post−translational protein modification
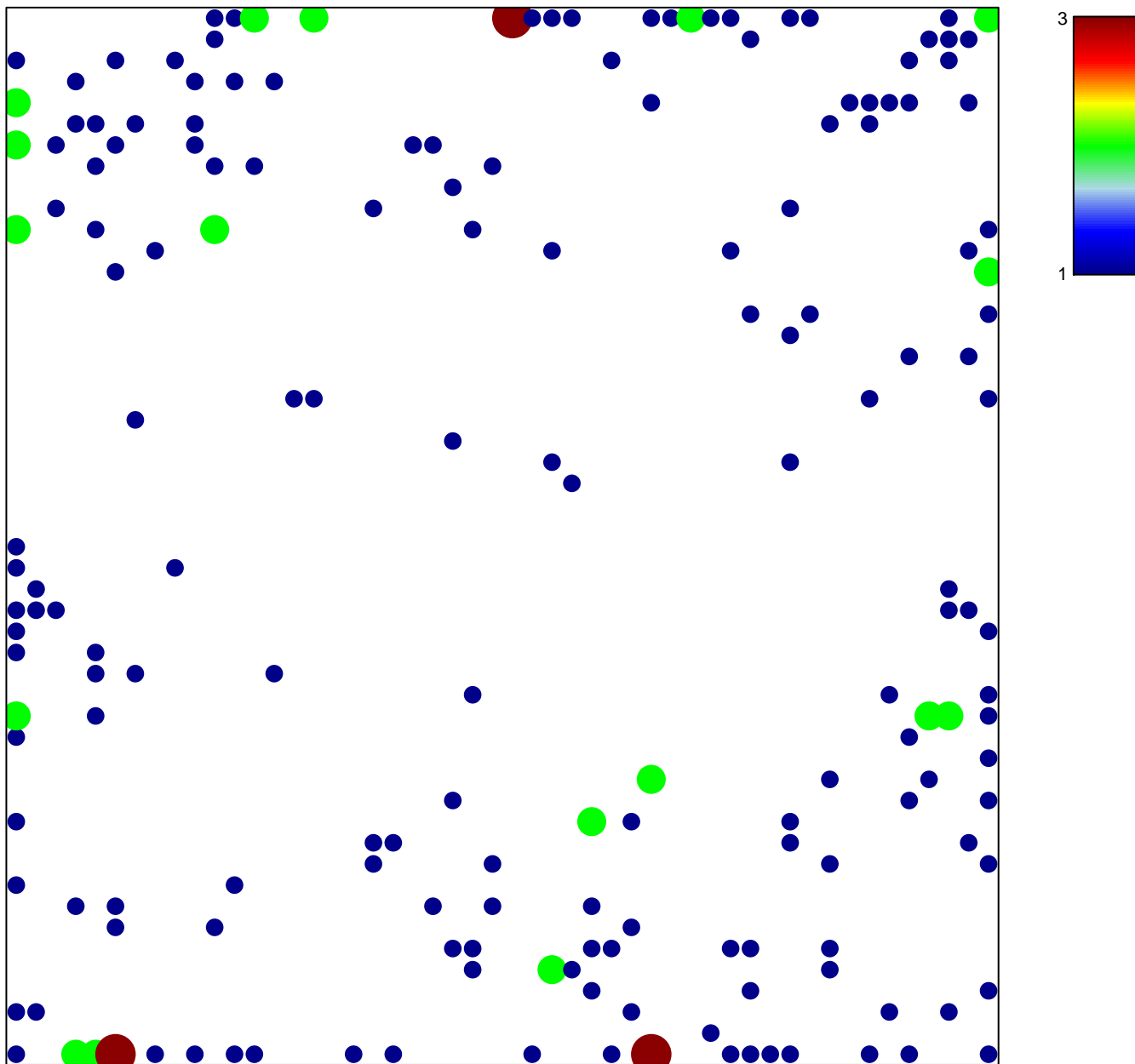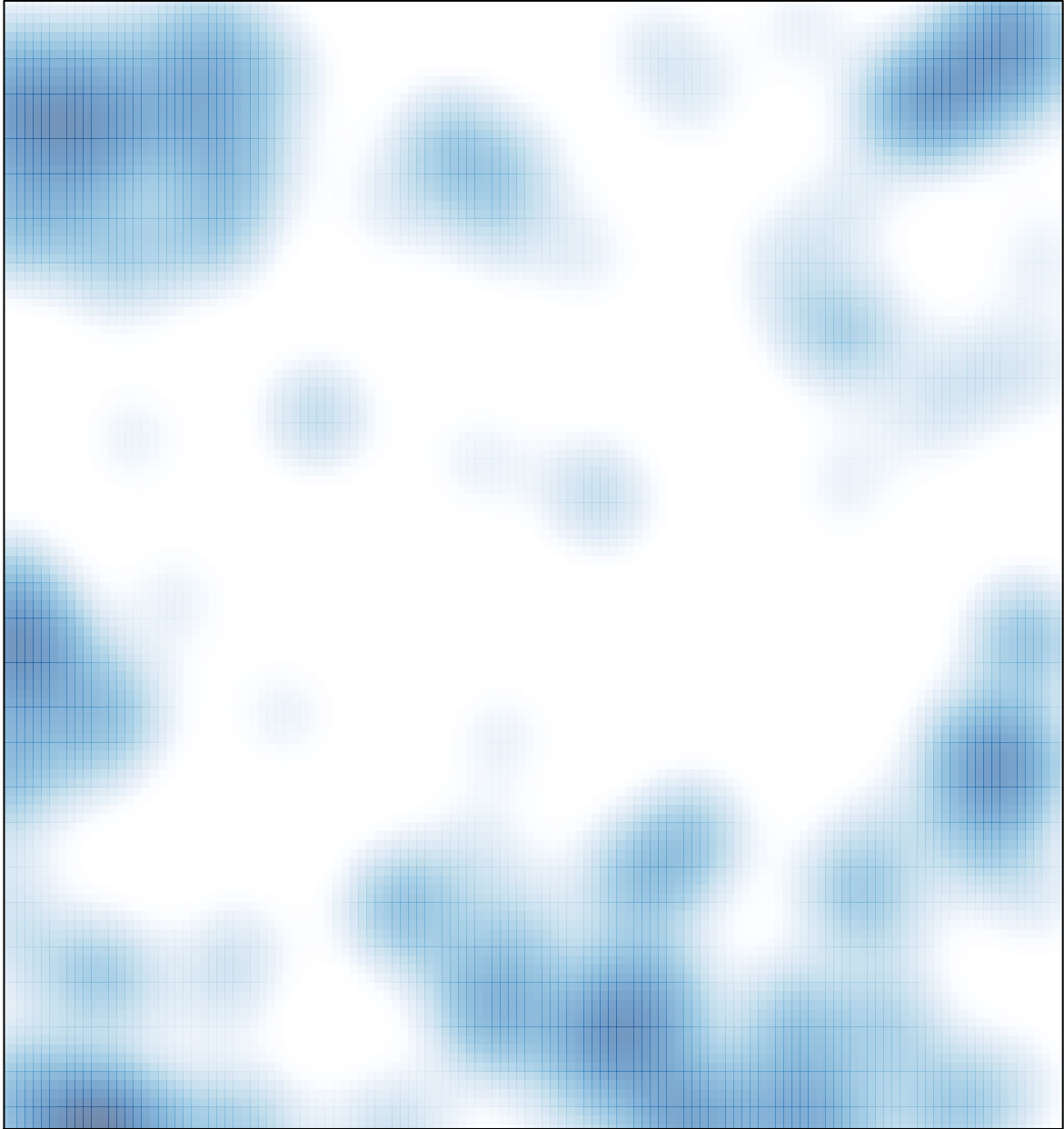
features = 204

chi−square p = 0.69

post–translational protein modification

features = 204 , max = 3