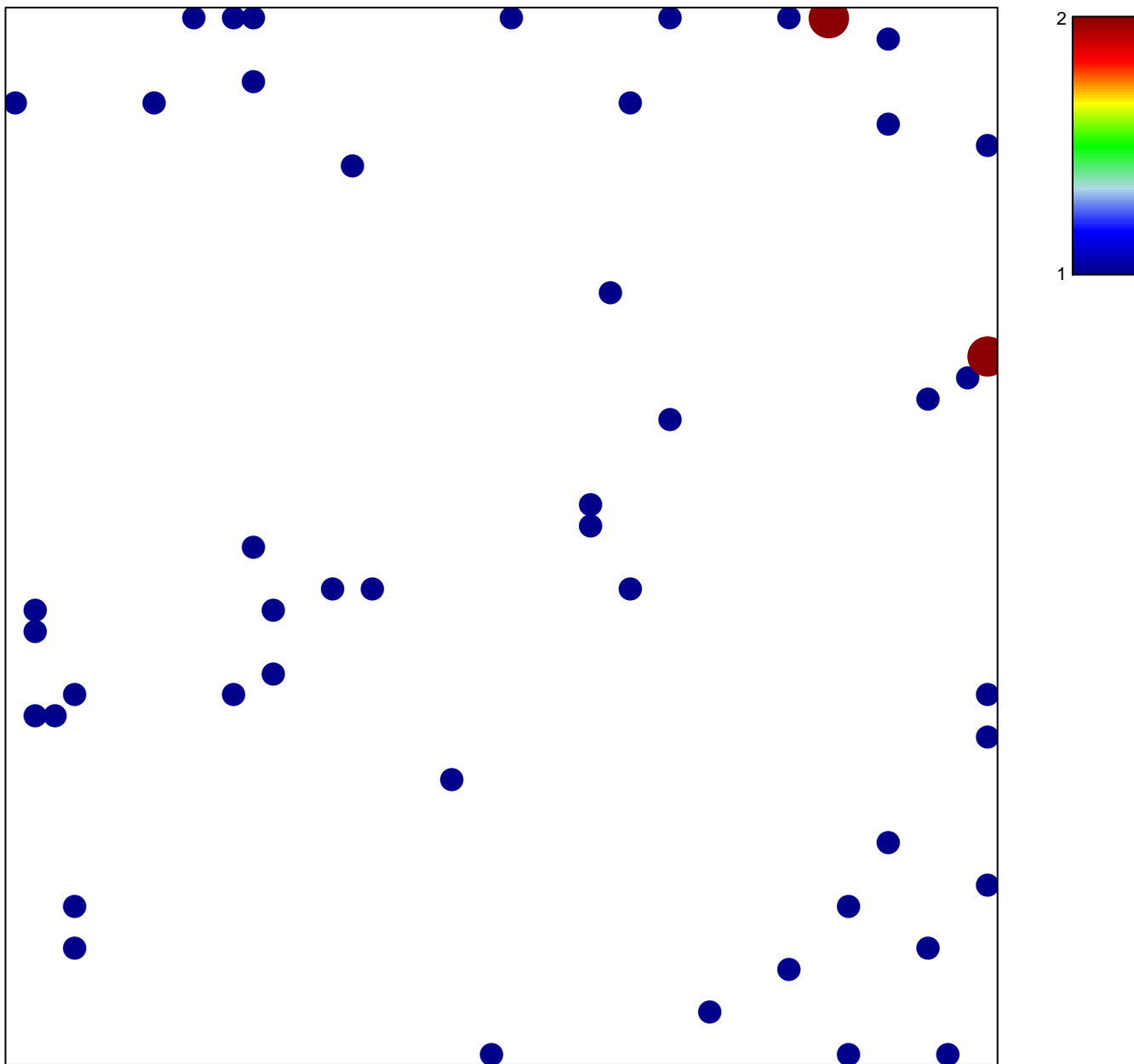
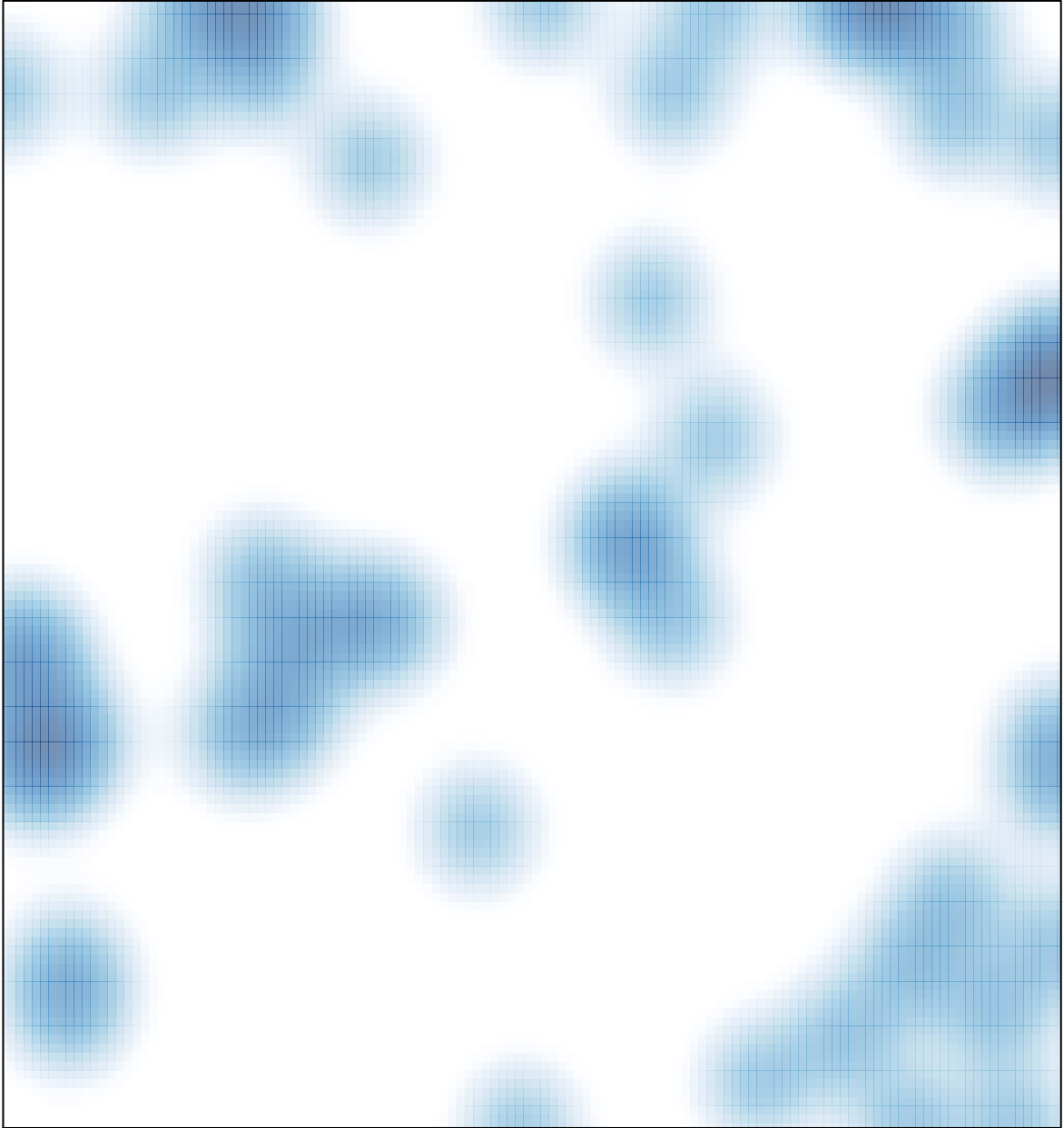


protein homotetramerization



features = 50
chi-square p = 0.8

protein homotetramerization



features = 50 , max = 2