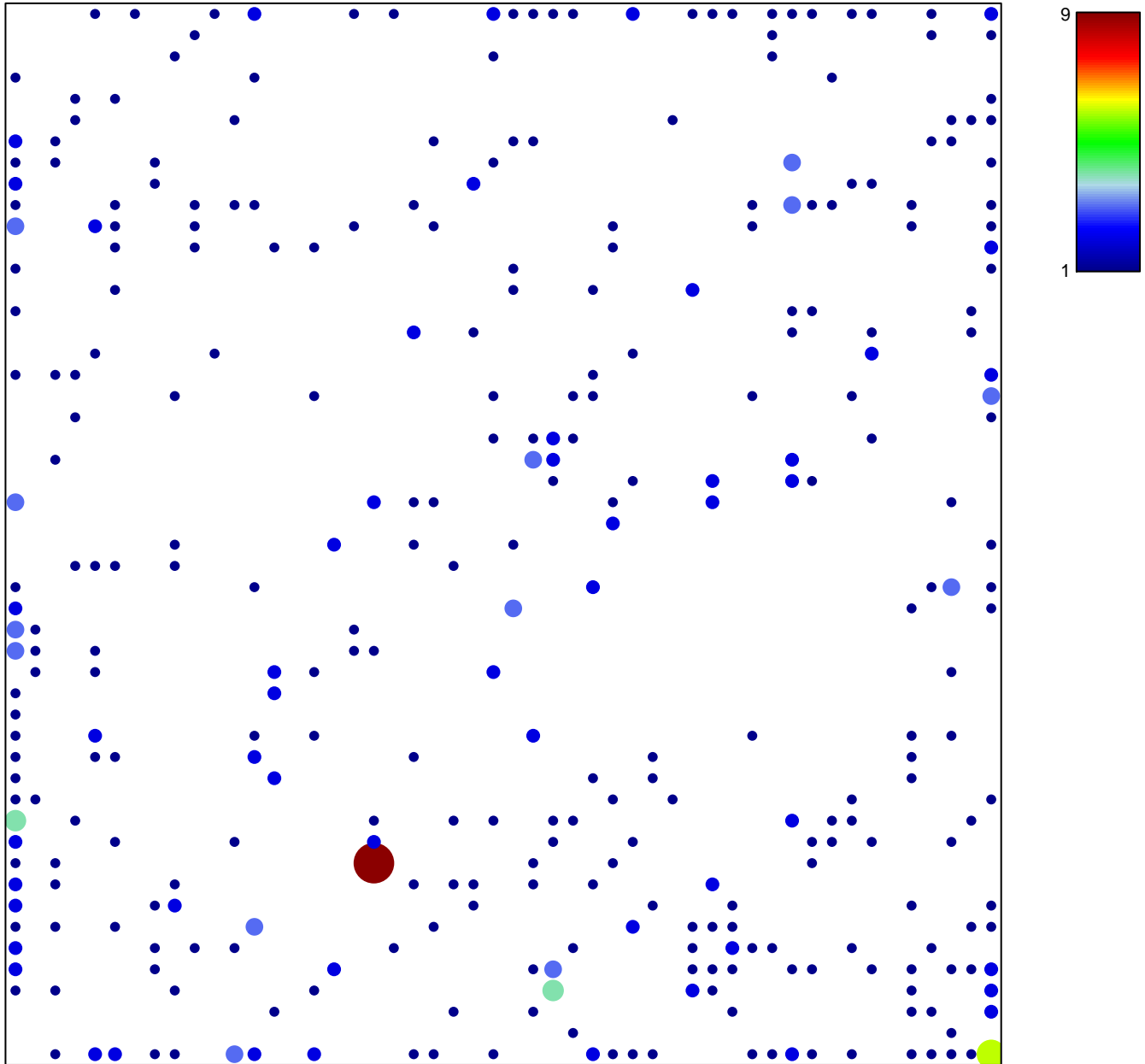
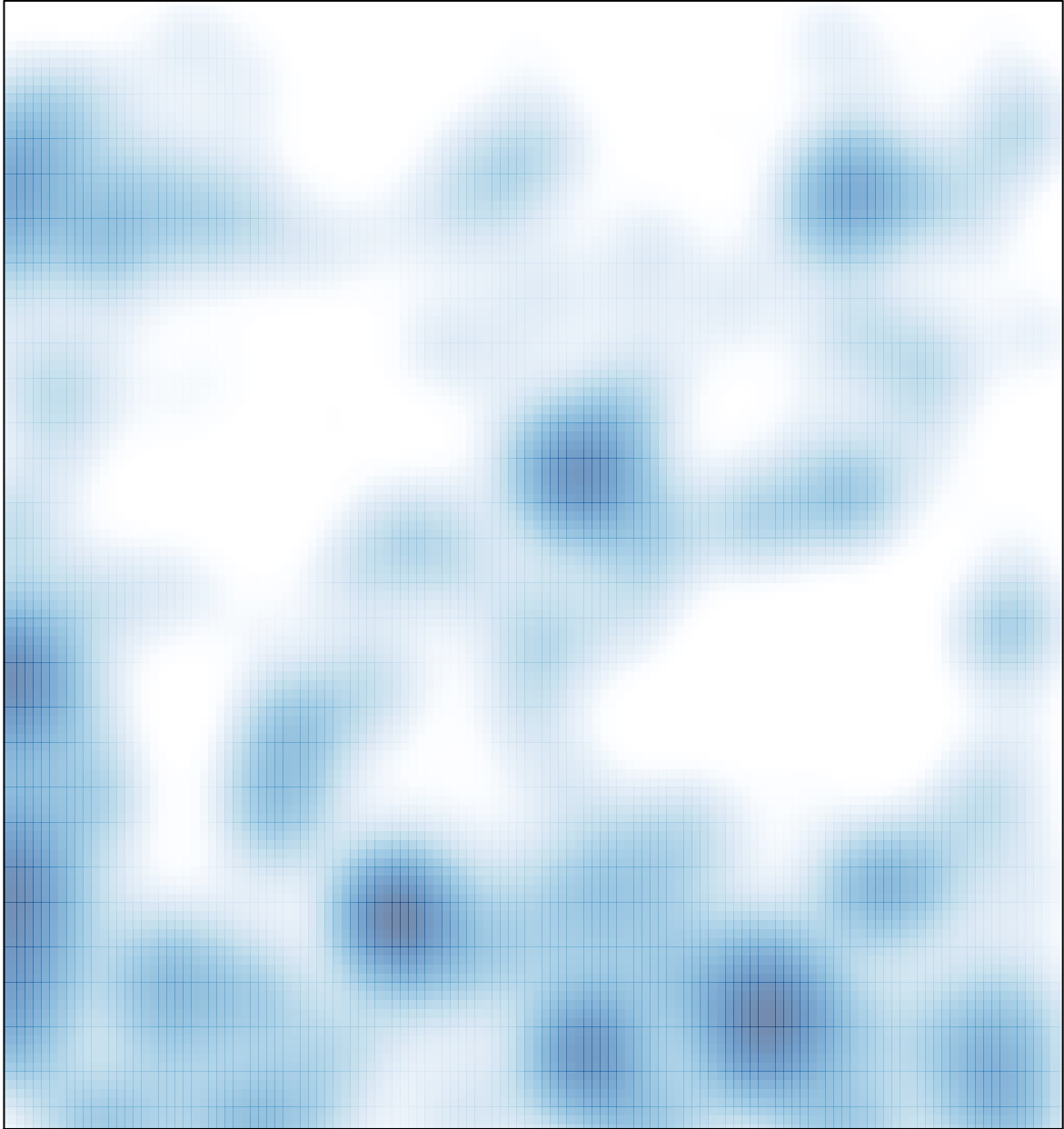


lipid binding



features = 430
chi-square p = 0.63

lipid binding



features = 430 , max = 9