post−translational protein modification
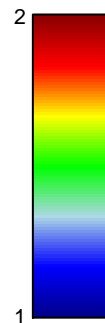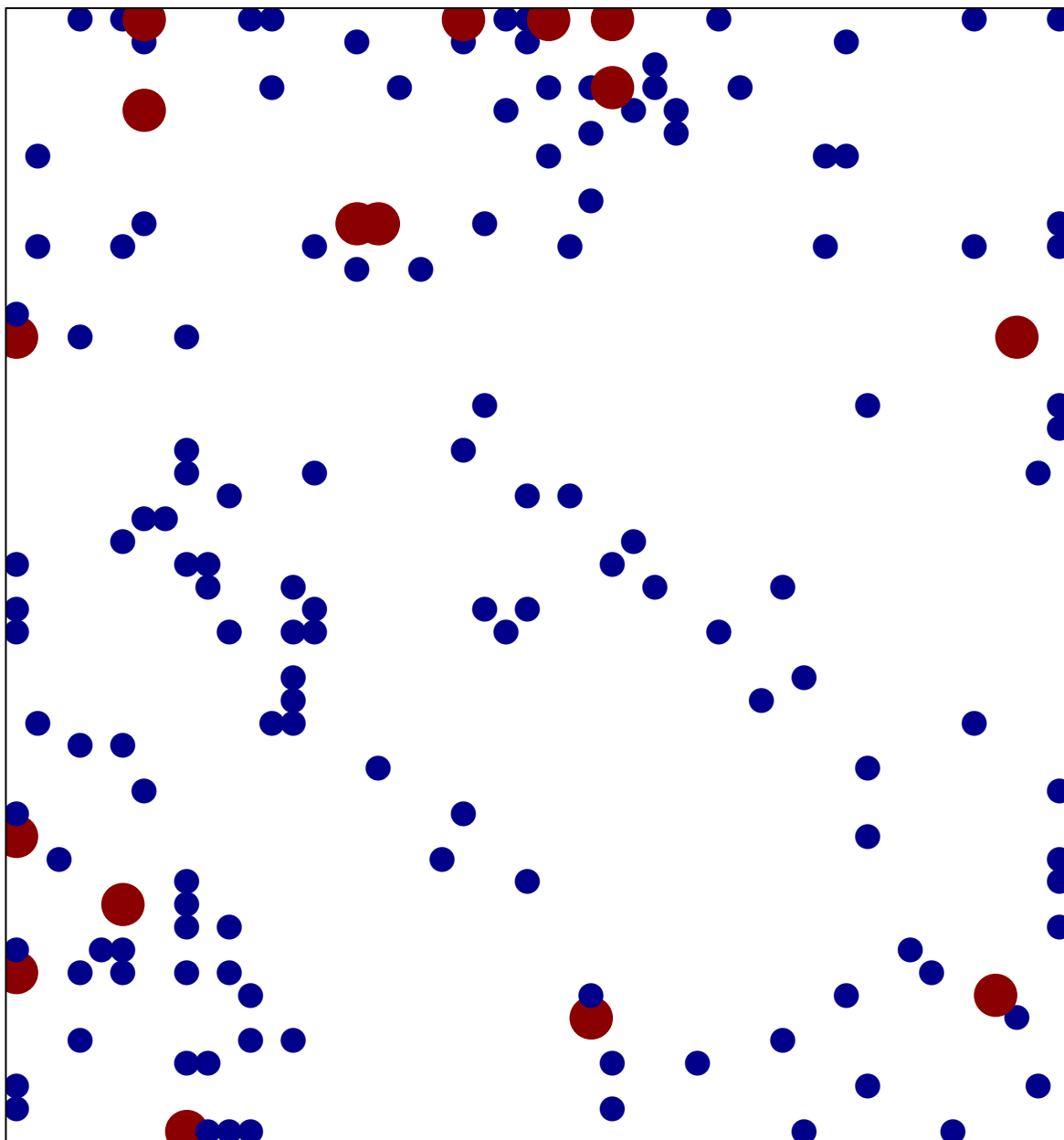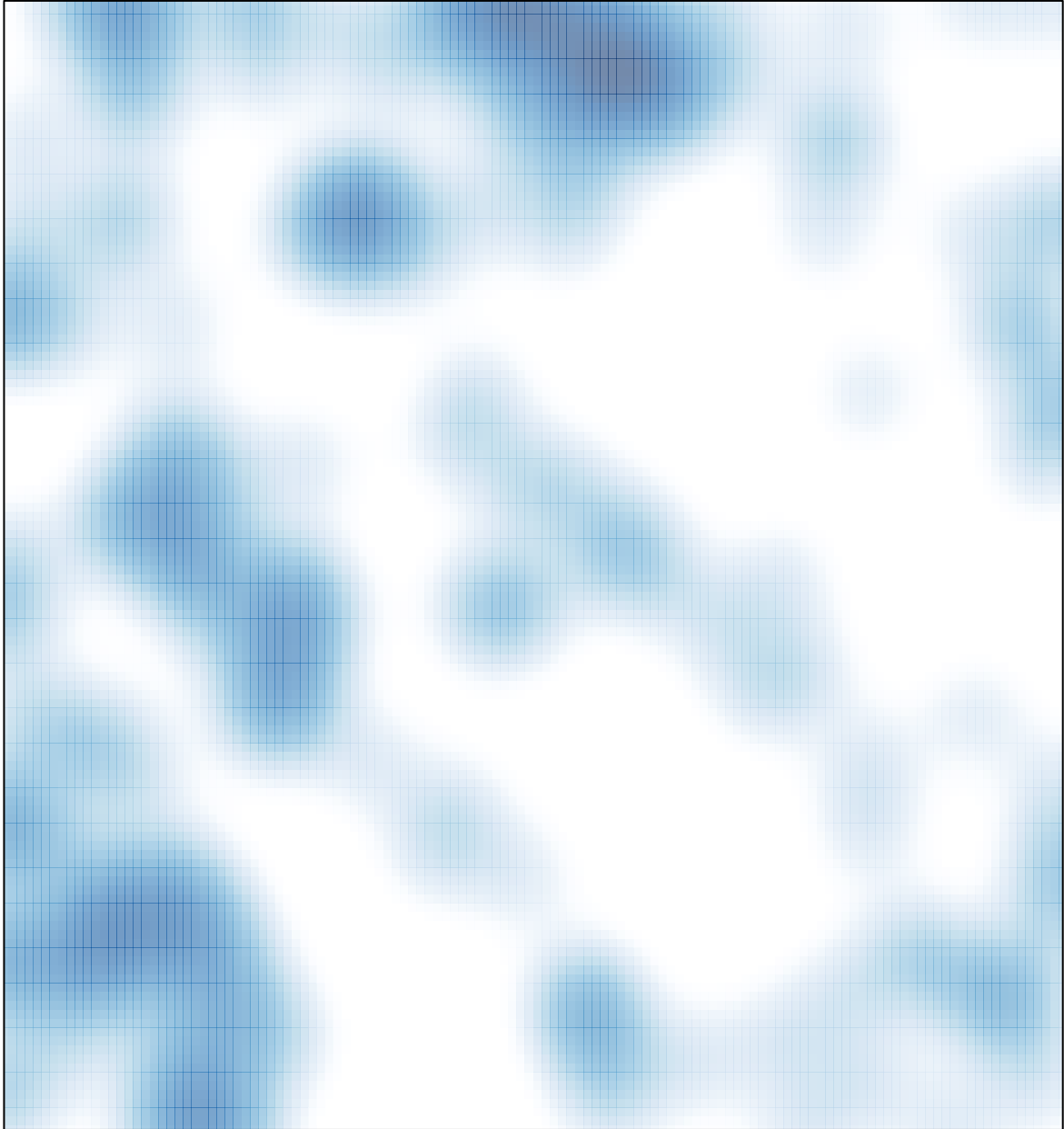

features = 170
chi−square p = 0.76

post–translational protein modification

features = 170 , max = 2