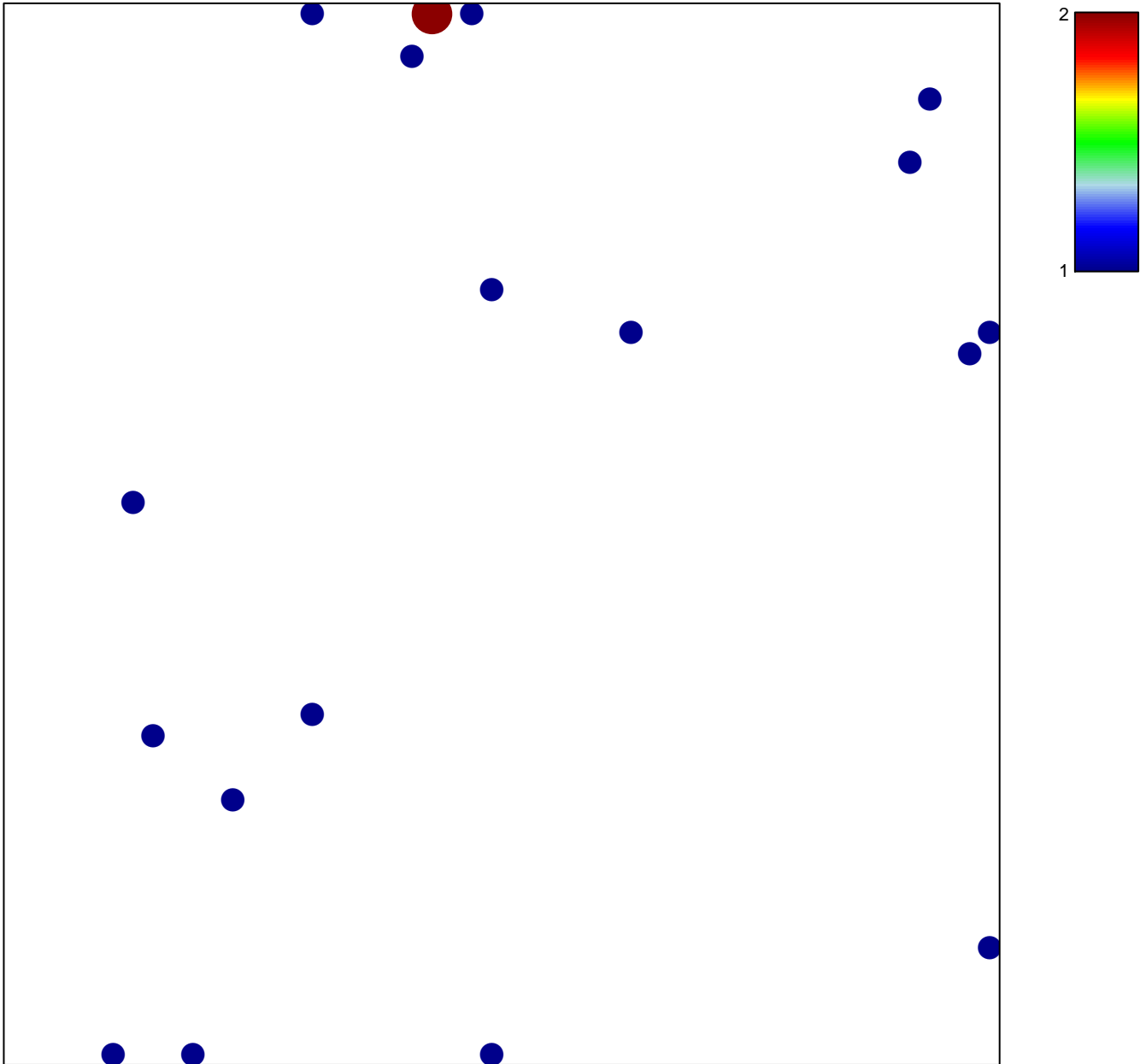
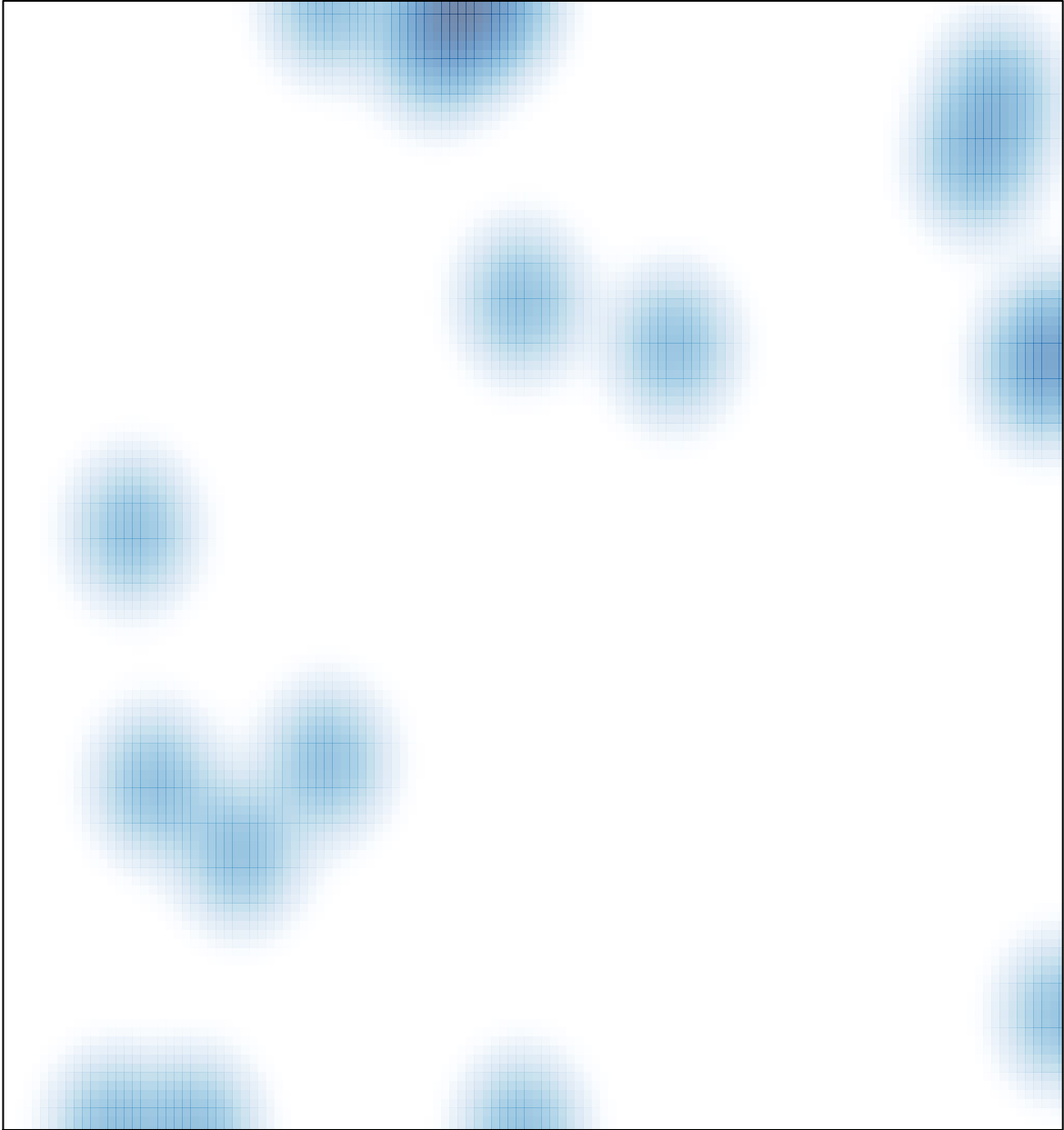


R-SMAD binding



features = 19
chi-square p = 0.82

R-SMAD binding



features = 19 , max = 2