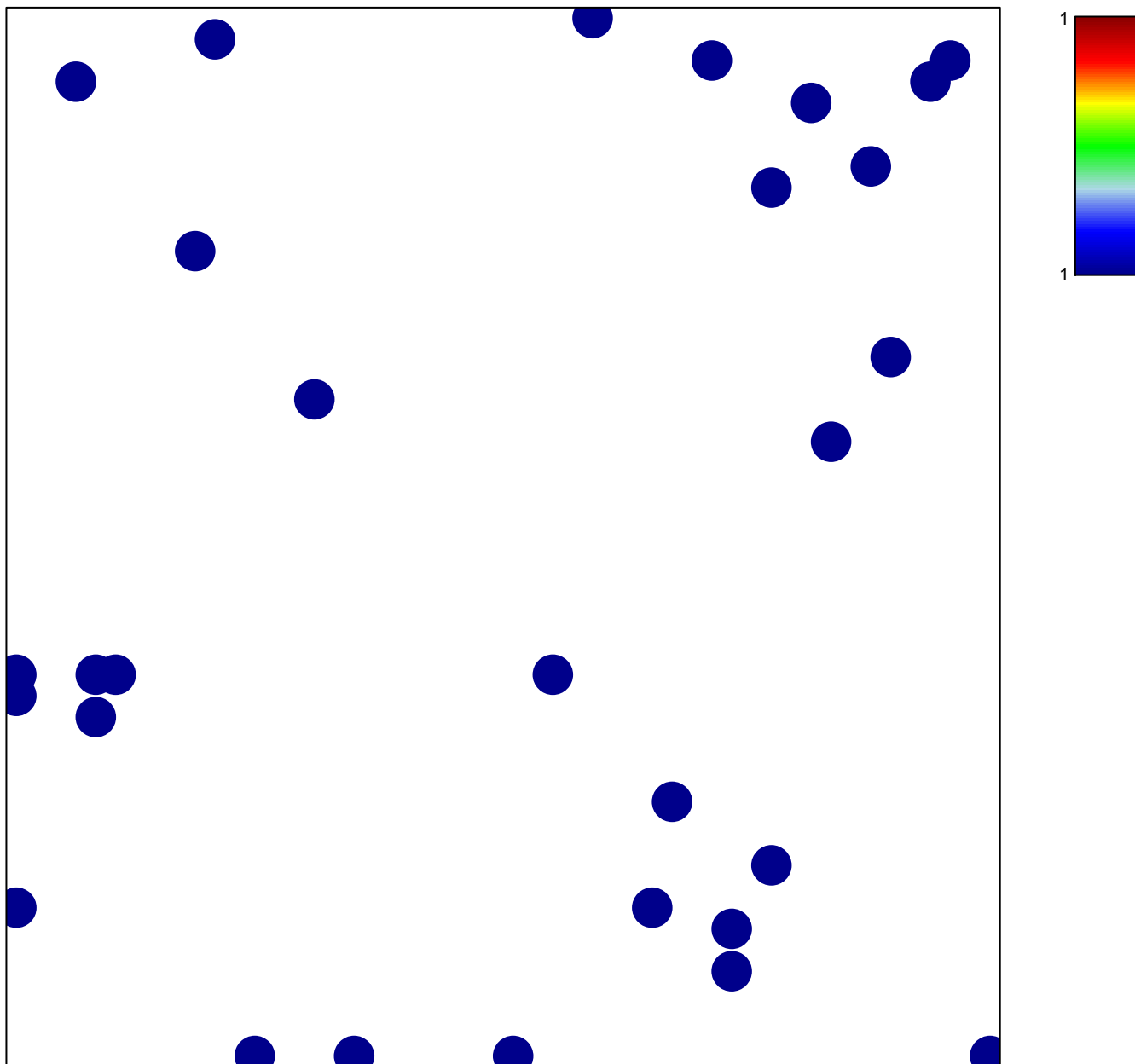
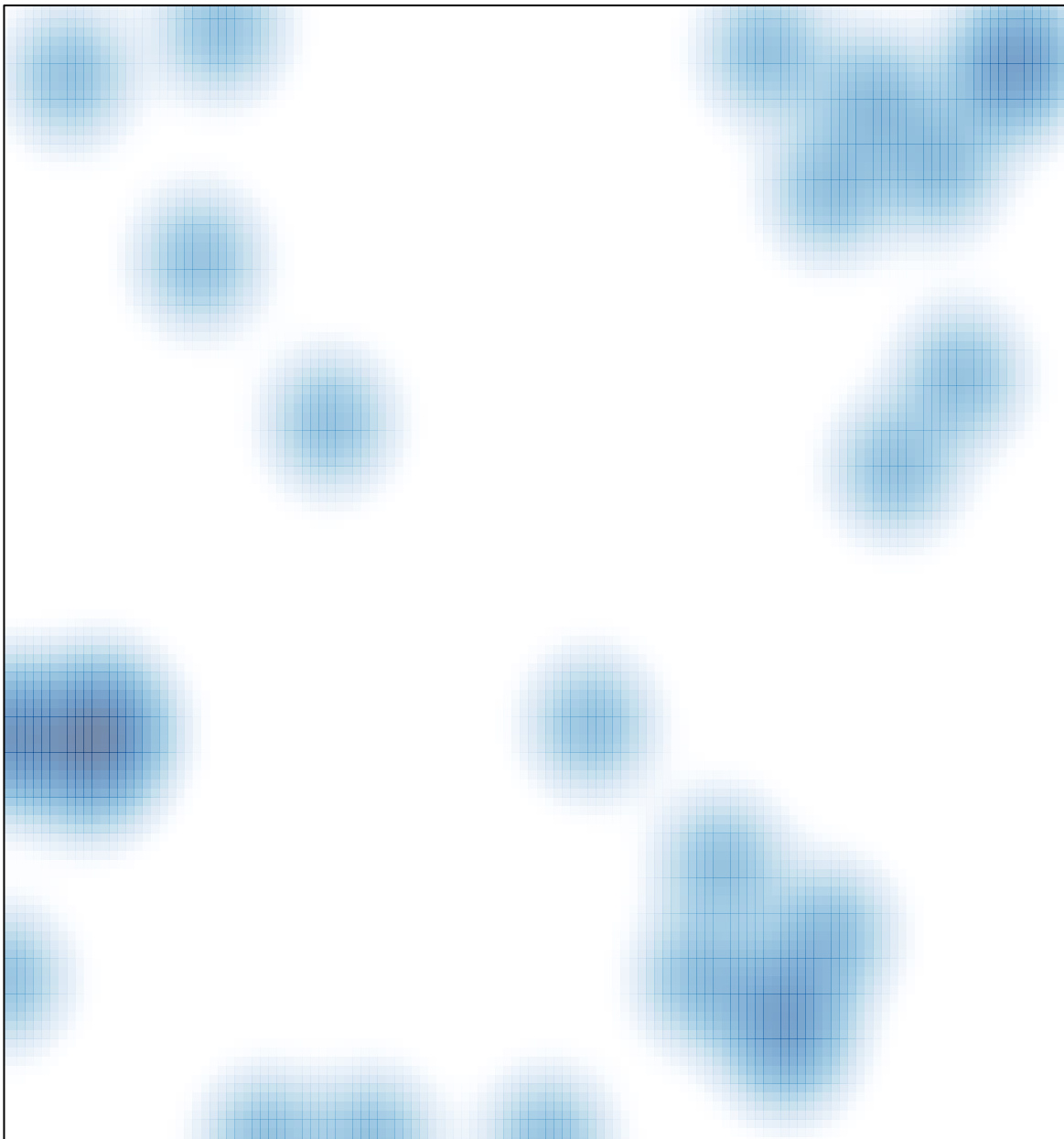


chaperone-mediated protein folding



features = 29
chi-square p = 0.83

chaperone-mediated protein folding



features = 29 , max = 1