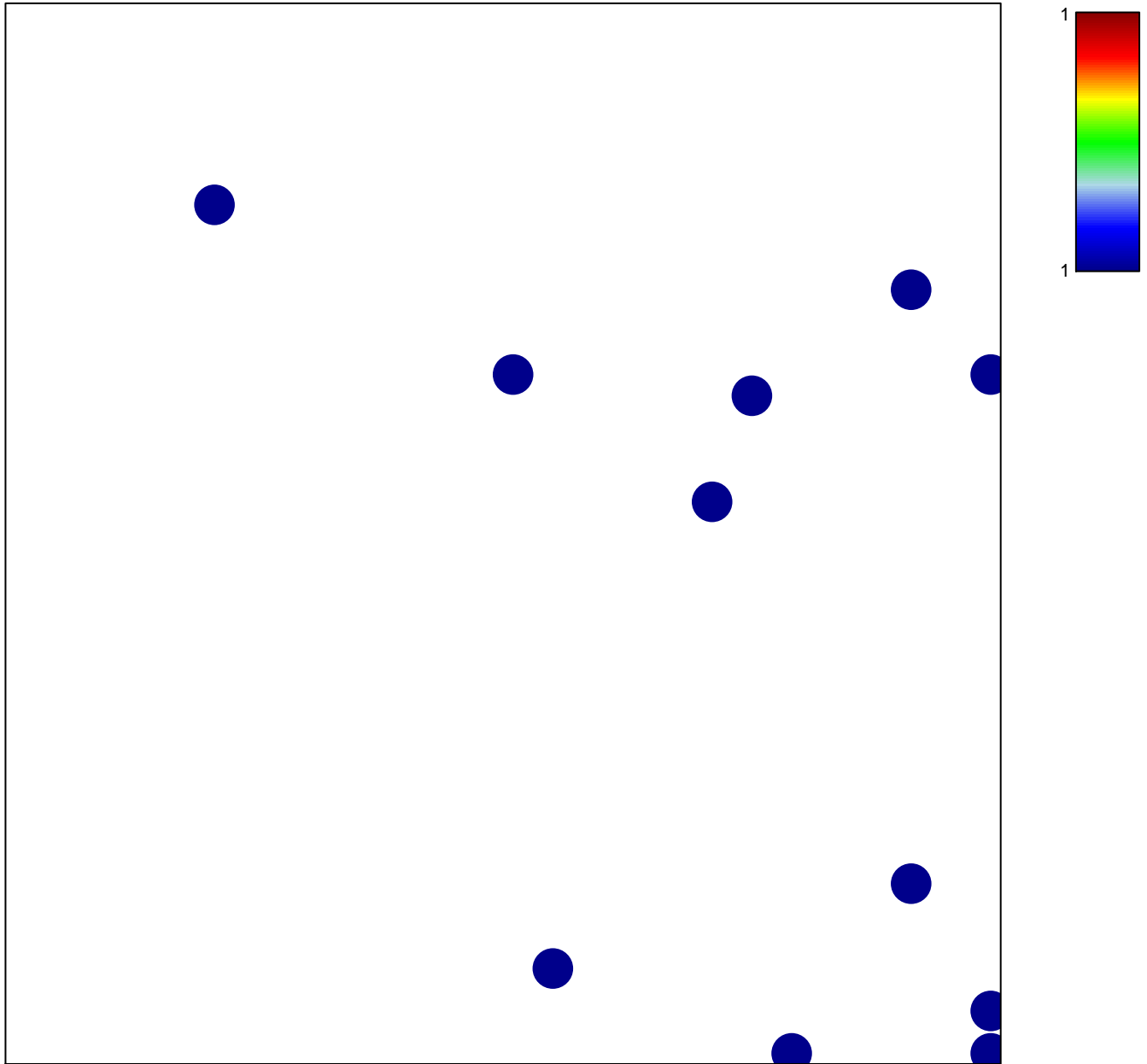
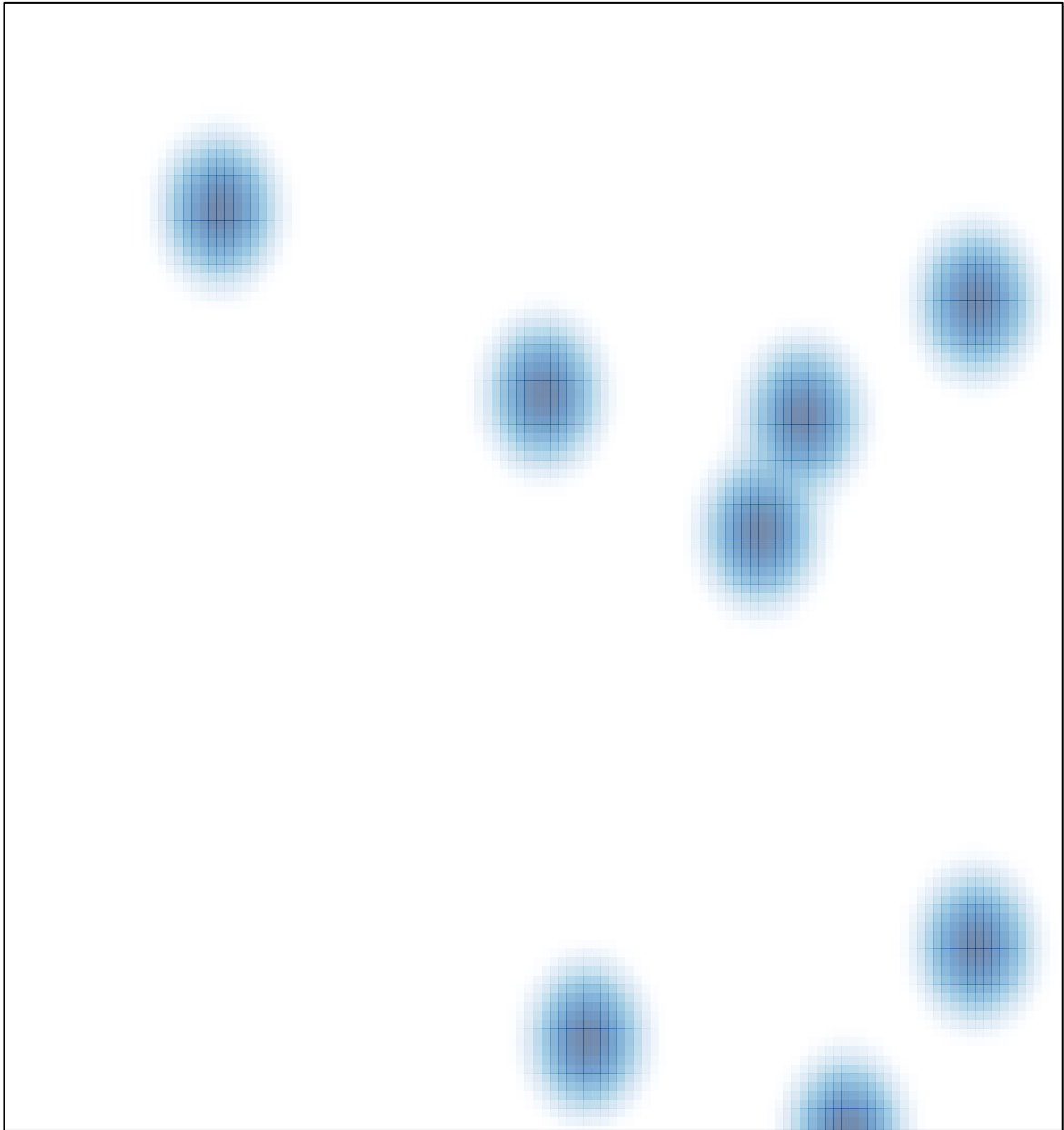


# S100 protein binding



# features = 11  
chi-square p = 0.82

# S100 protein binding



# features = 11 , max = 1