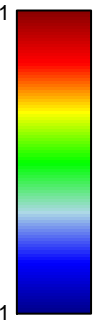
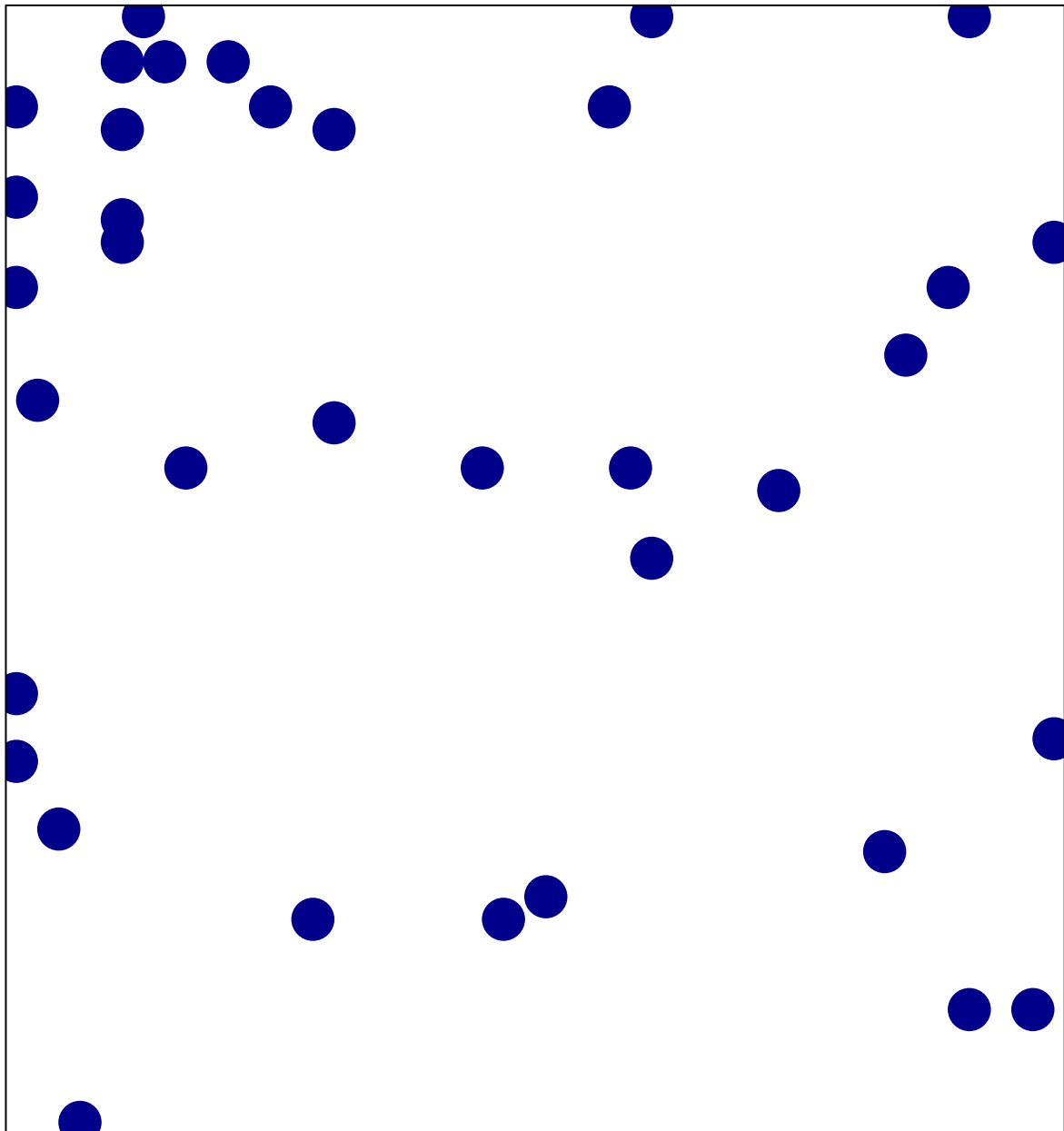
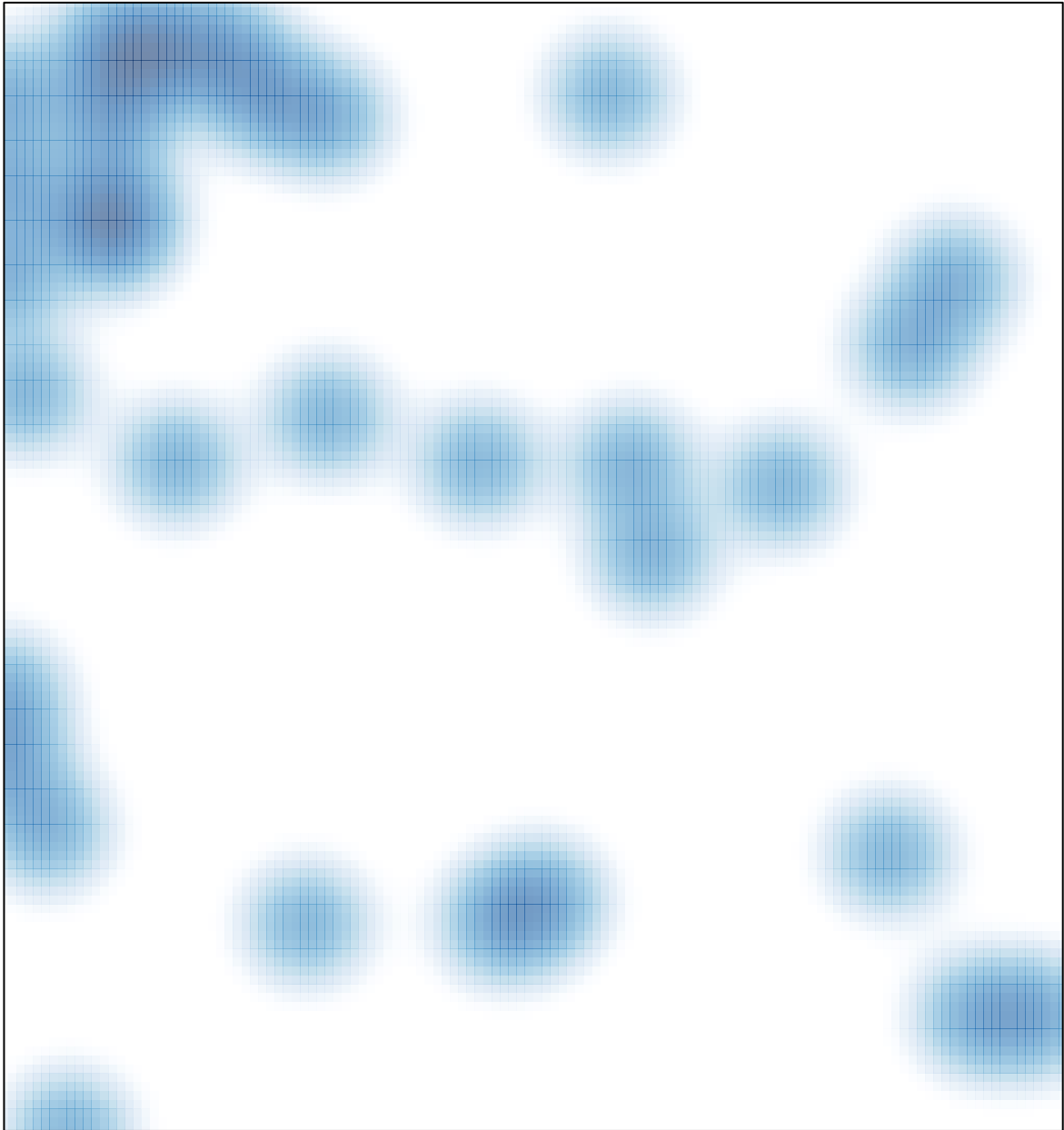


# 'de novo' posttranslational protein folding



# features = 36  
chi-square p = 0.82

# 'de novo' posttranslational protein folding



# features = 36 , max = 1