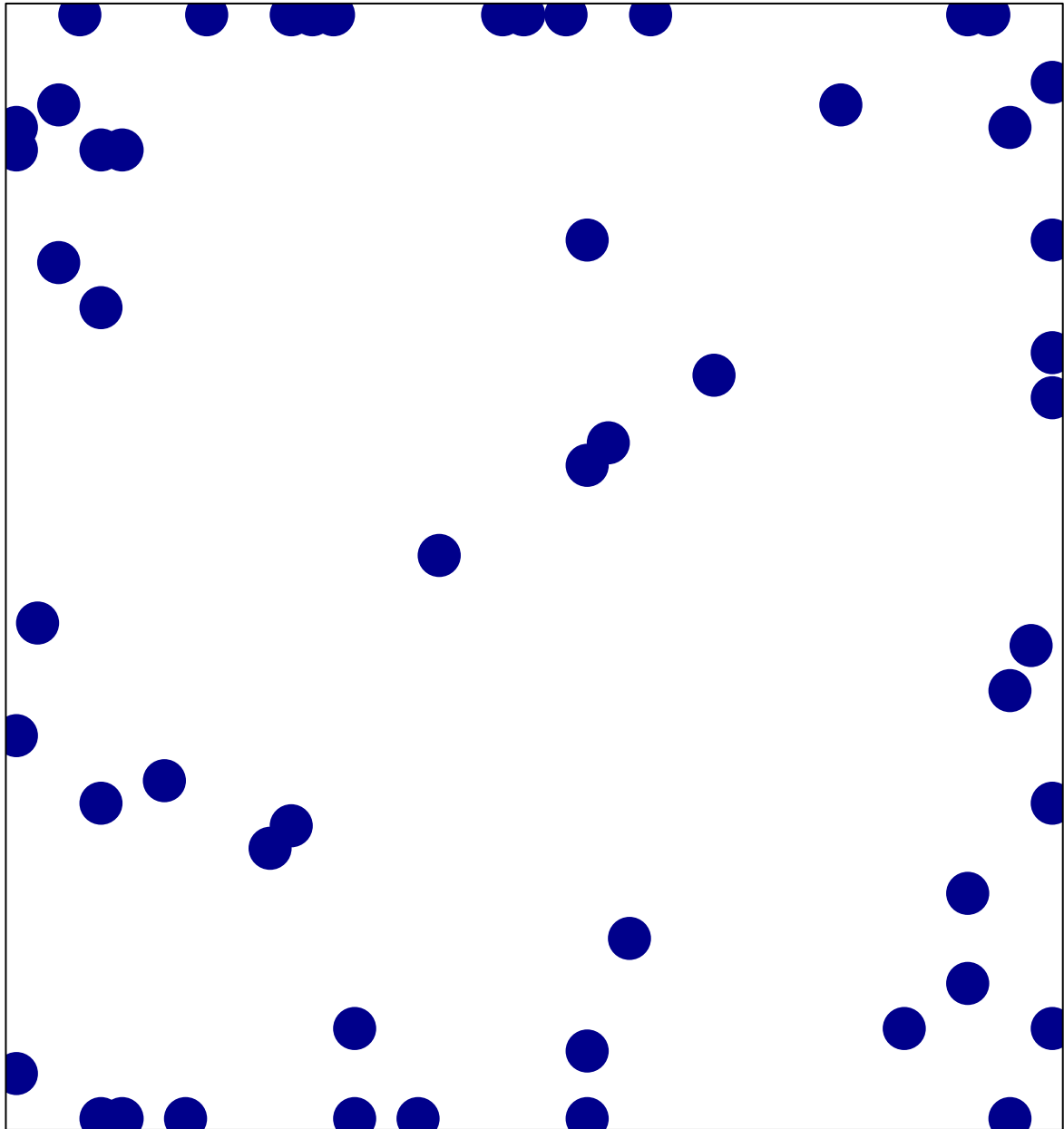
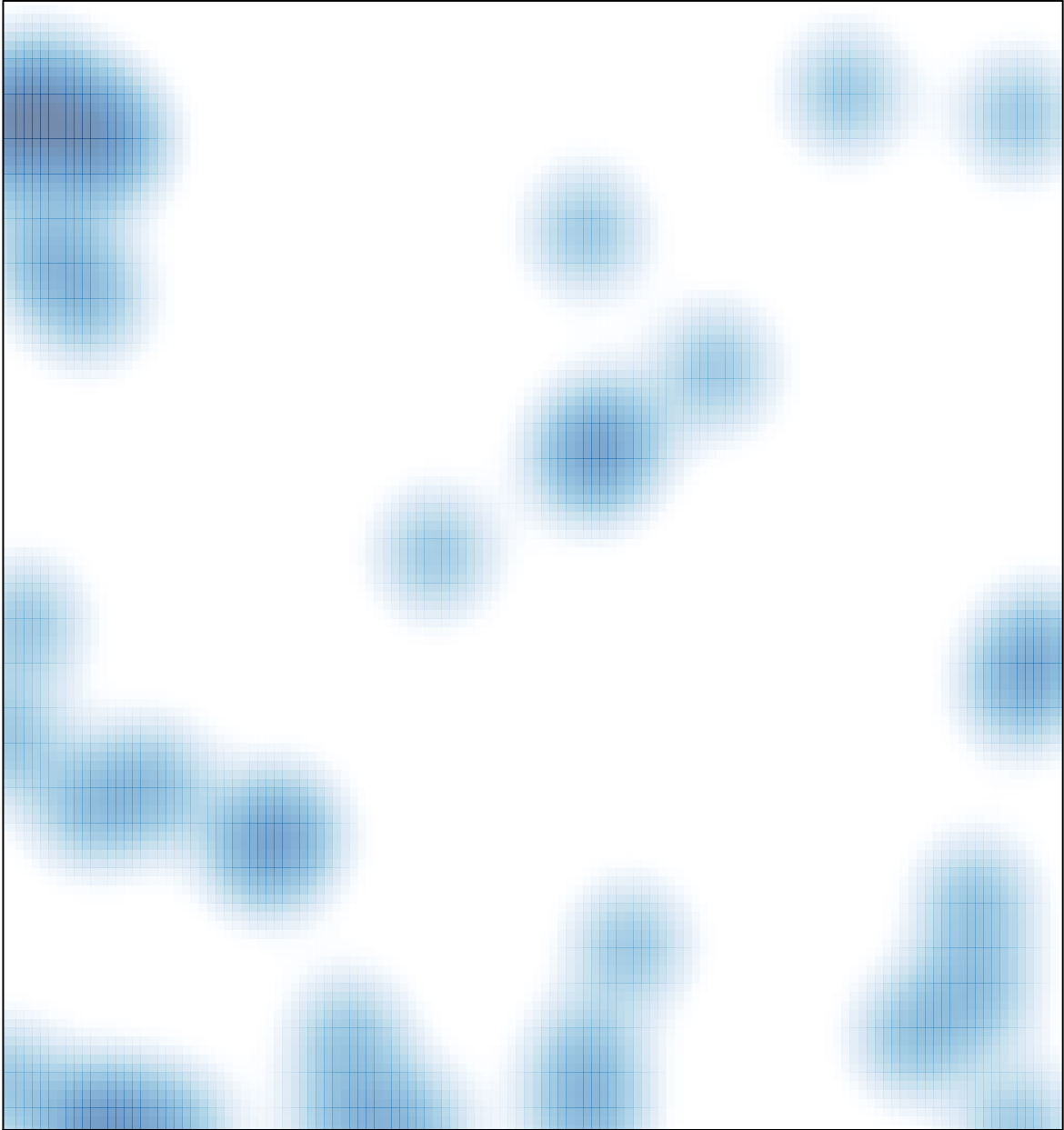


protein homotetramerization



features = 53
chi-square p = 0.76

protein homotetramerization



features = 53 , max = 1