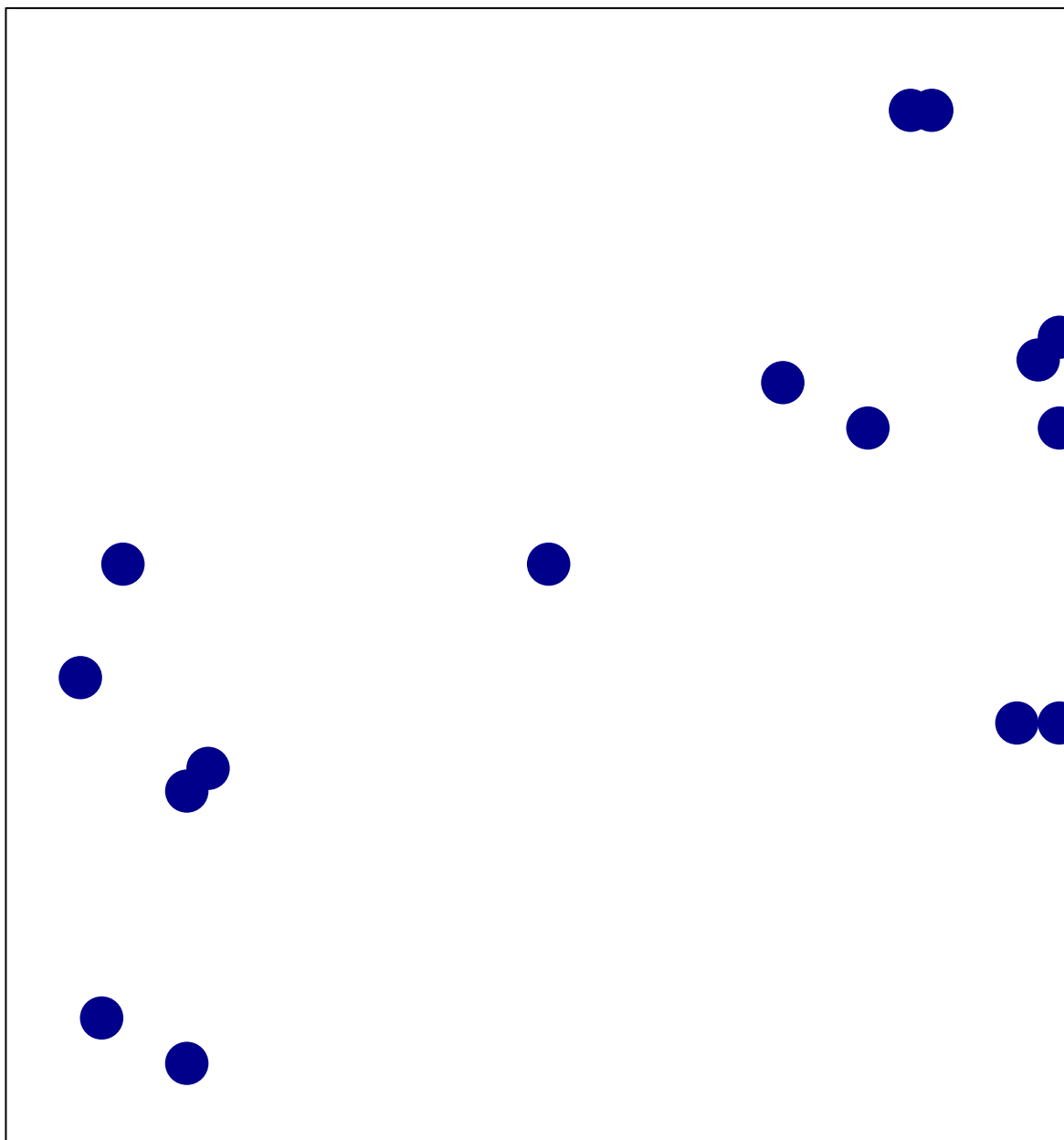
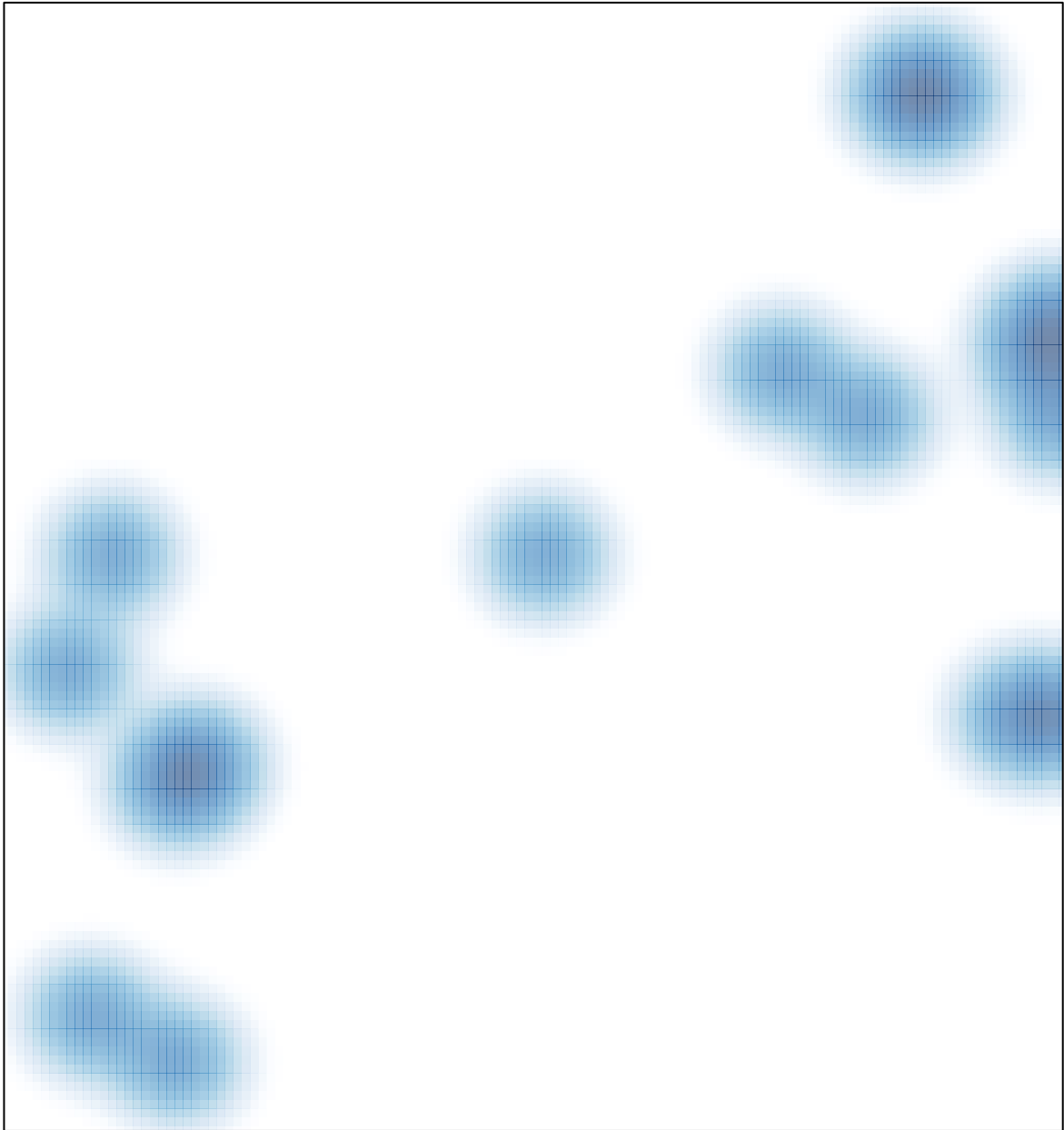


# nucleosome disassembly



# features = 16  
chi-square p = 0.83

# nucleosome disassembly



# features = 16 , max = 1