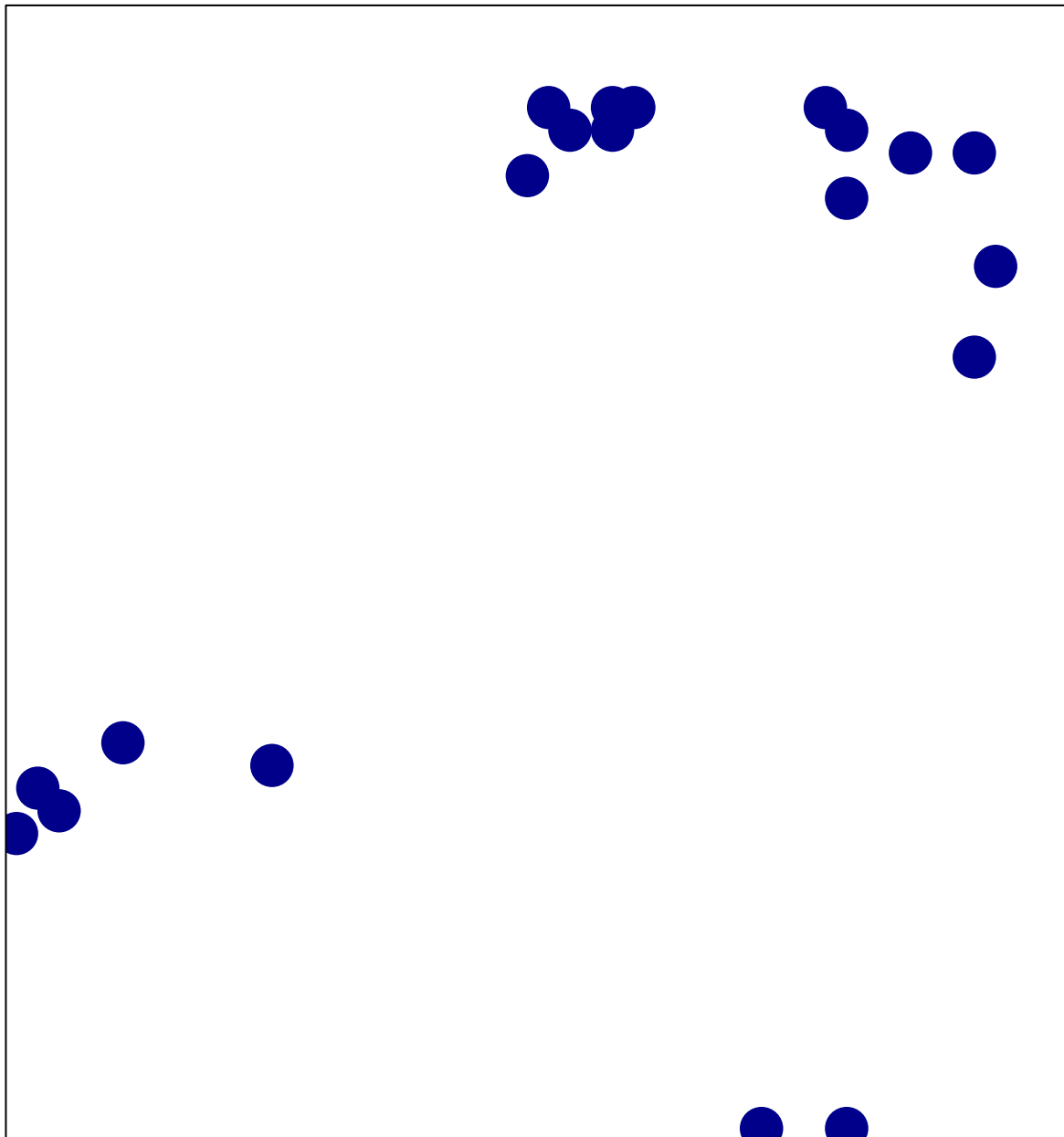


# DNA duplex unwinding



# features = 20  
chi-square p = 0.81

# DNA duplex unwinding



# features = 20 , max = 1