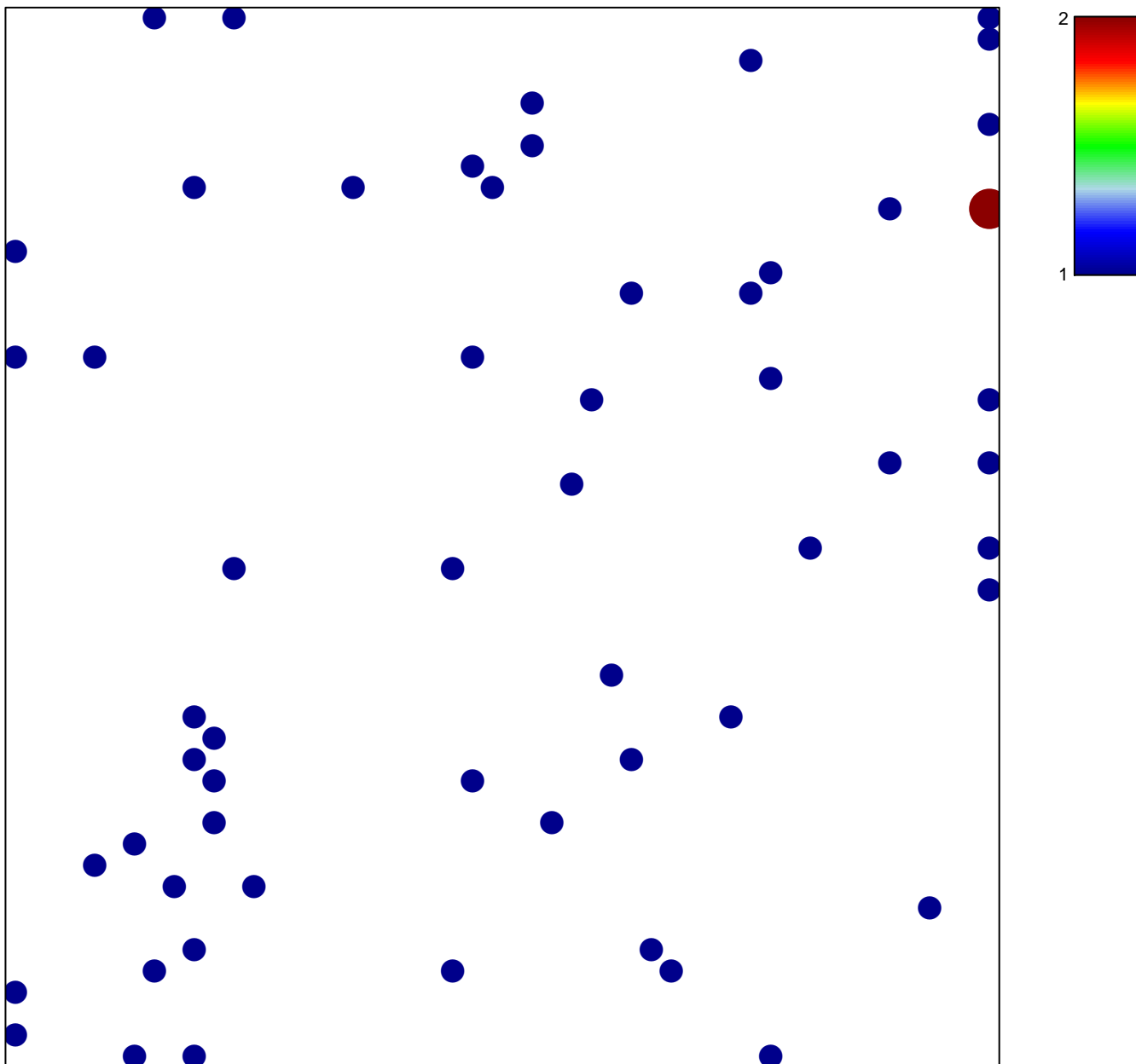
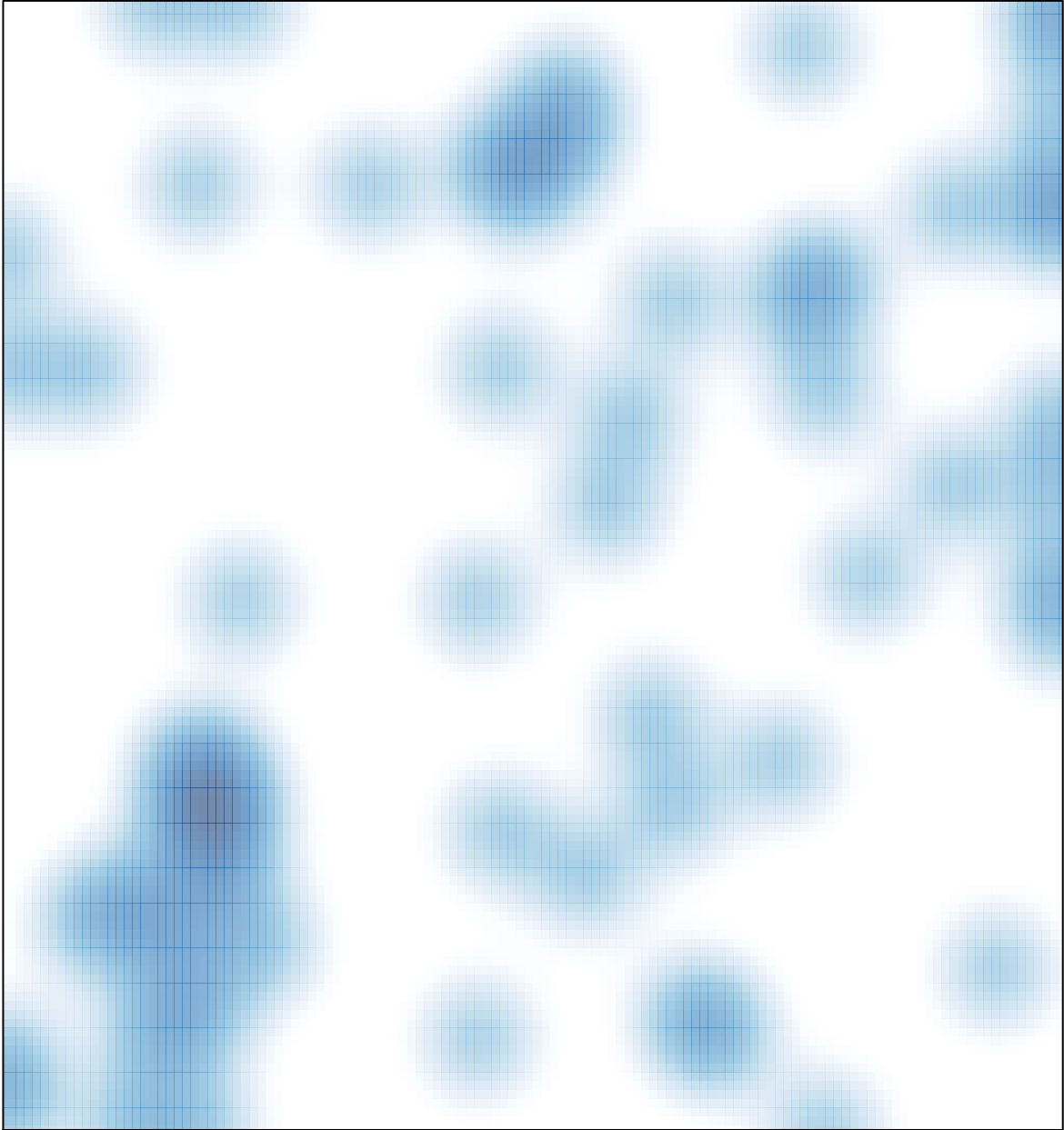


protein localization



features = 58
chi-square p = 0.8

protein localization



features = 58 , max = 2