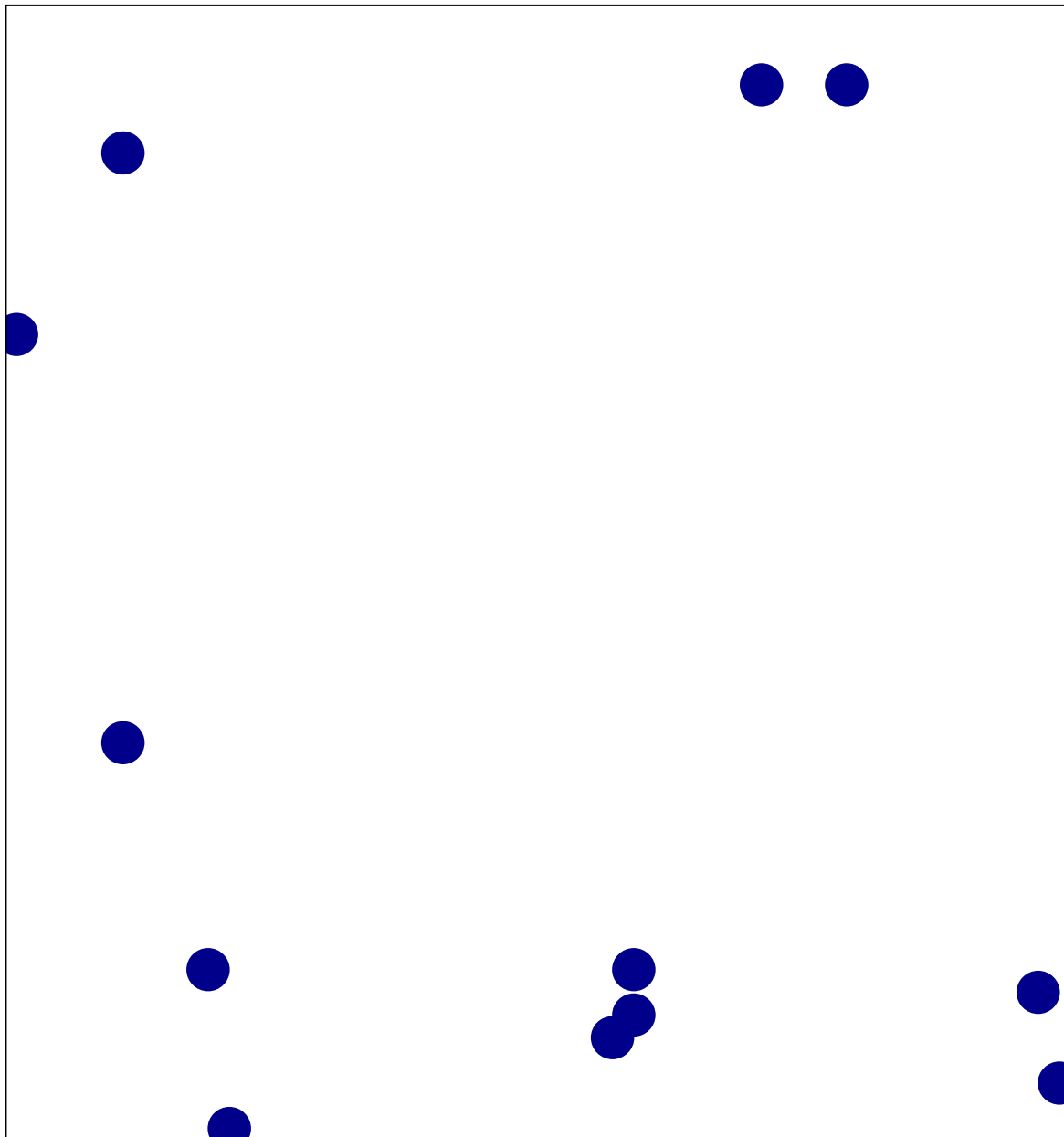
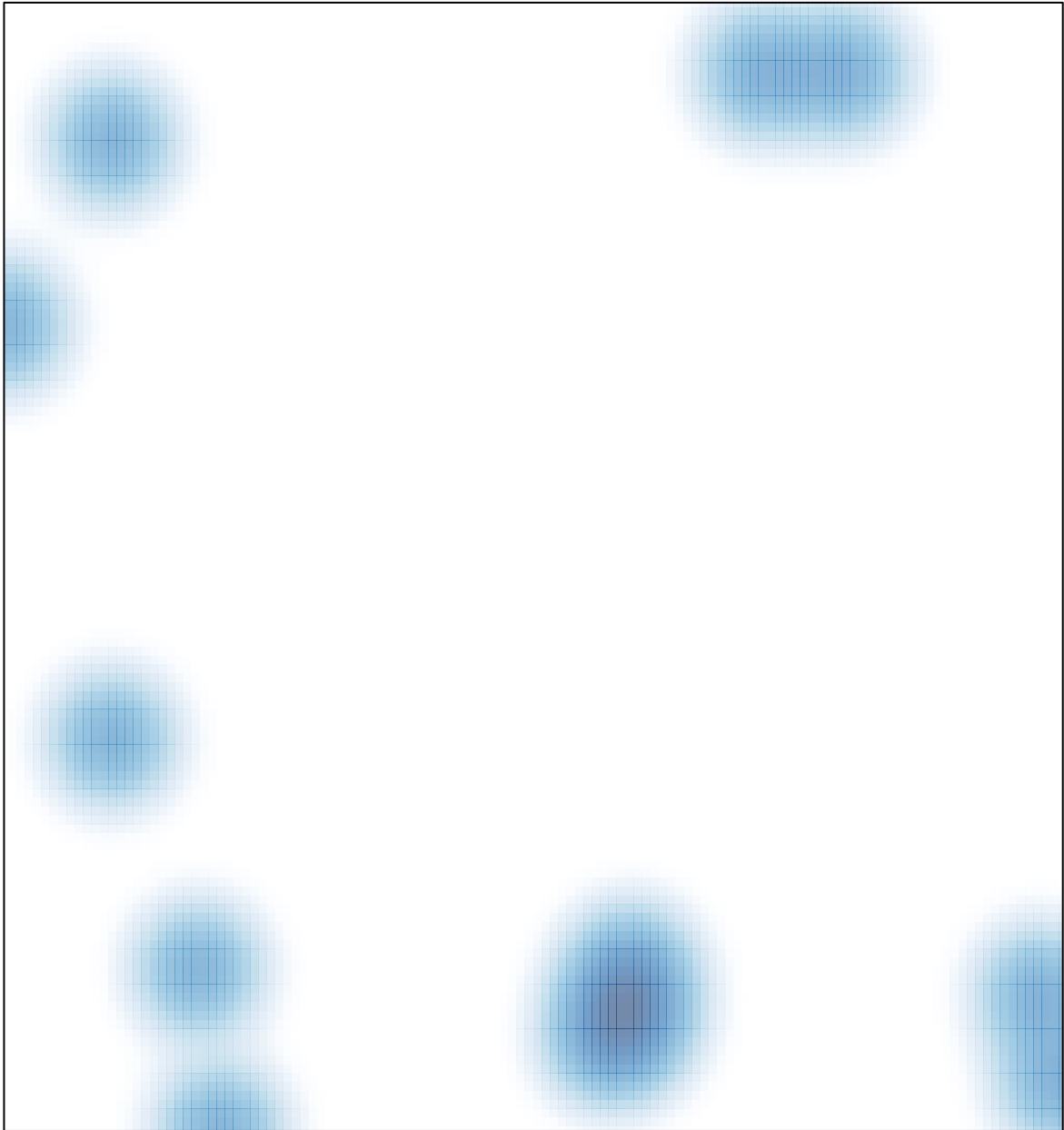


# lipopolysaccharide binding



# features = 12  
chi-square p = 0.83

# lipopolysaccharide binding



# features = 12 , max = 1