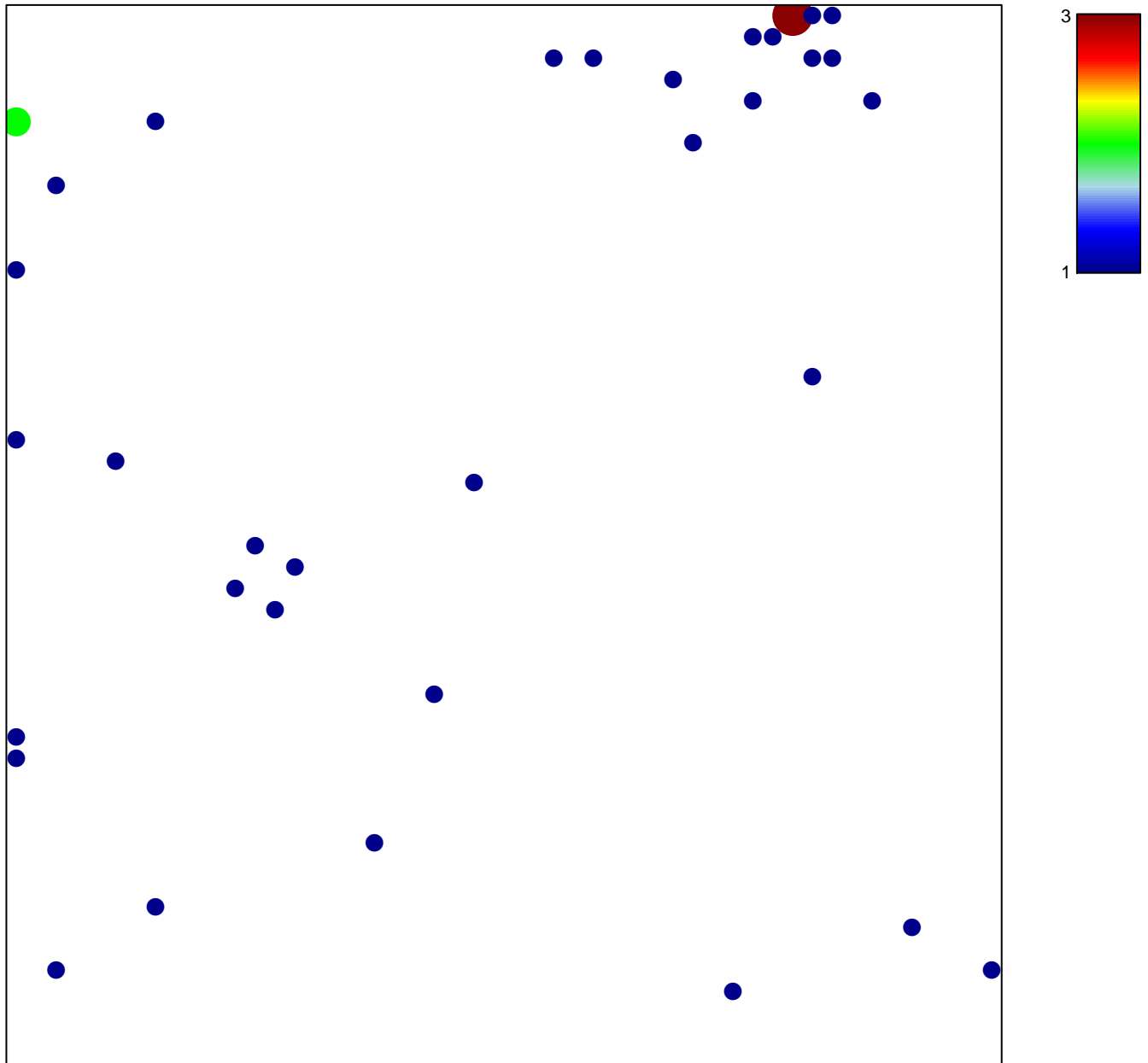
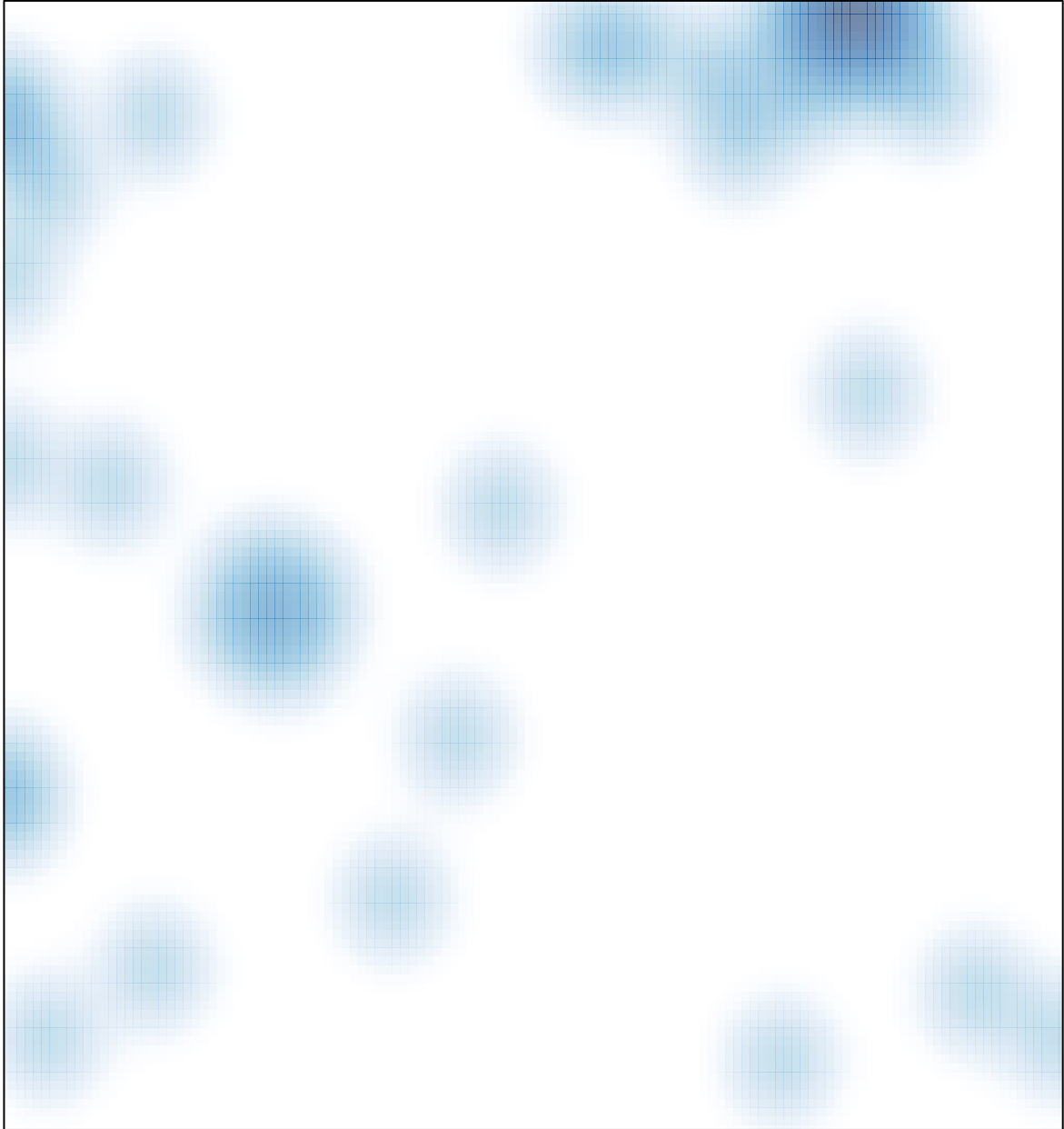


# 'de novo' posttranslational protein folding



# features = 37  
chi-square p = 0.8

# 'de novo' posttranslational protein folding



# features = 37 , max = 3