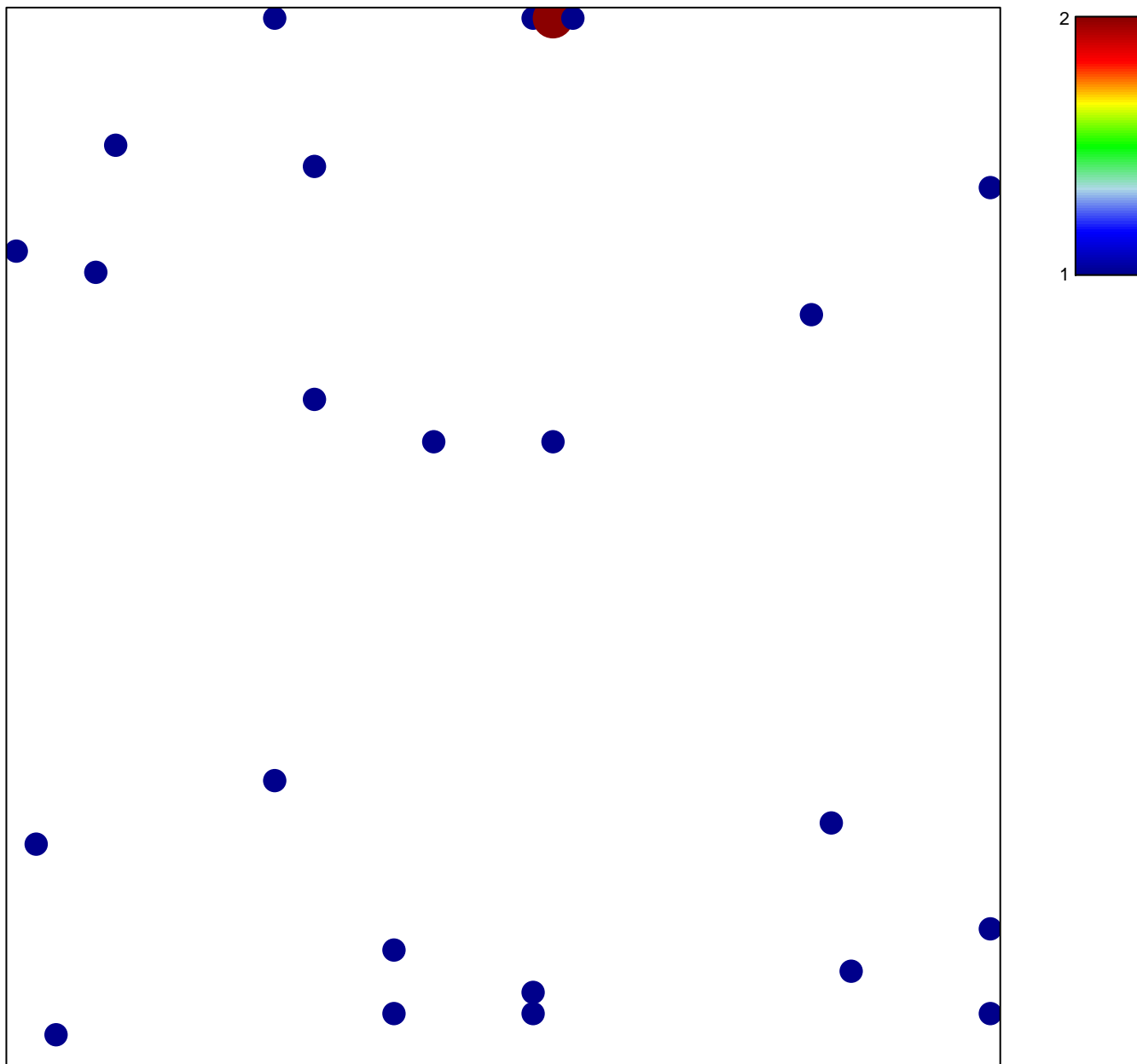


protein tetramerization



features = 25
chi-square p = 0.8

protein tetramerization



features = 25 , max = 2