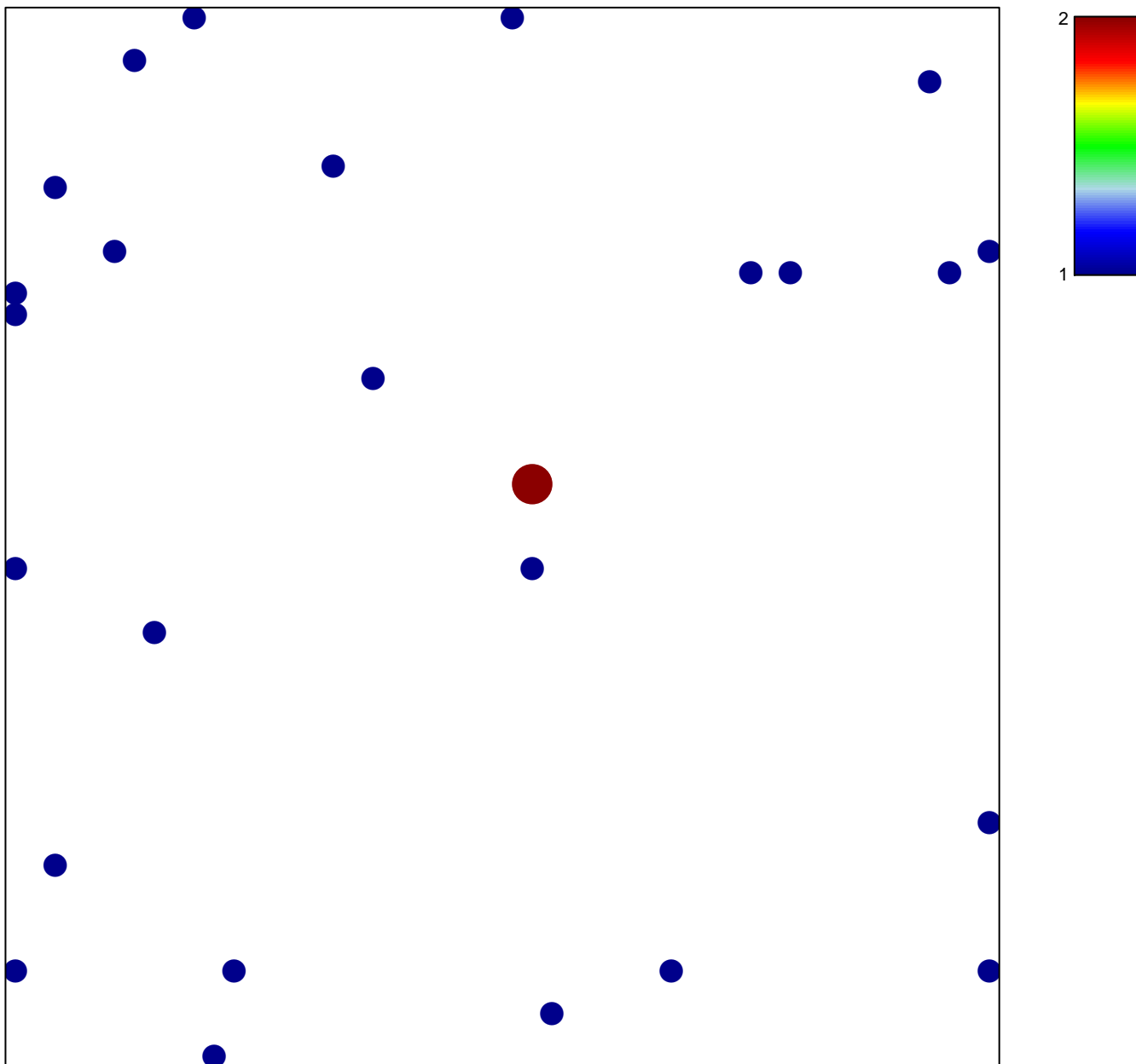
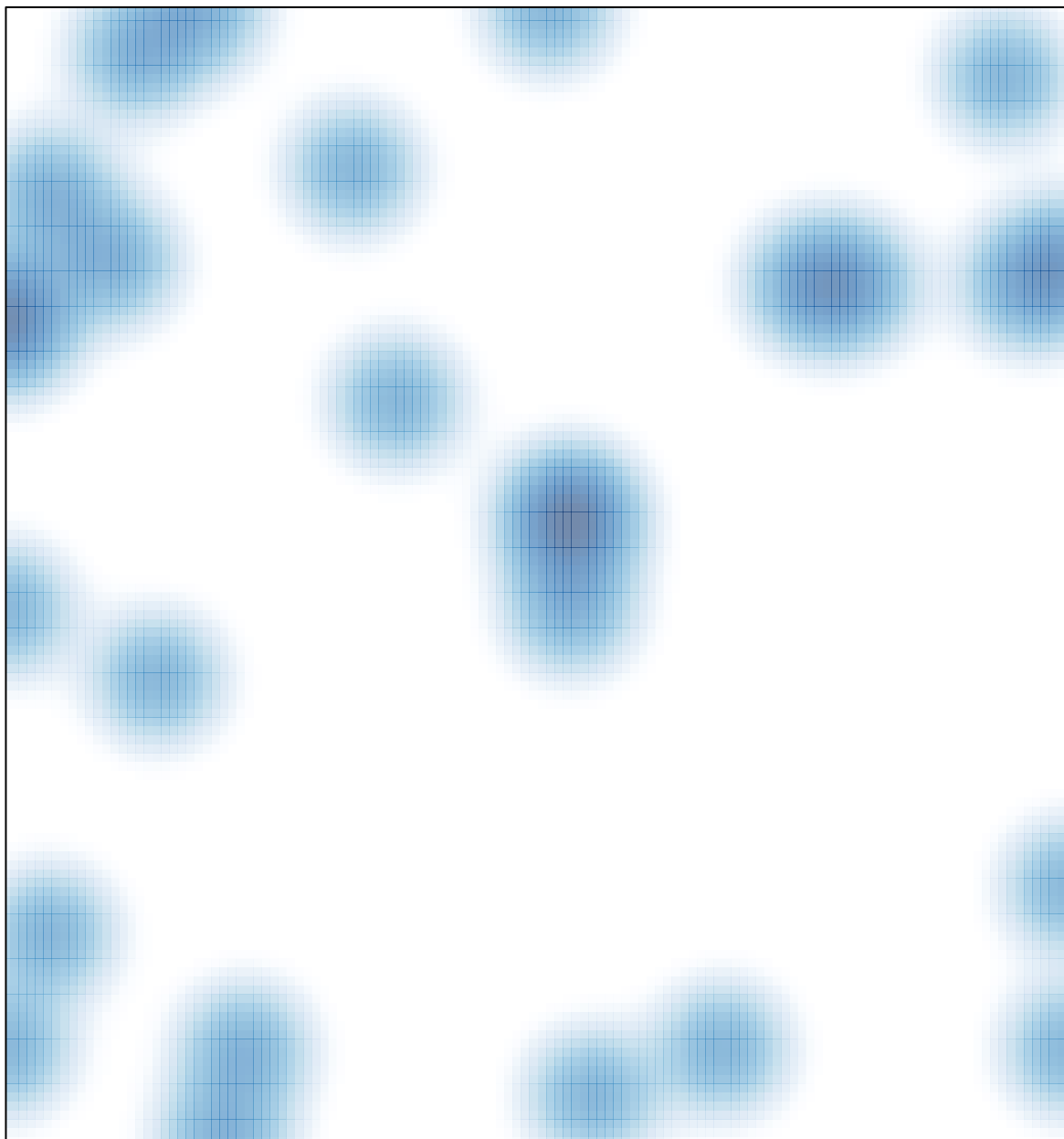


# phosphatidylinositol-4,5-bisphosphate binding



# features = 27  
chi-square p = 0.81

# phosphatidylinositol-4,5-bisphosphate binding



# features = 27 , max = 2