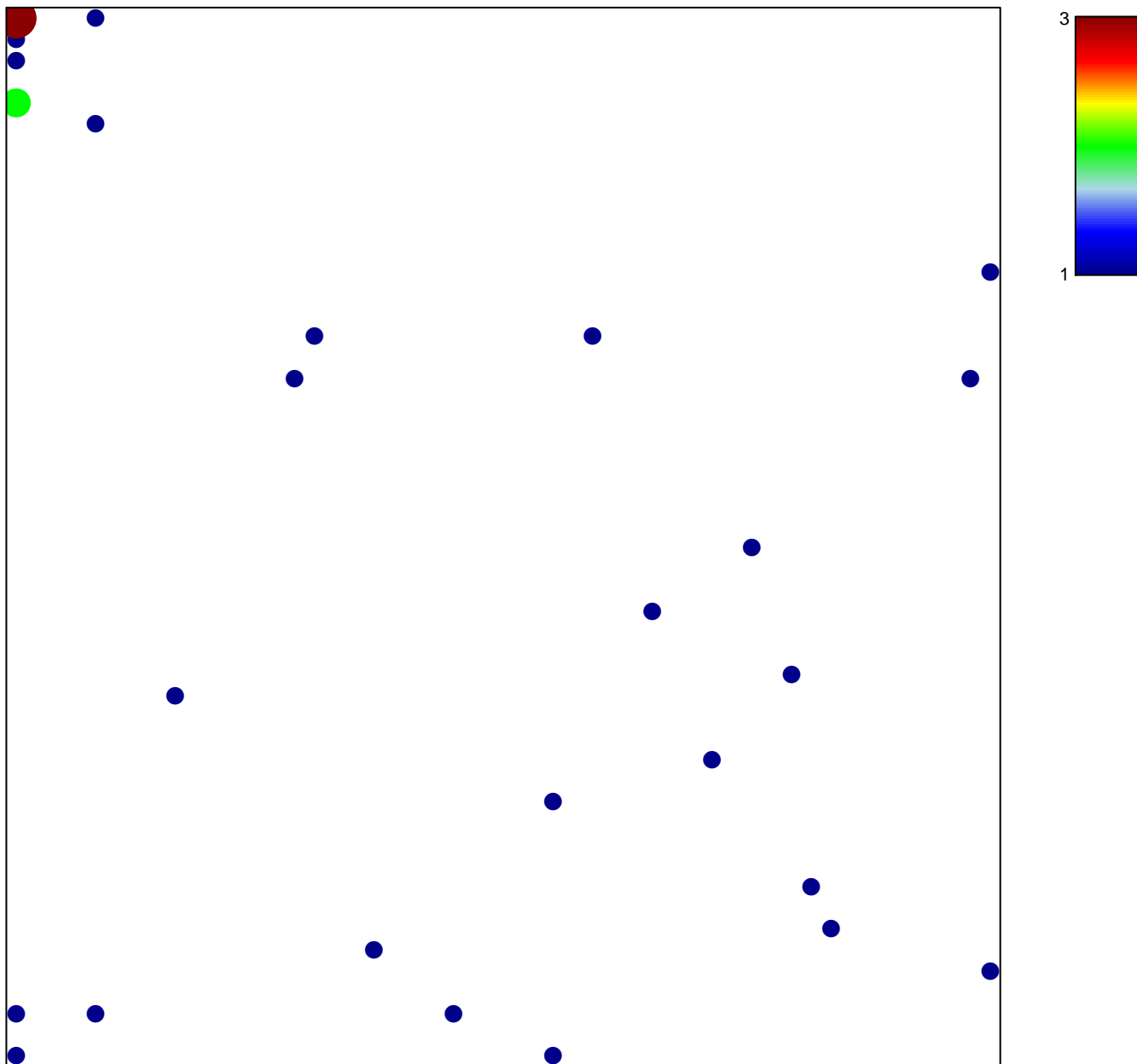
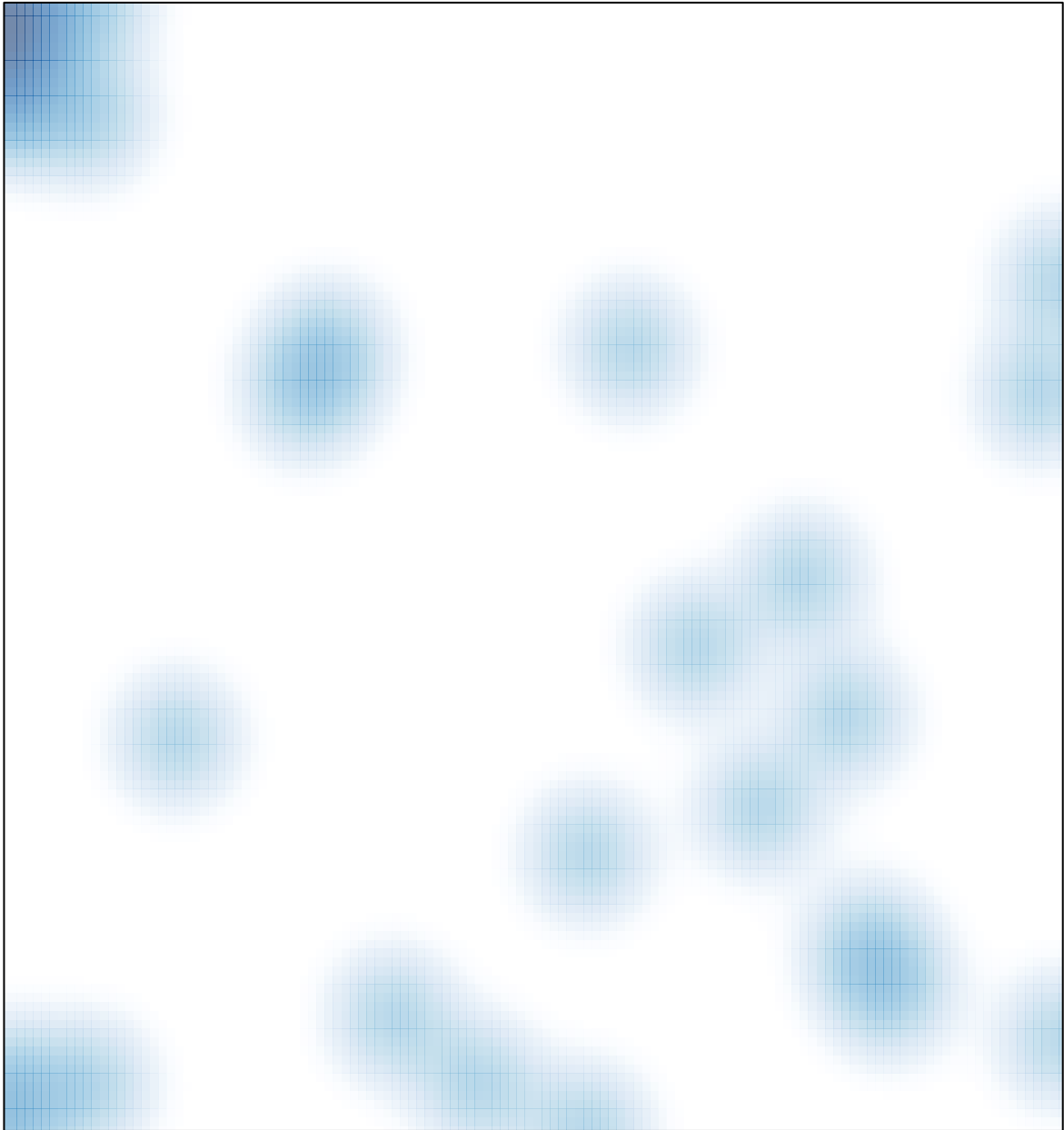


# regulation of proteolysis



# features = 29  
chi-square p = 0.76

# regulation of proteolysis



# features = 29 , max = 3