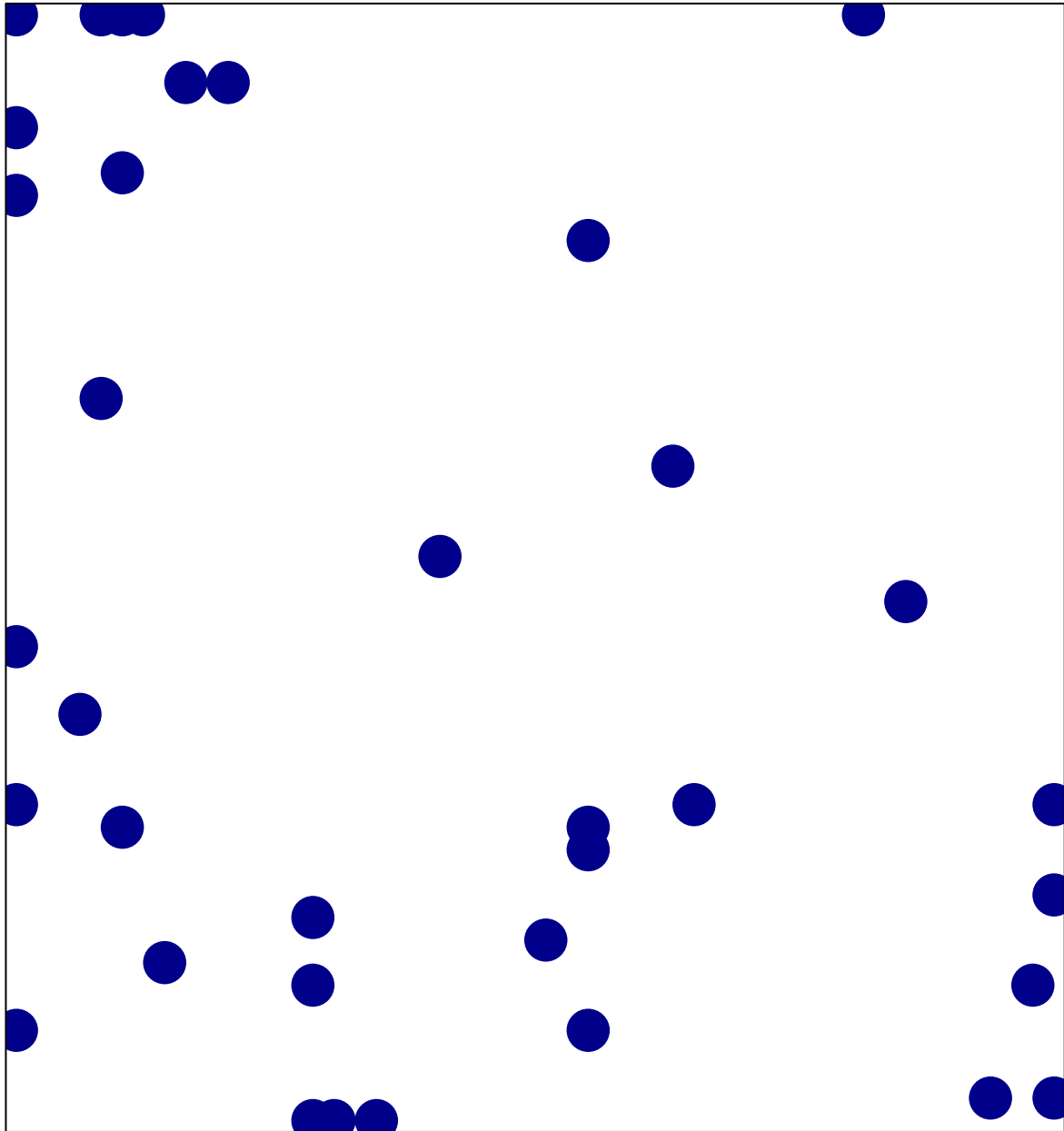


# tubulin binding



# features = 36  
chi-square p = 0.79

# tubulin binding



# features = 36 , max = 1