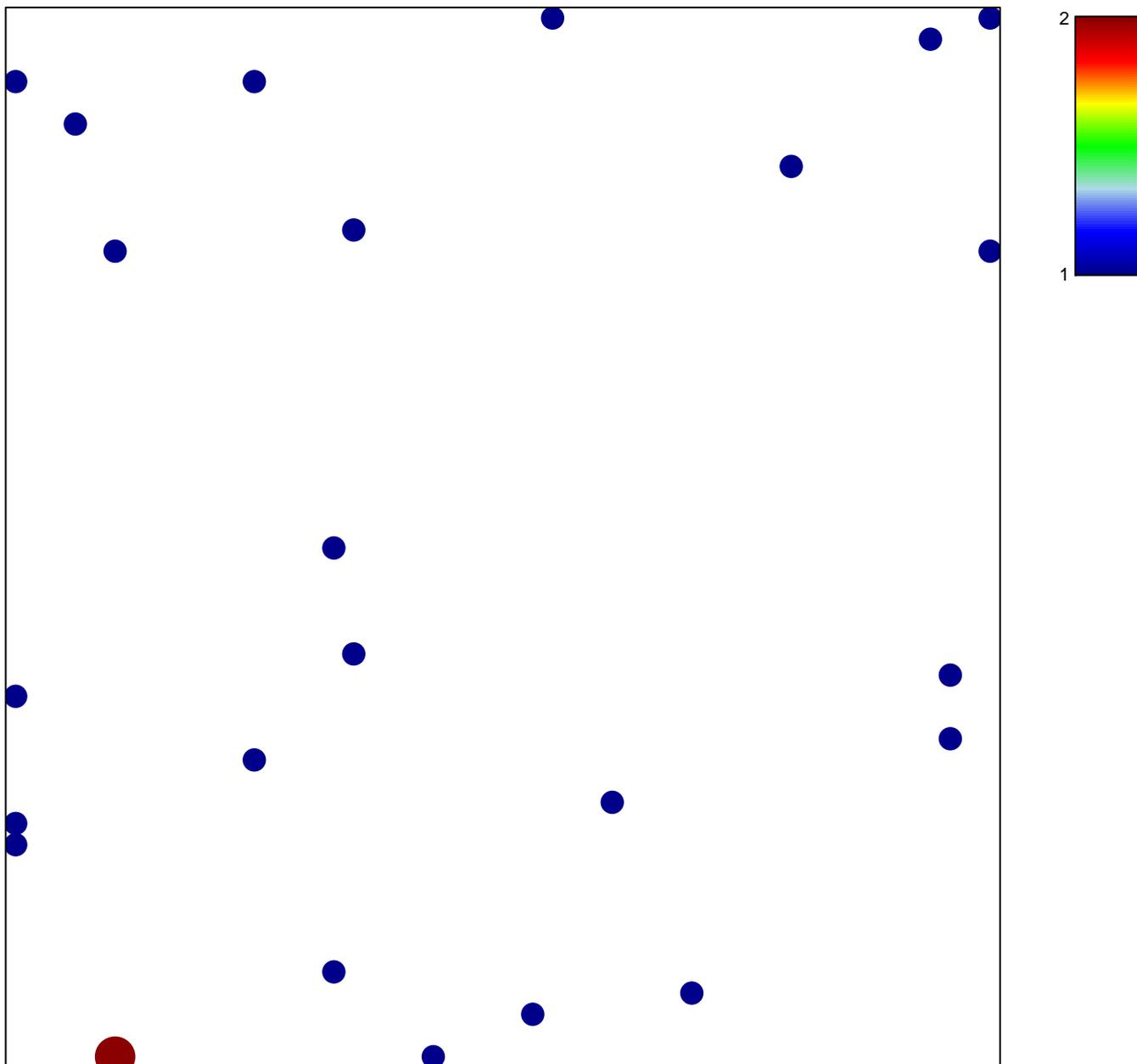
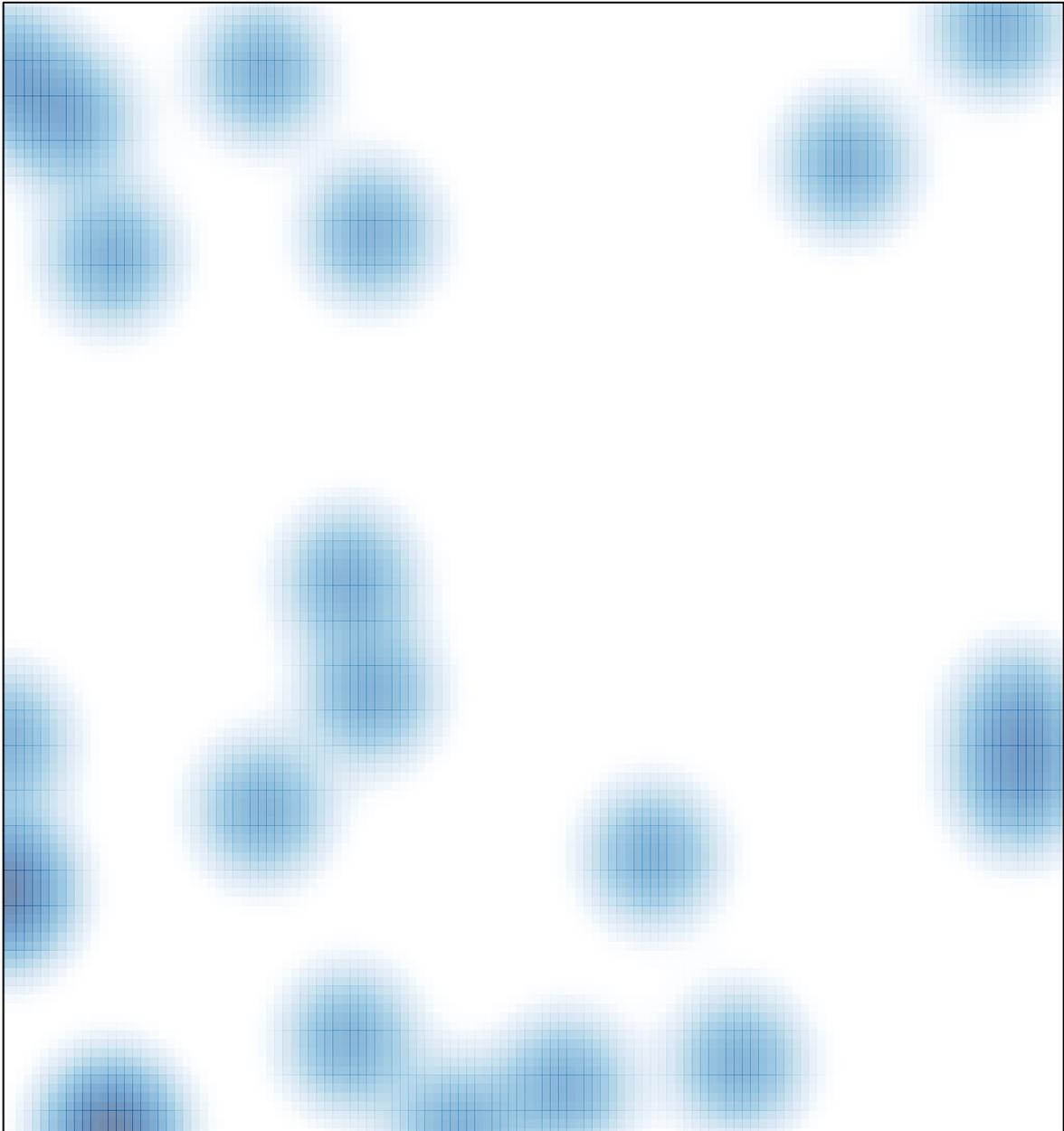


# phosphatidylinositol-3-phosphate binding



# features = 25  
chi-square p = 0.82

# phosphatidylinositol-3-phosphate binding



# features = 25 , max = 2