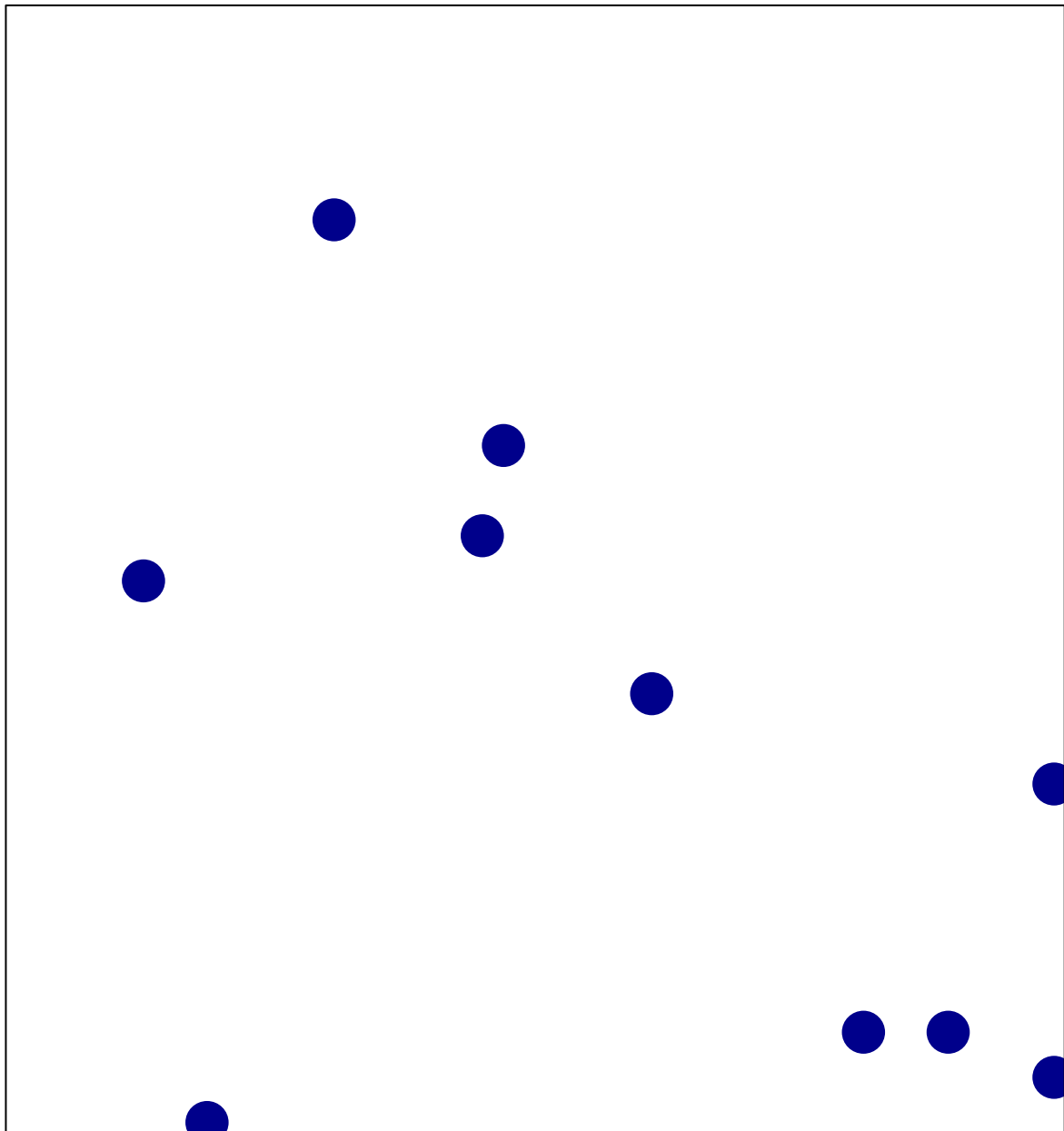
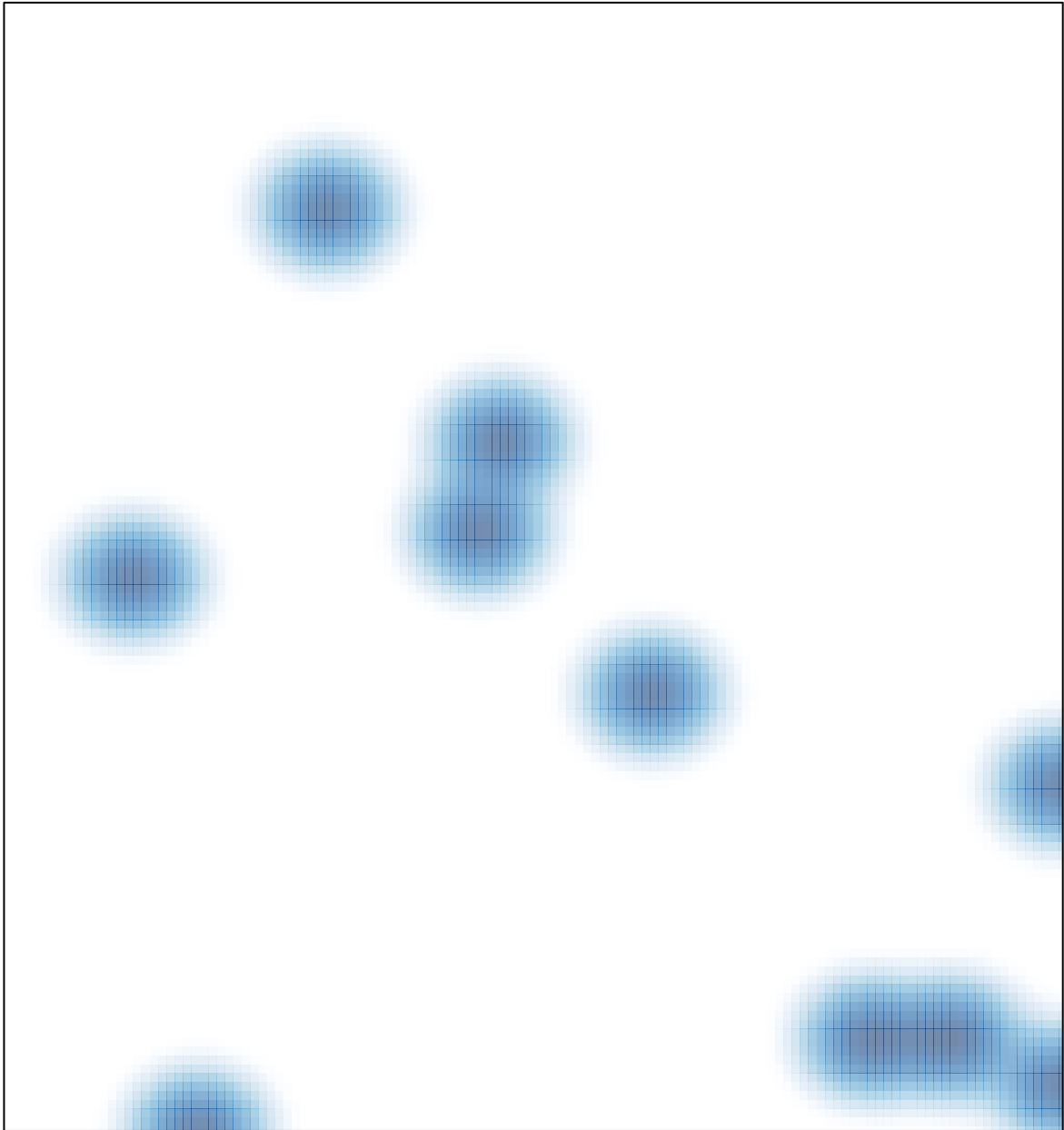


# diacylglycerol binding



# features = 10  
chi-square p = 0.84

# diacylglycerol binding



# features = 10 , max = 1