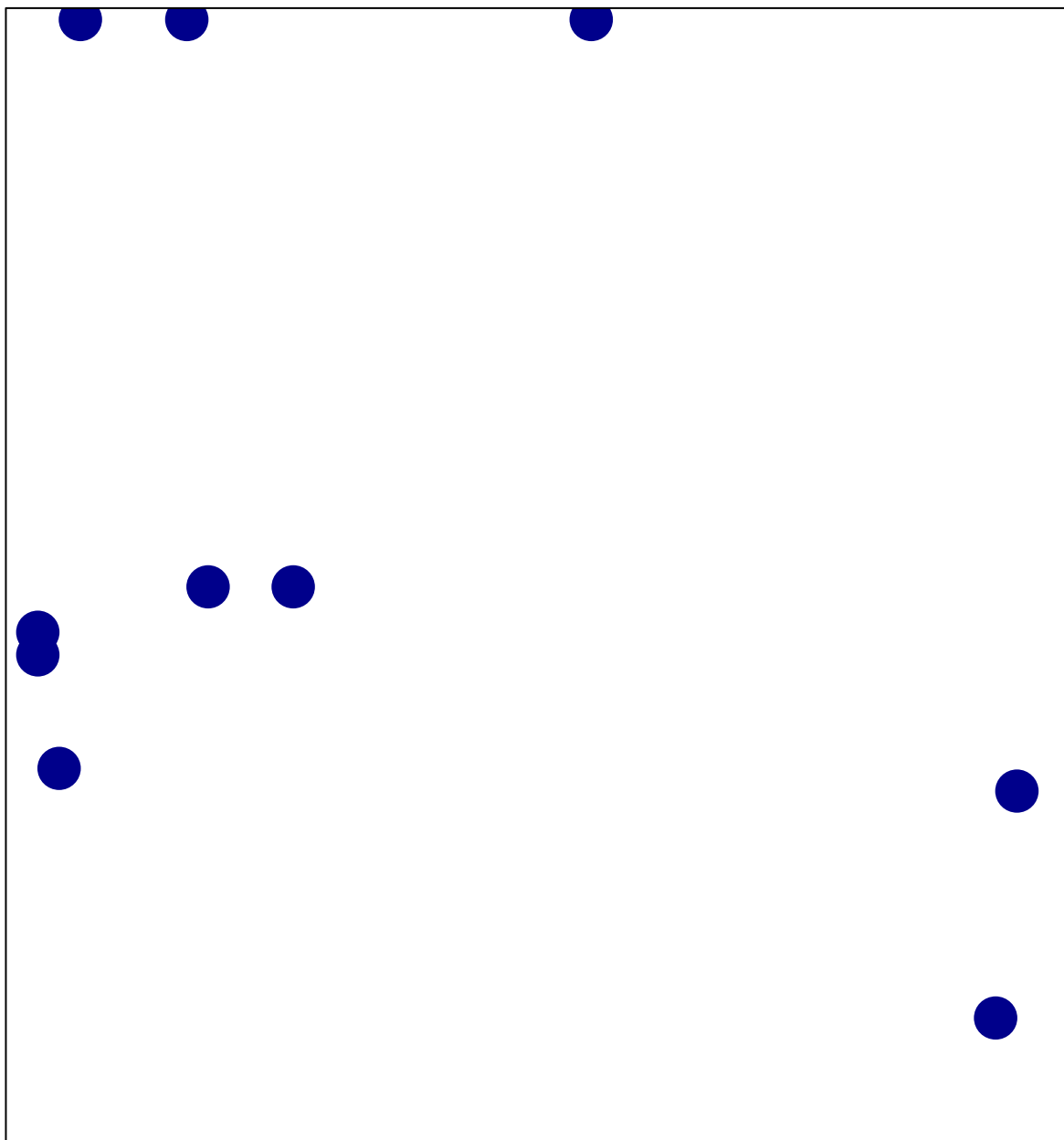


# monosaccharide binding



# features = 10  
chi-square p = 0.82

# monosaccharide binding



# features = 10 , max = 1