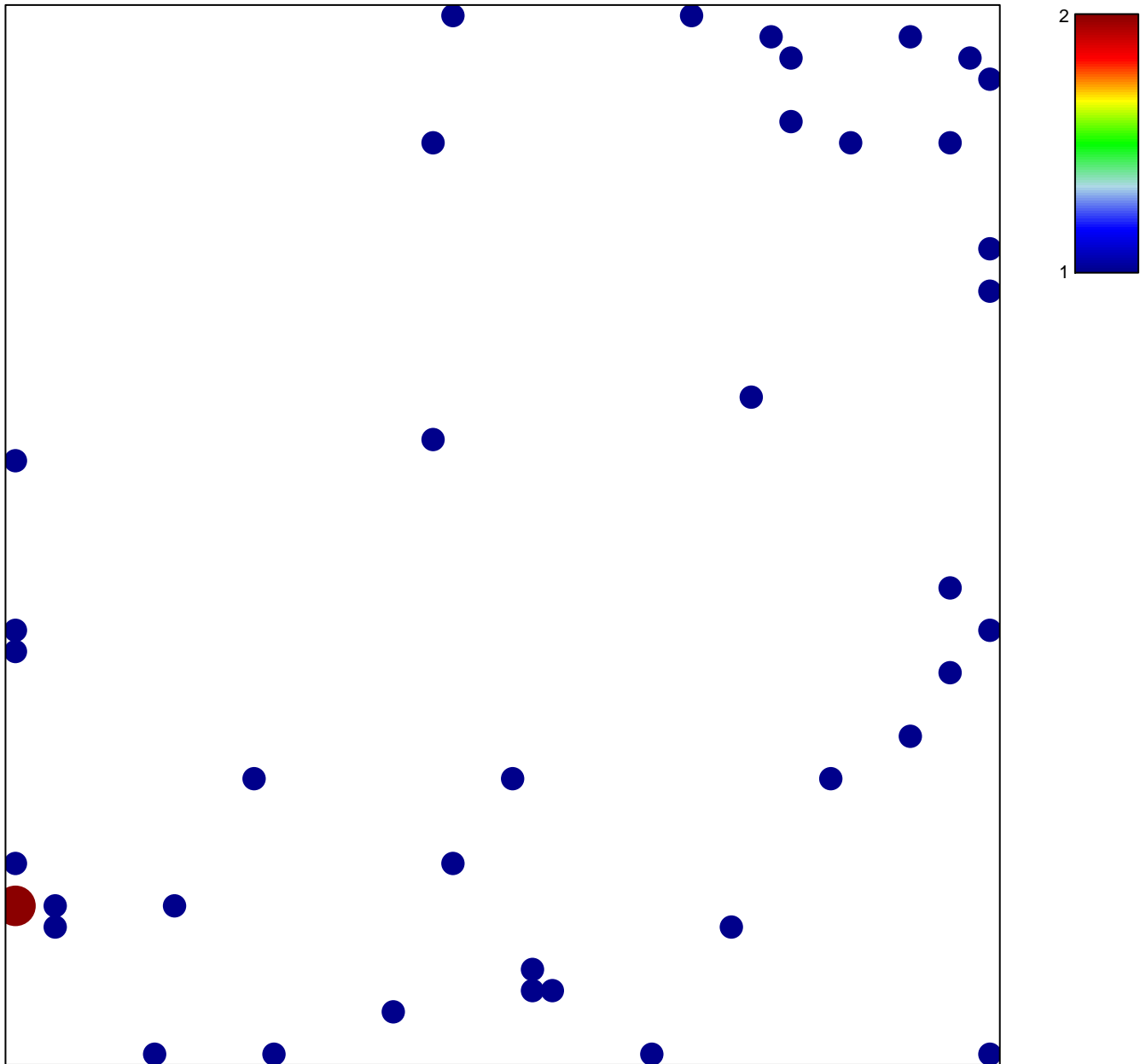
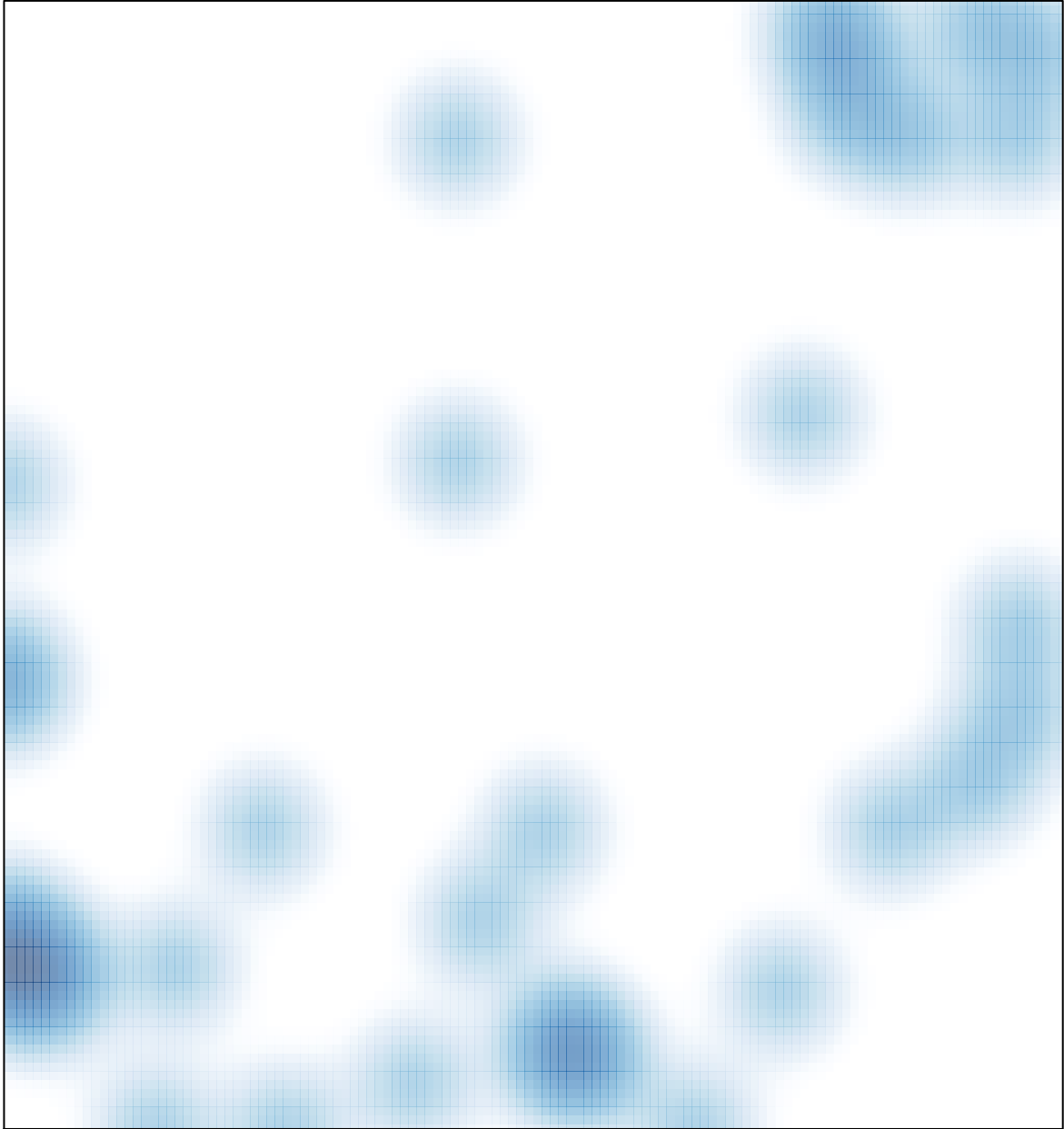


ATPase binding



features = 41
chi-square p = 0.8

ATPase binding



features = 41 , max = 2