protein heterotetramerization
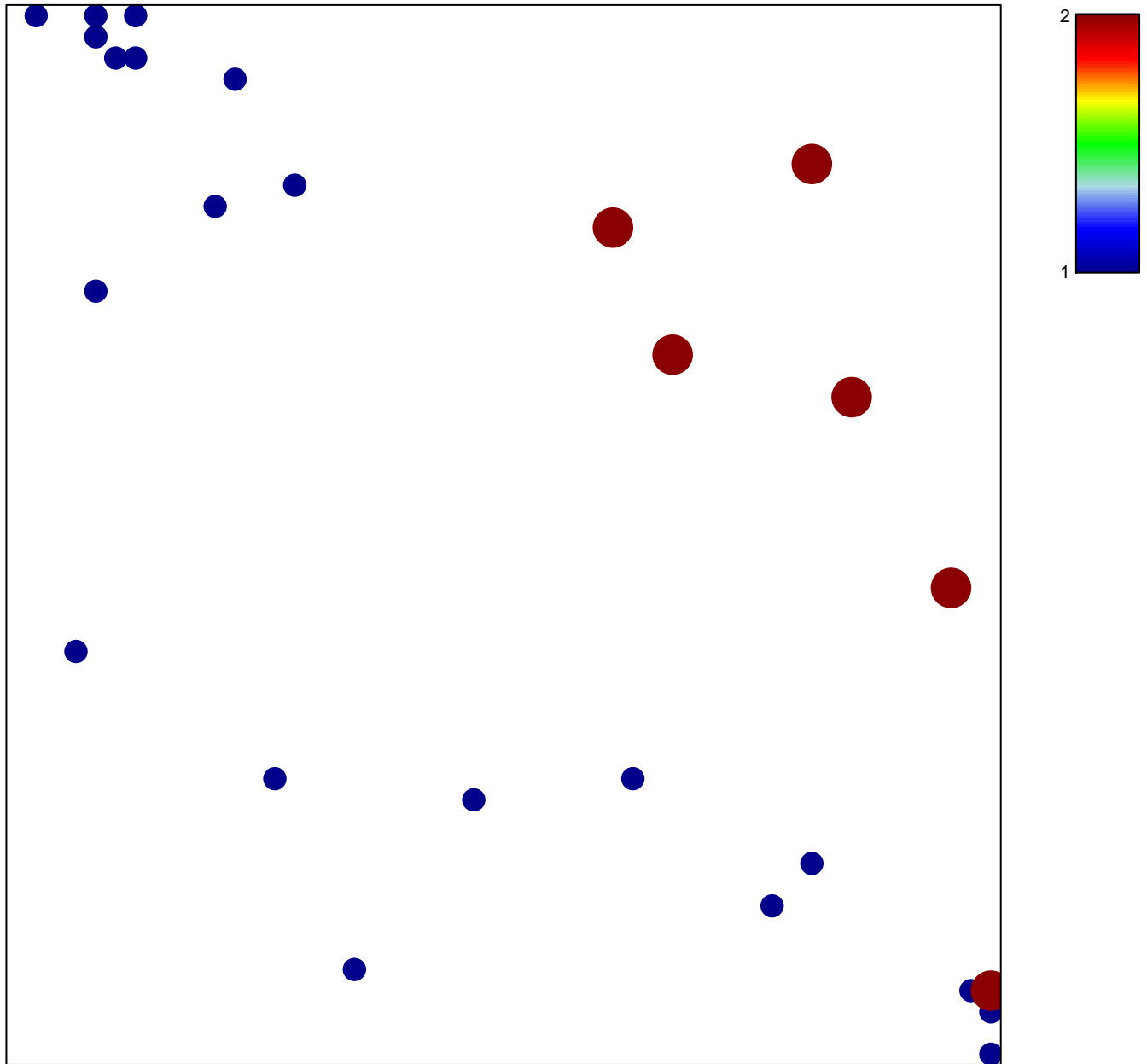
features = 32
chi−square p = 0.82

protein heterotetramerization

features = 32 , max = 2