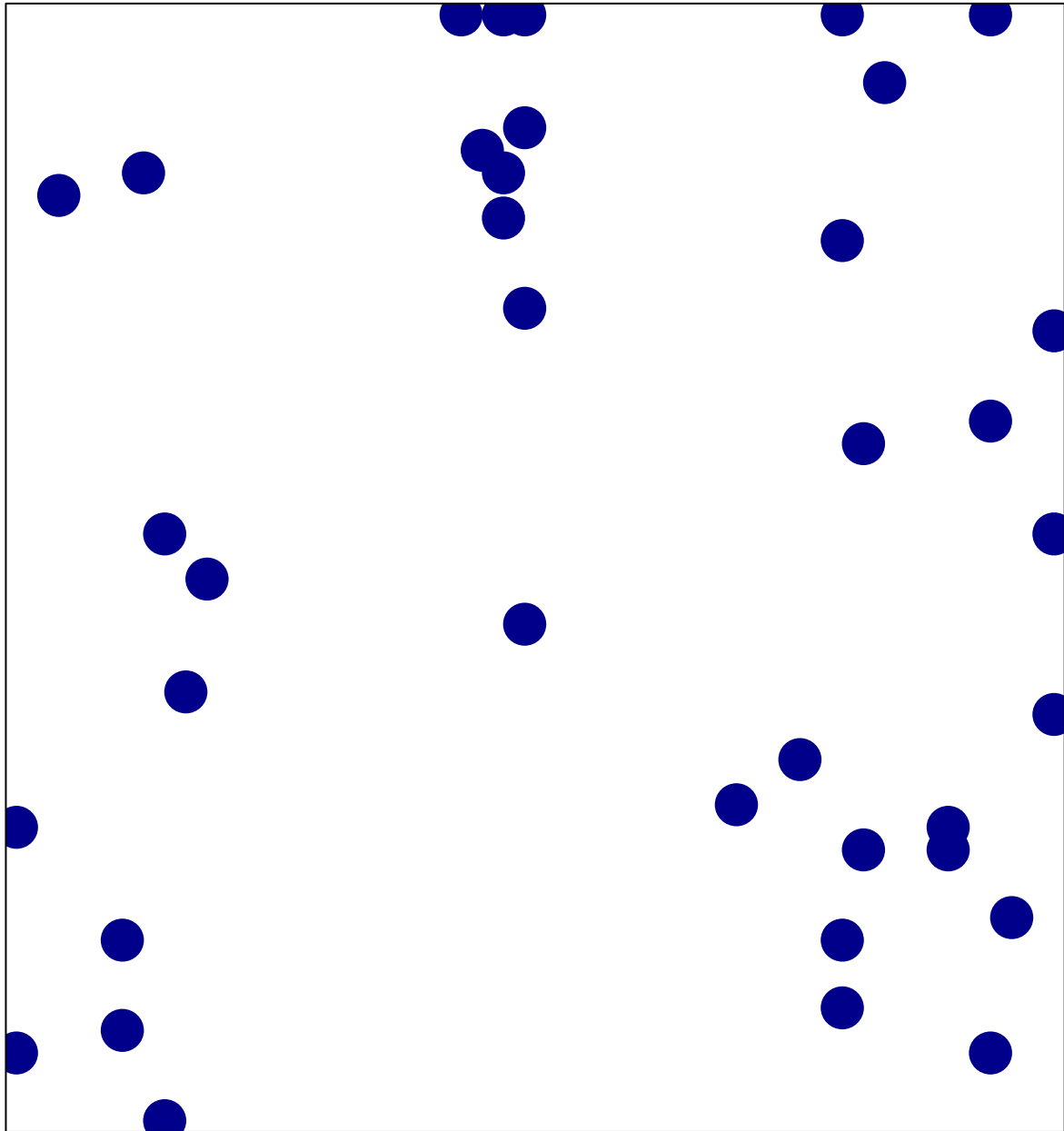
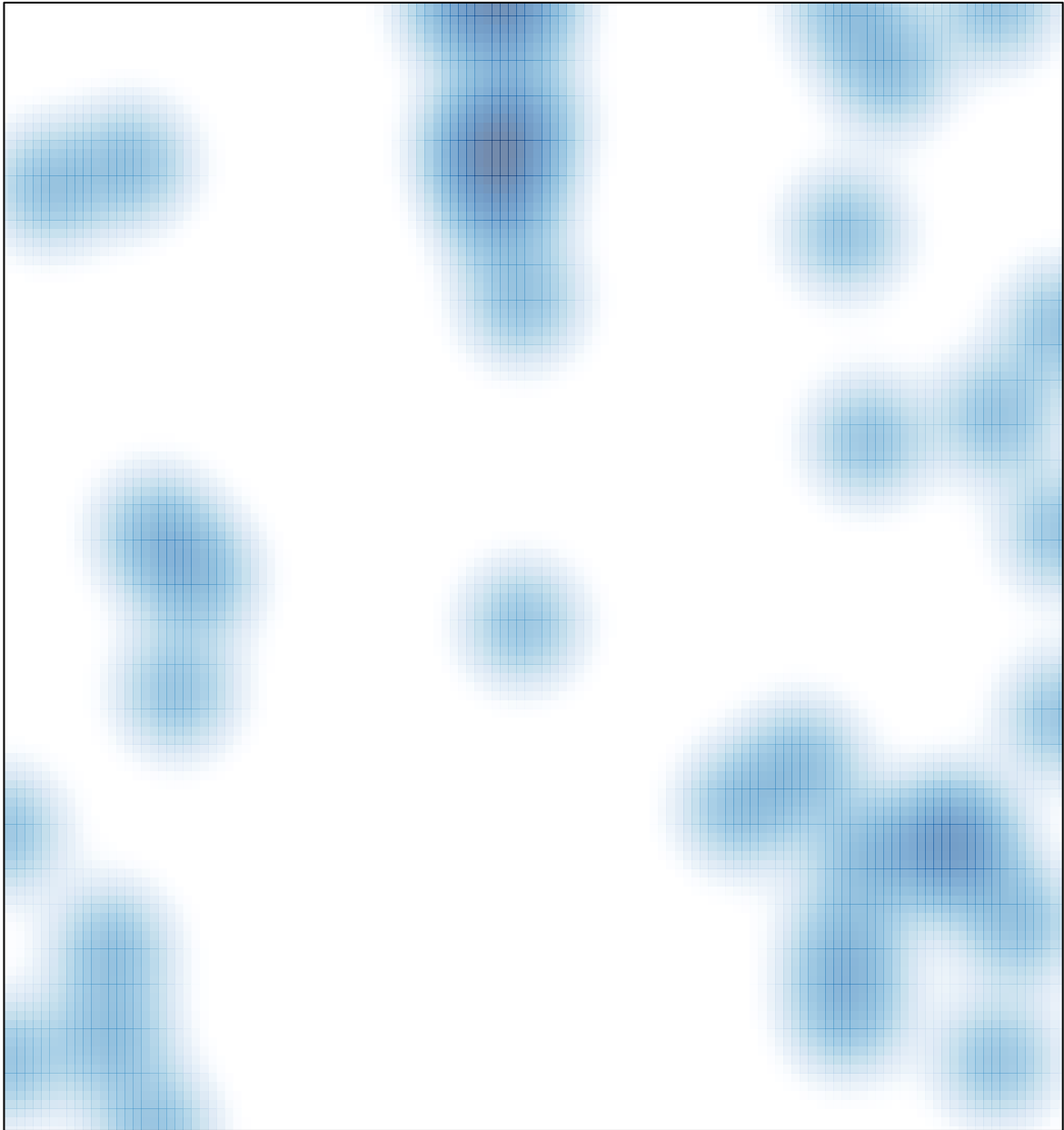


# androgen receptor binding



# features = 37  
chi-square p = 0.81

# androgen receptor binding



# features = 37 , max = 1