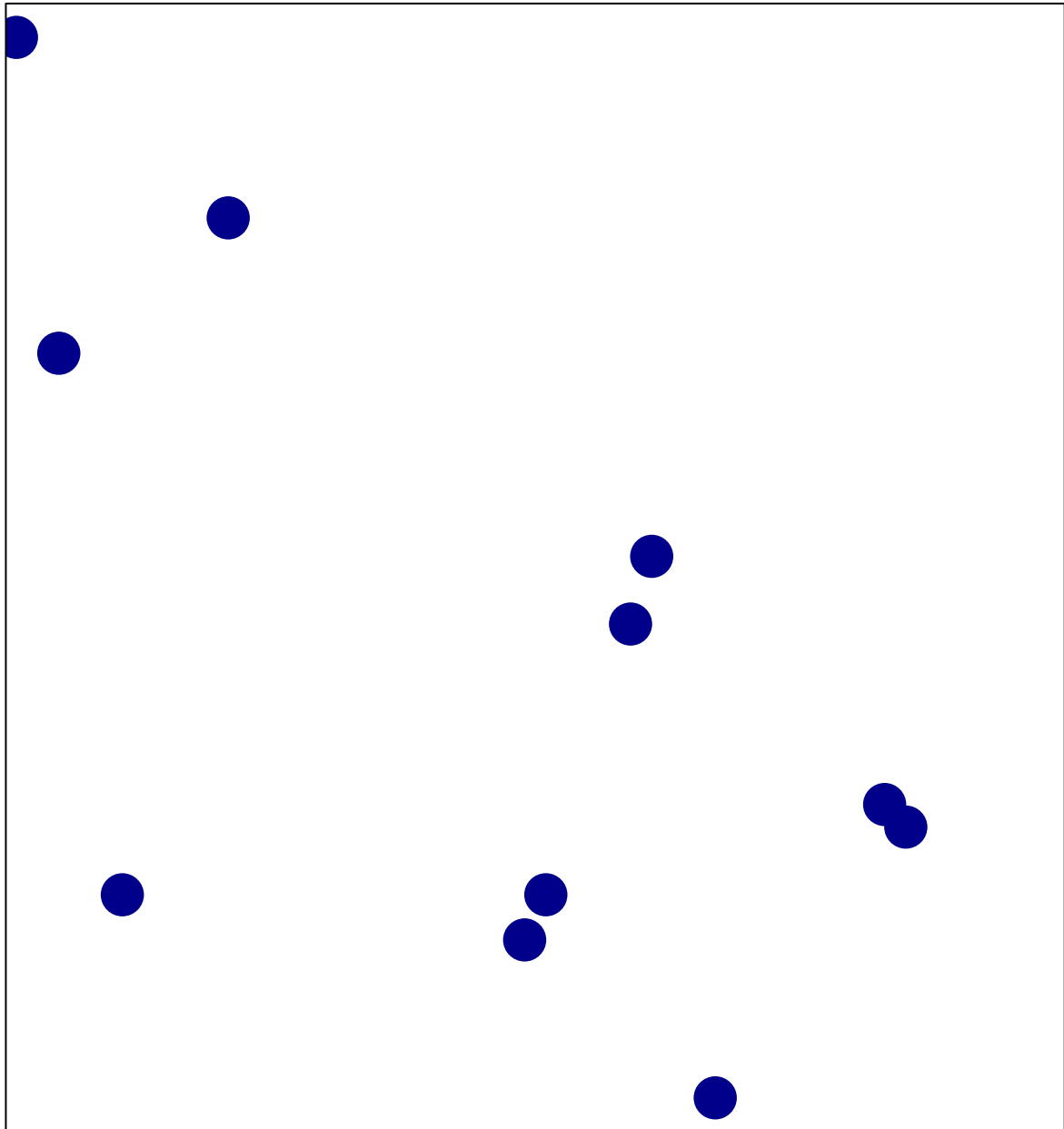
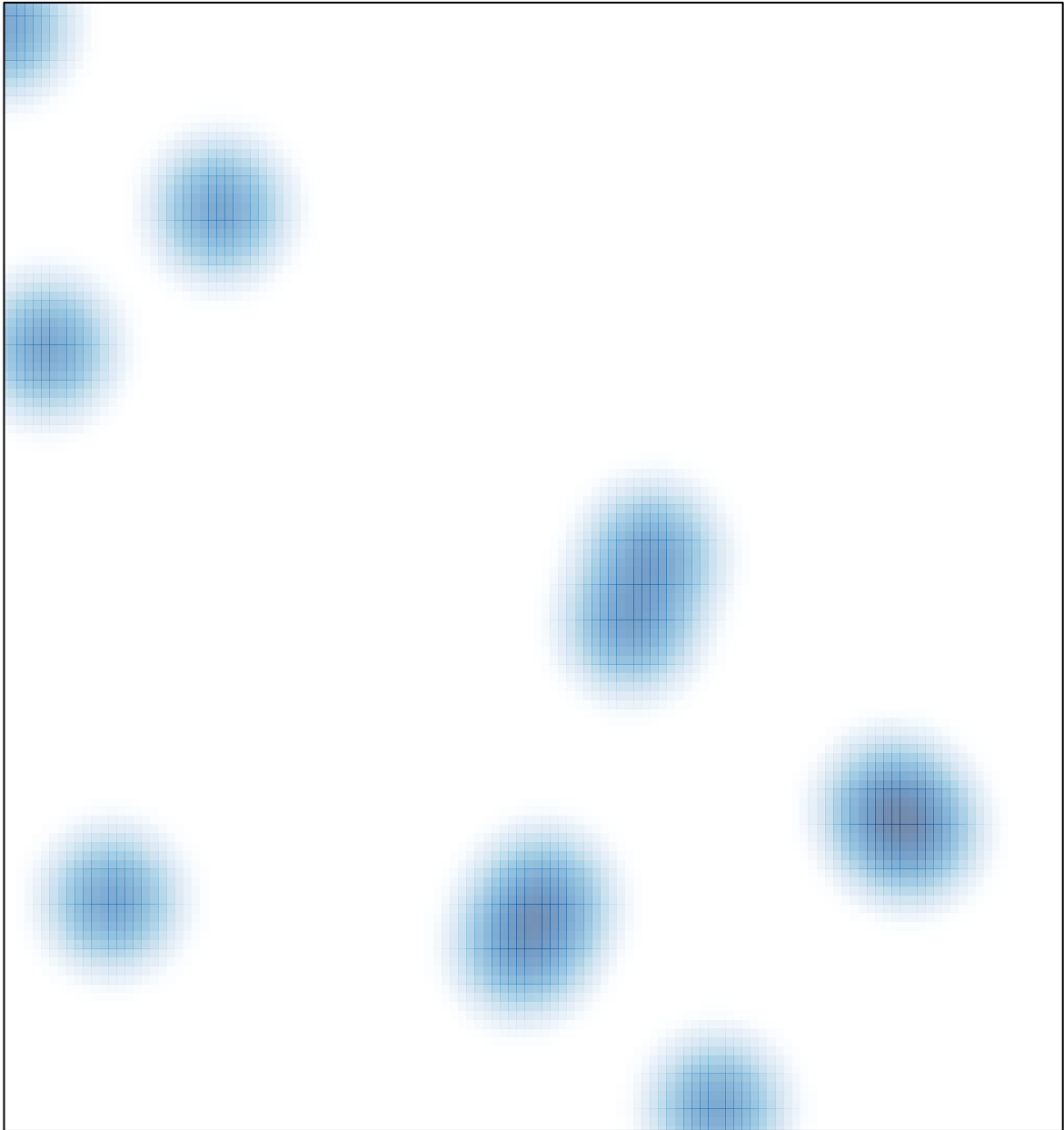


# mismatched DNA binding



# features = 11  
chi-square p = 0.85

# mismatched DNA binding



# features = 11 , max = 1