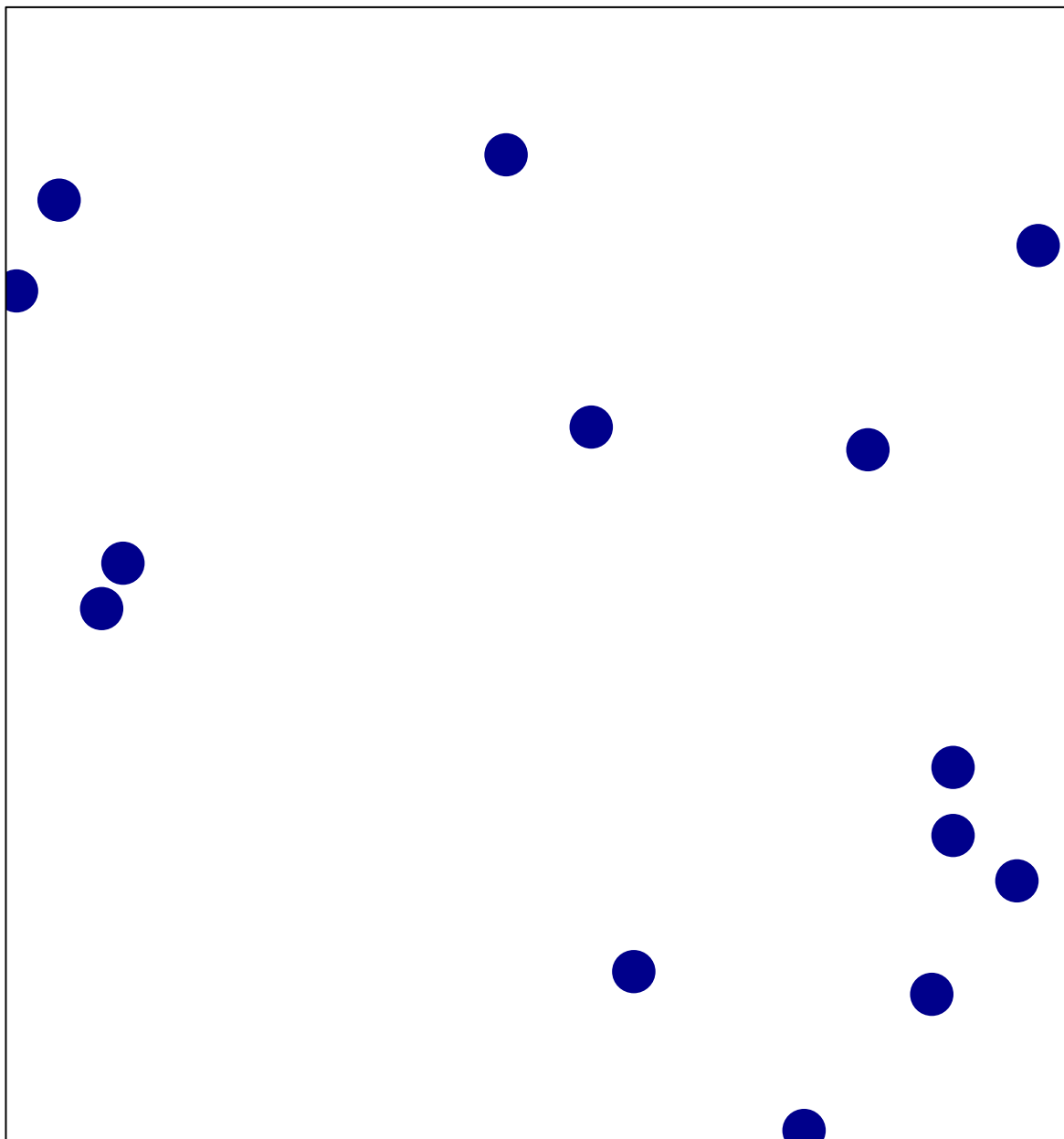
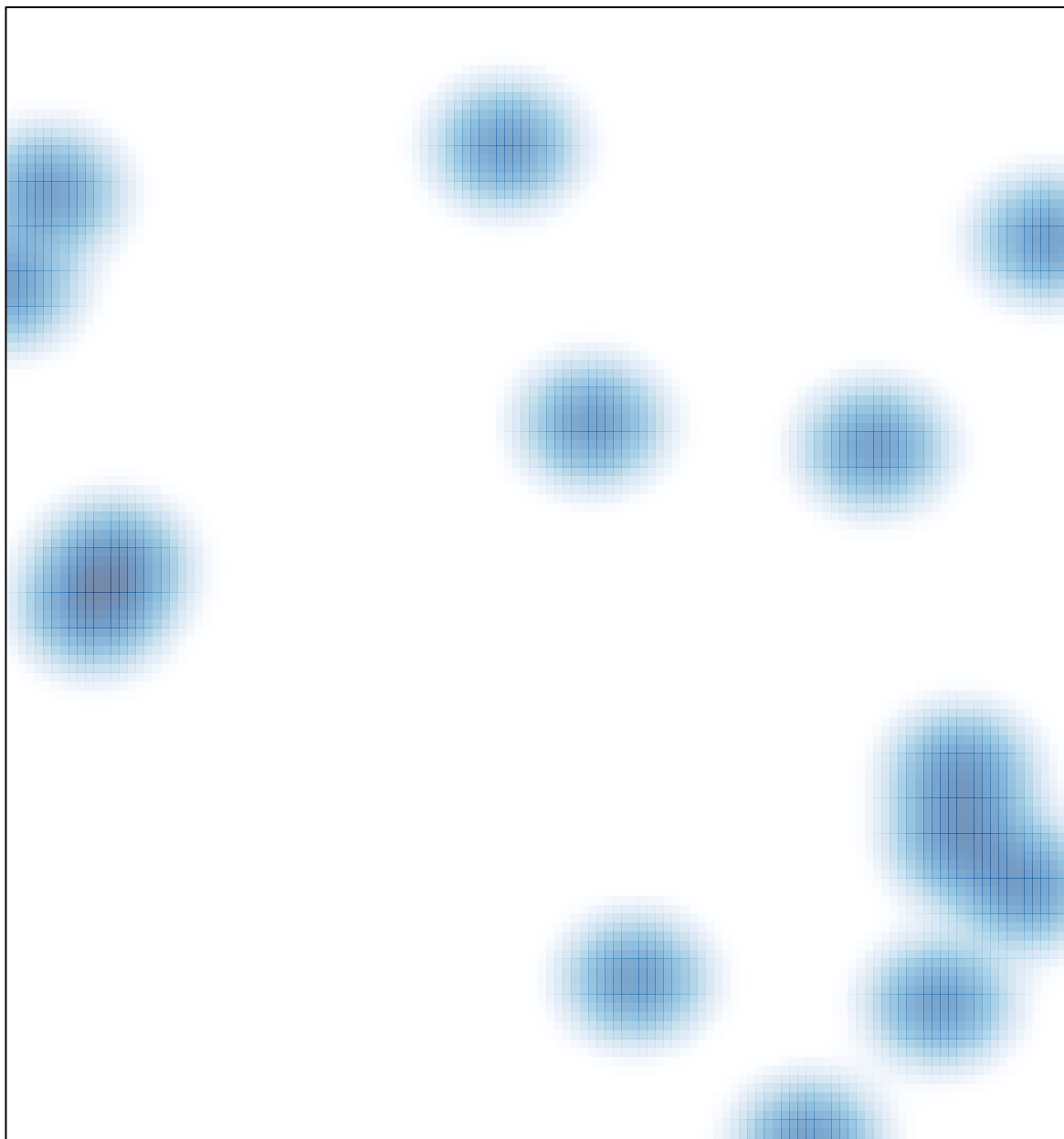


# retinoid X receptor binding



# features = 14  
chi-square p = 0.86

# retinoid X receptor binding



# features = 14 , max = 1