ribonucleoprotein complex assembly
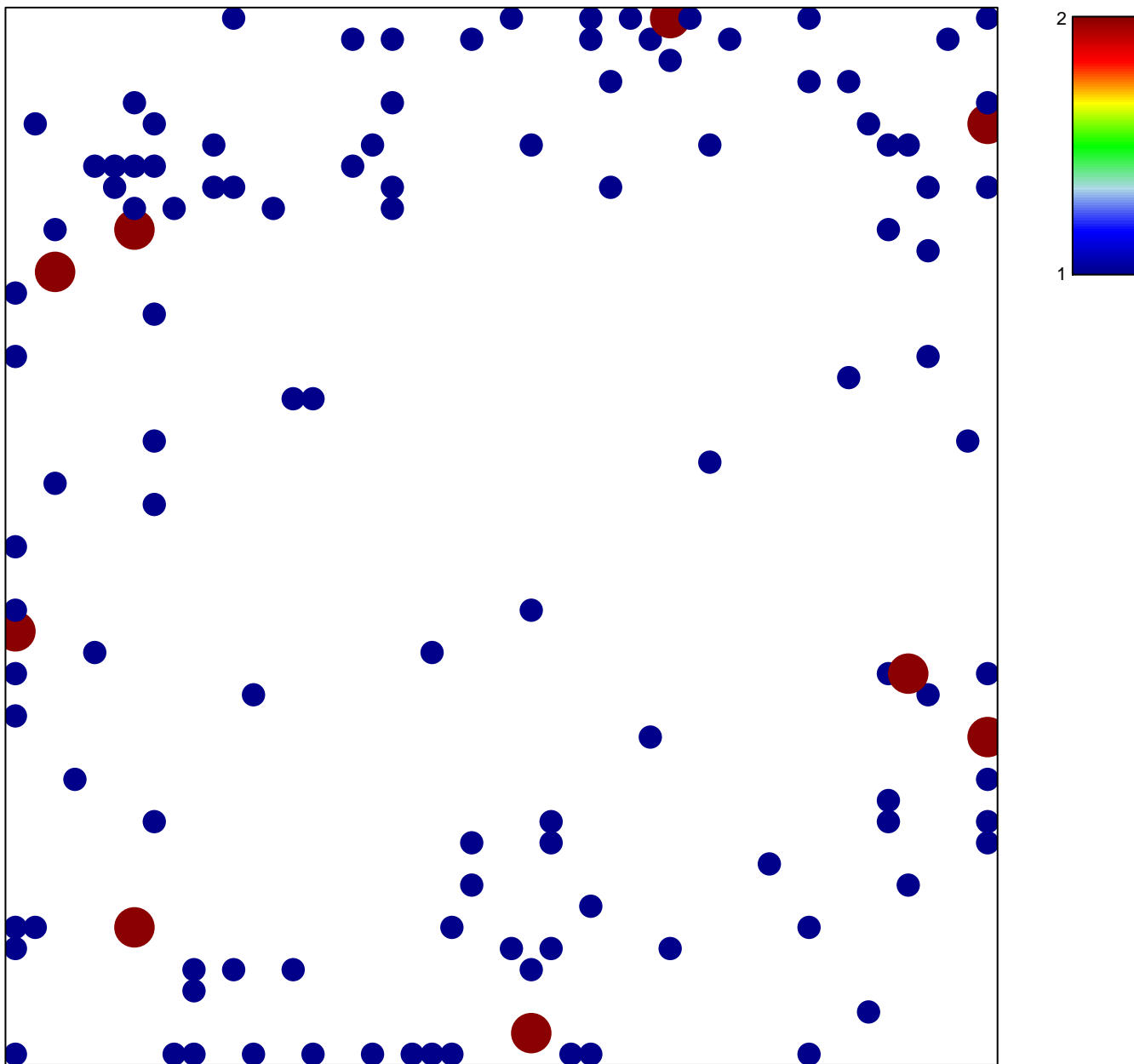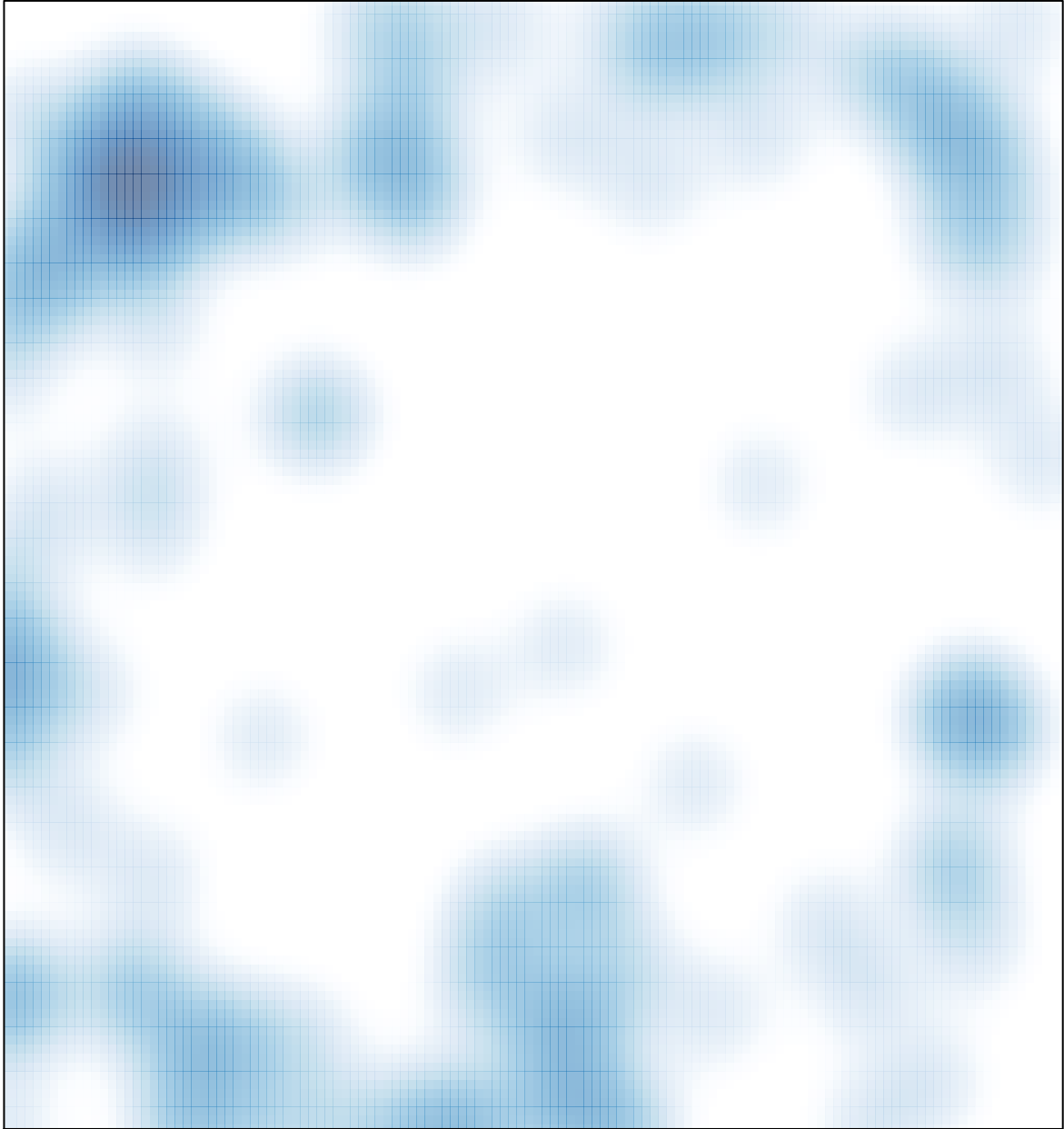

features = 131
chi−square p = 0.77

ribonucleoprotein complex assembly

features = 131 , max = 2