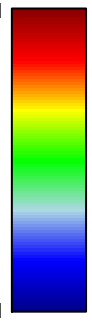
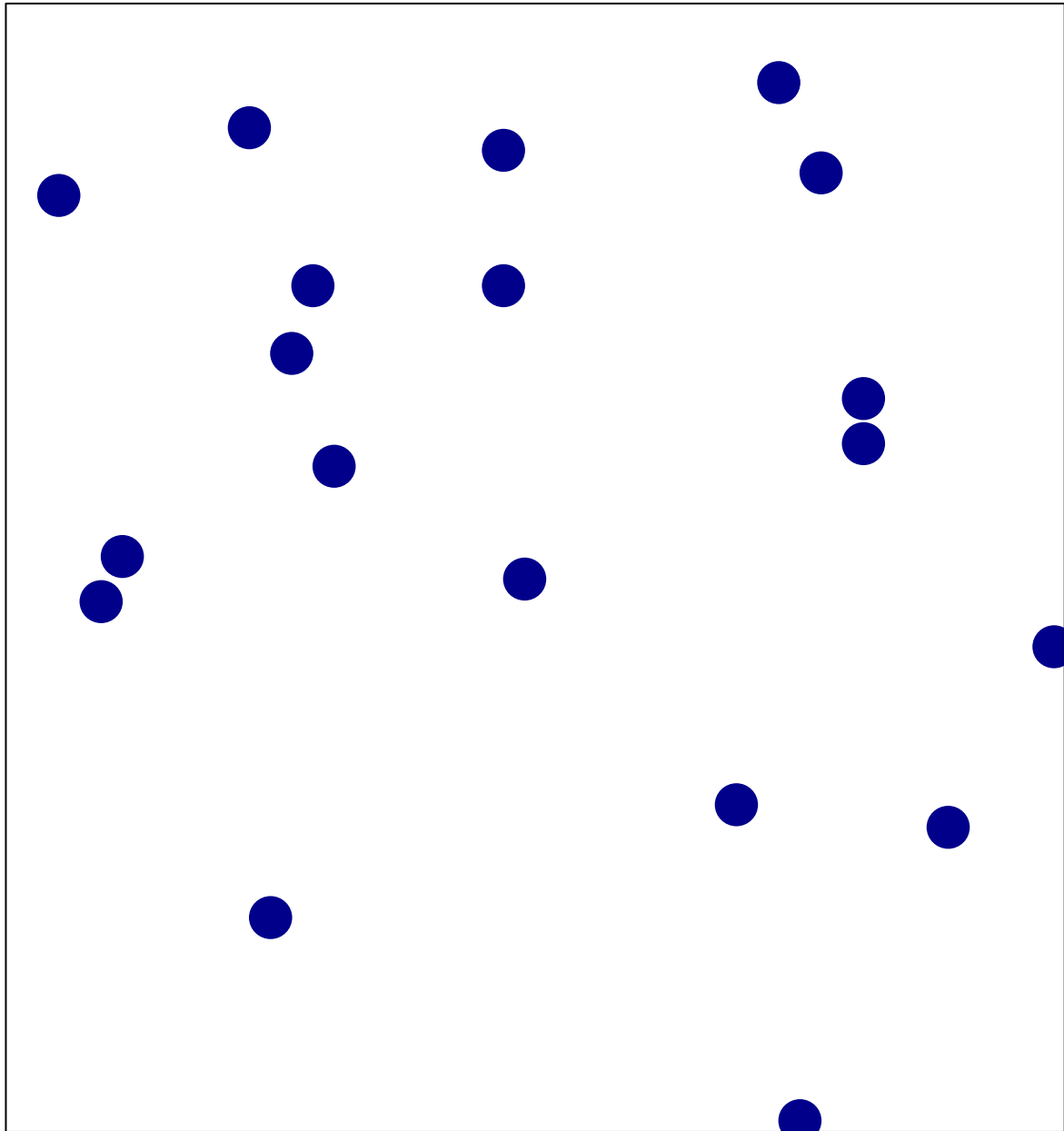
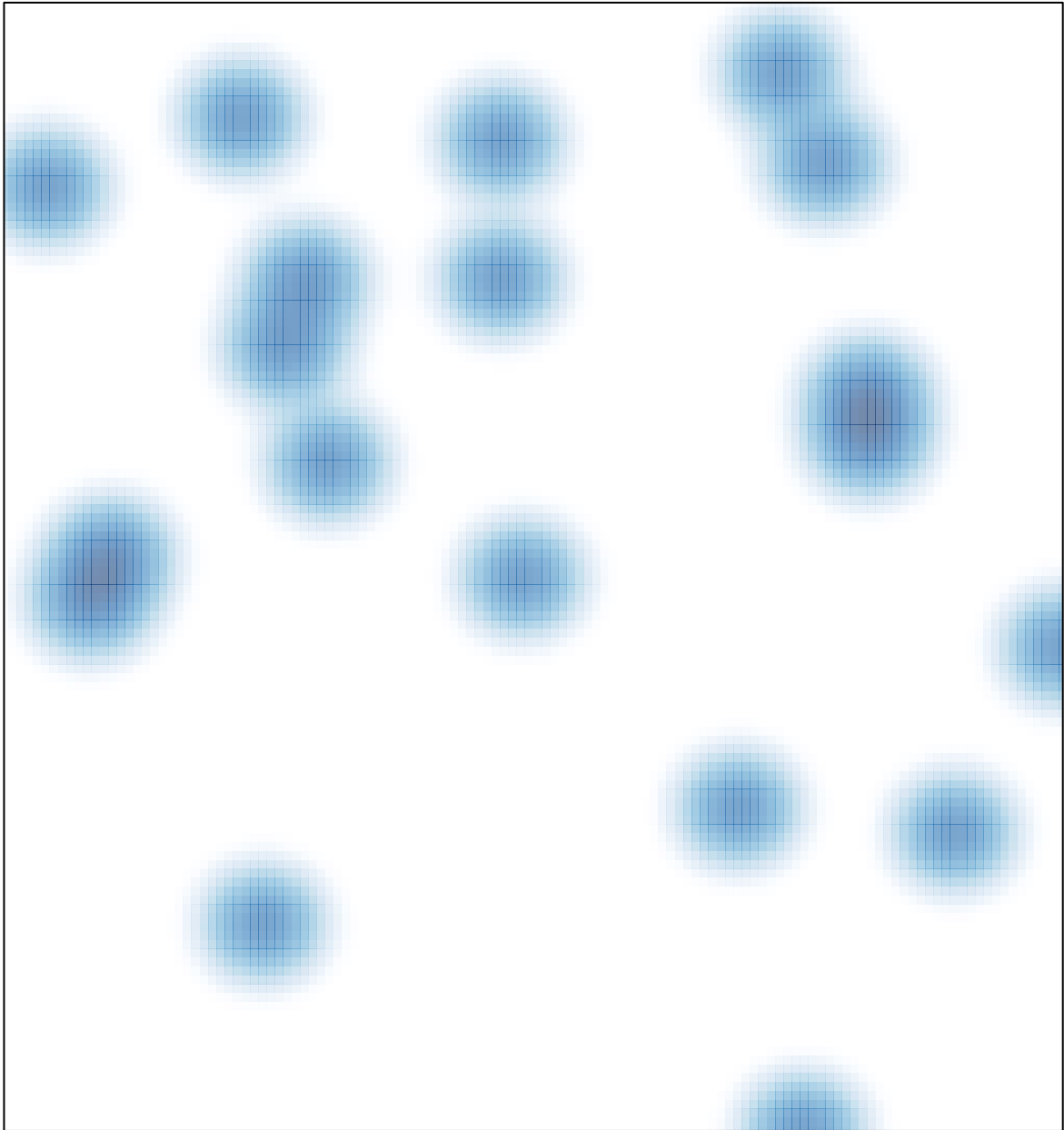


# retinoic acid receptor binding



# features = 19  
chi-square p = 0.86

# retinoic acid receptor binding



# features = 19 , max = 1