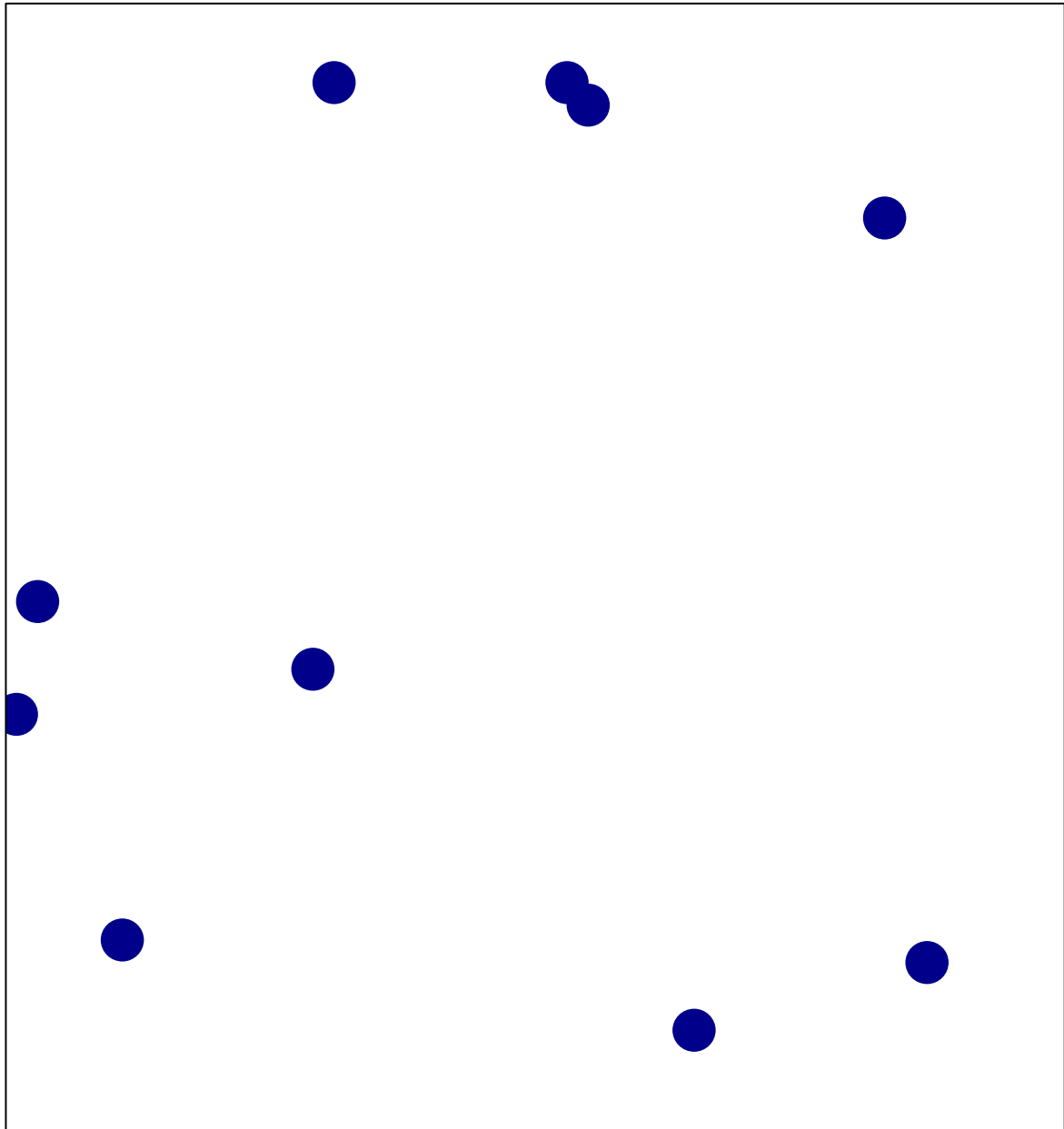
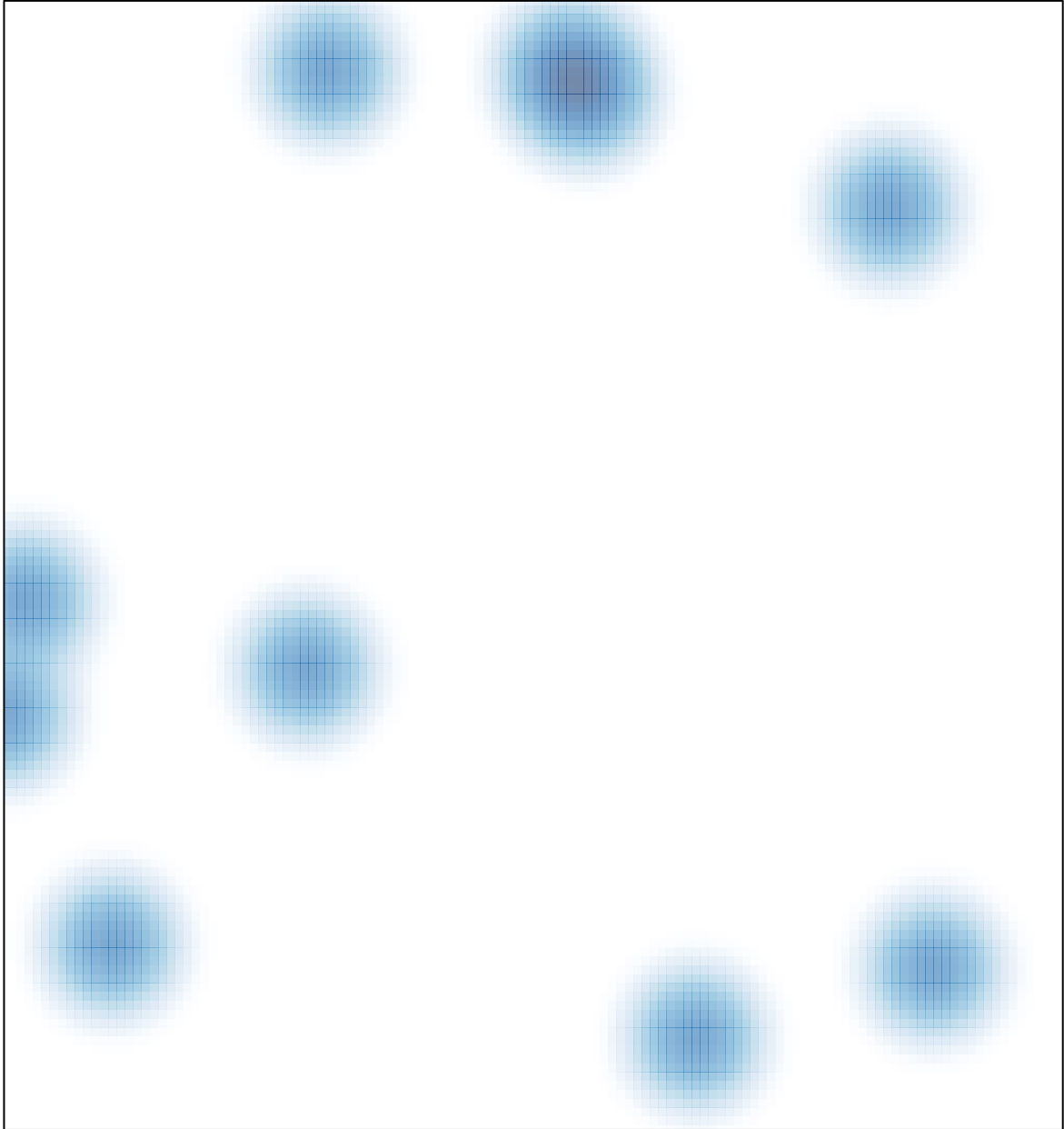


# chaperone-mediated protein folding



# features = 10  
chi-square p = 0.85

# chaperone-mediated protein folding



# features = 10 , max = 1