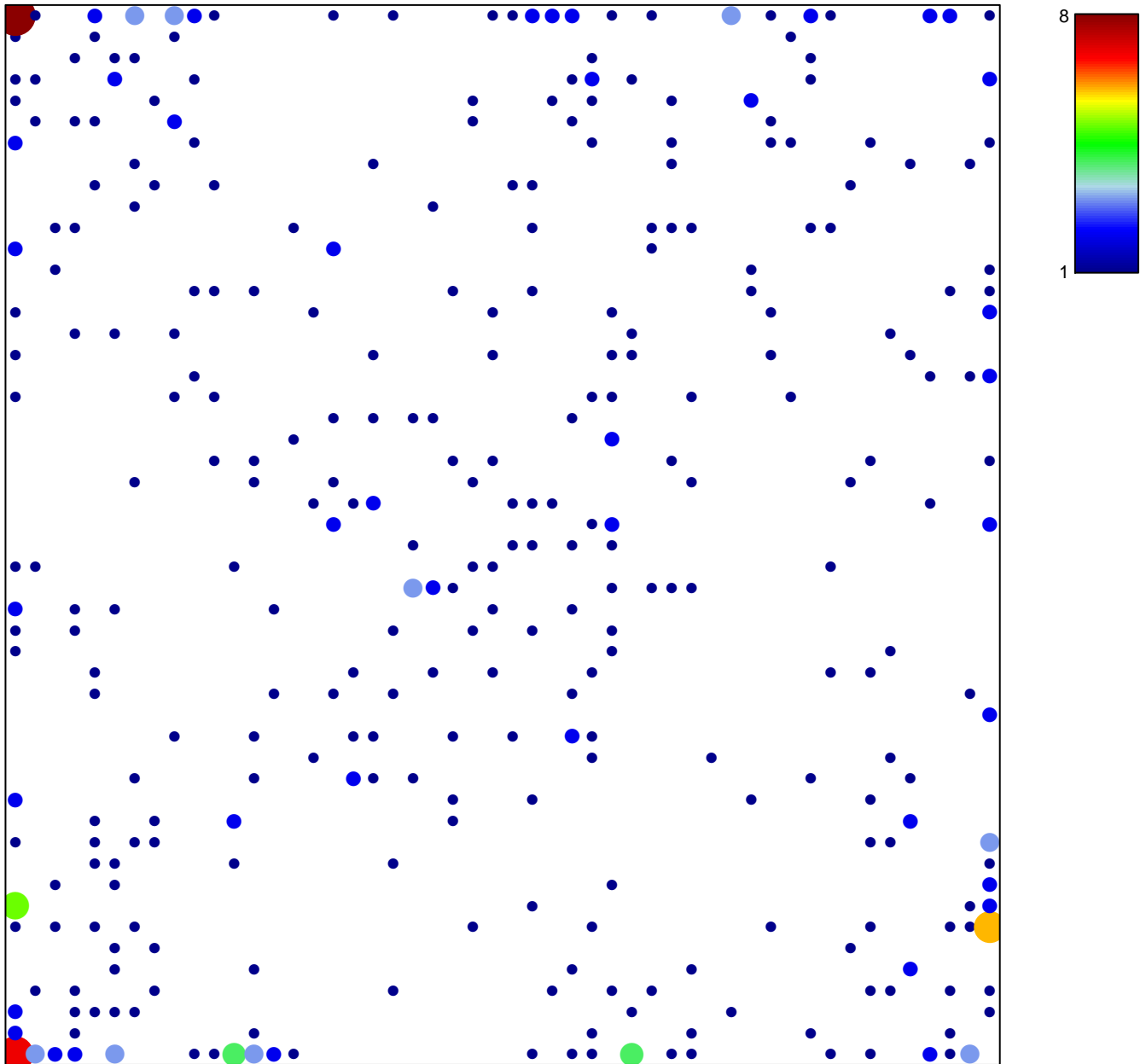
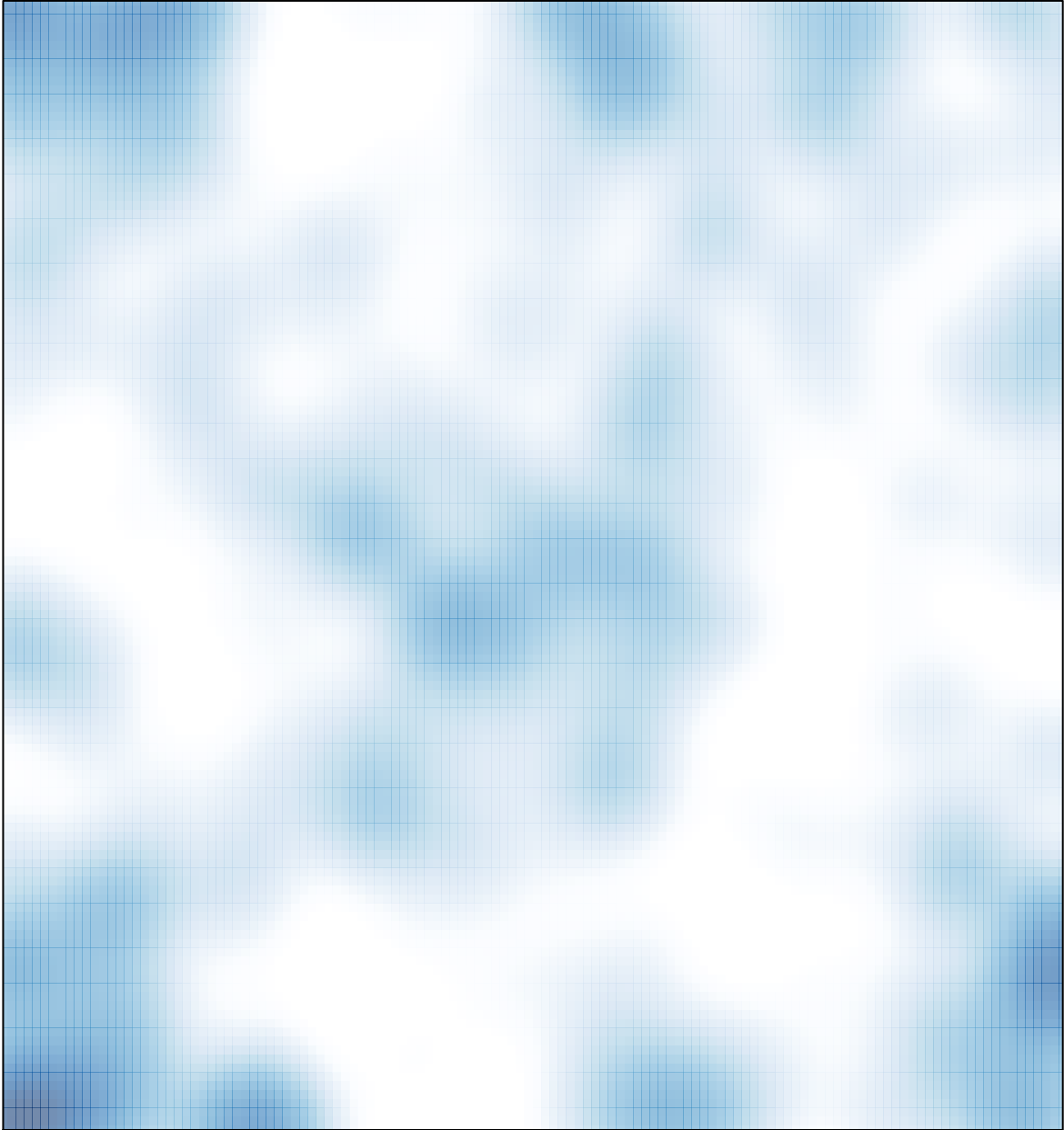


# proteolysis



# features = 408  
chi-square p = 0.61

# proteolysis



# features = 408 , max = 8