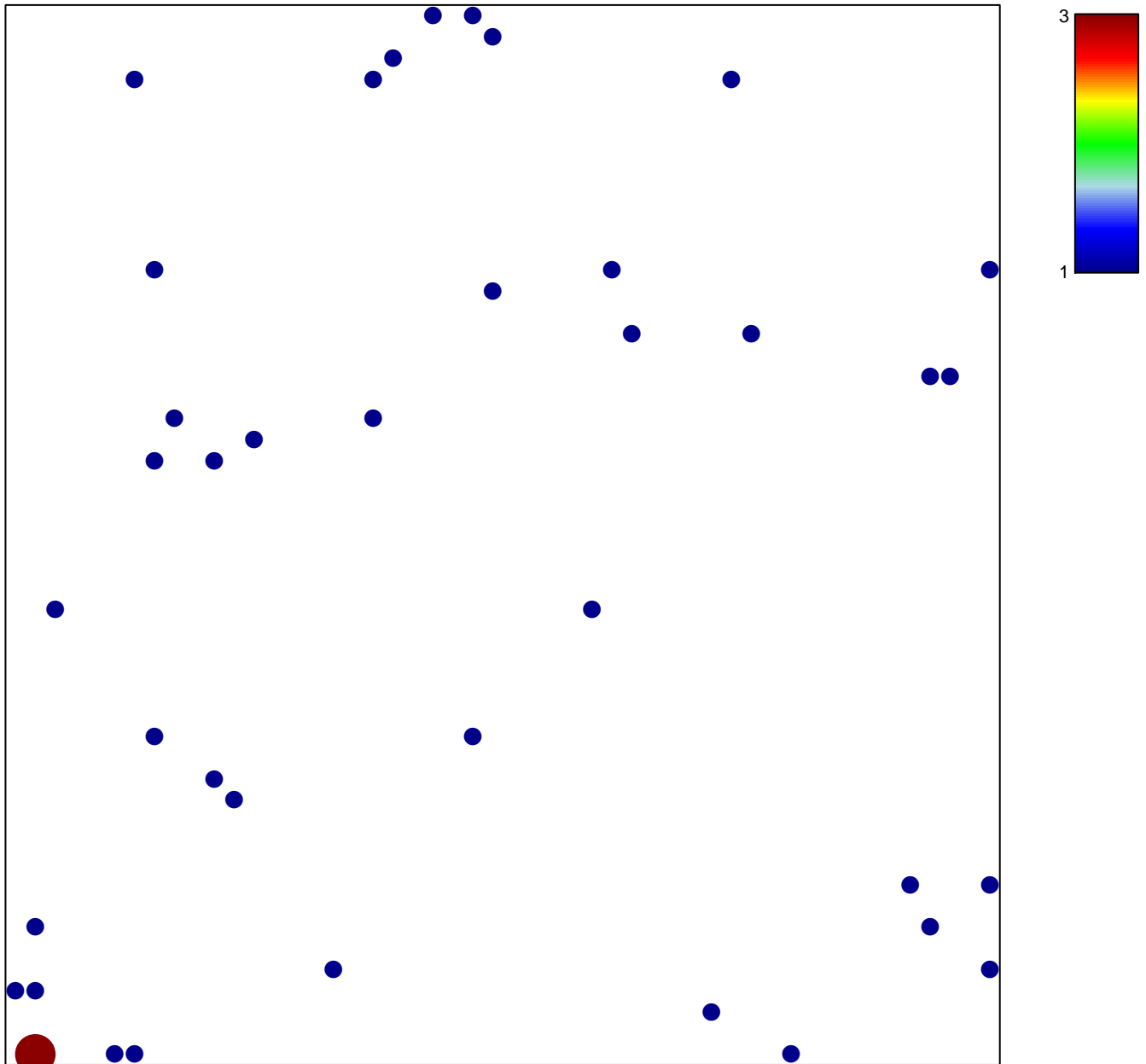
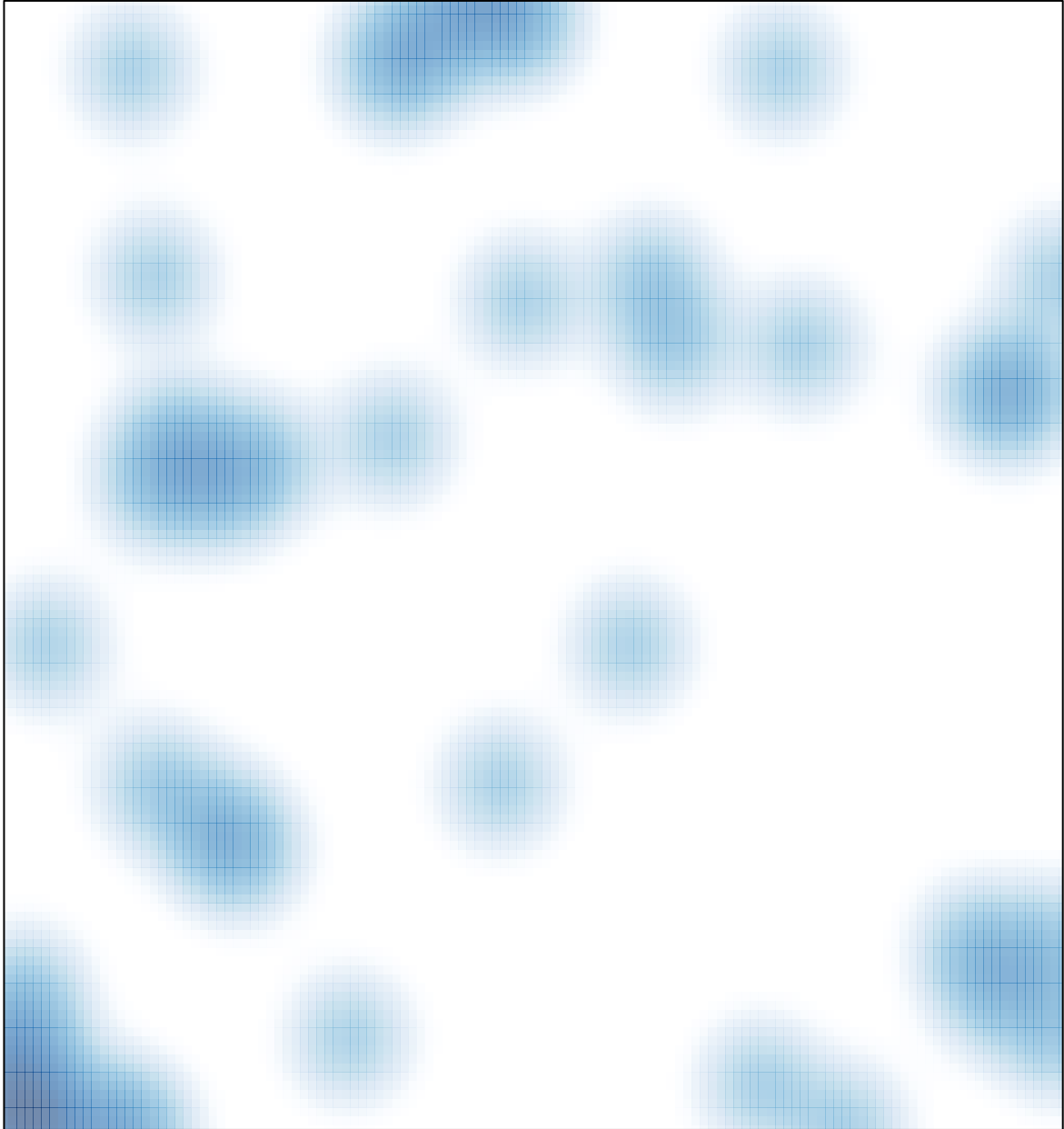


SMAD binding



features = 41
chi-square p = 0.81

SMAD binding



features = 41 , max = 3