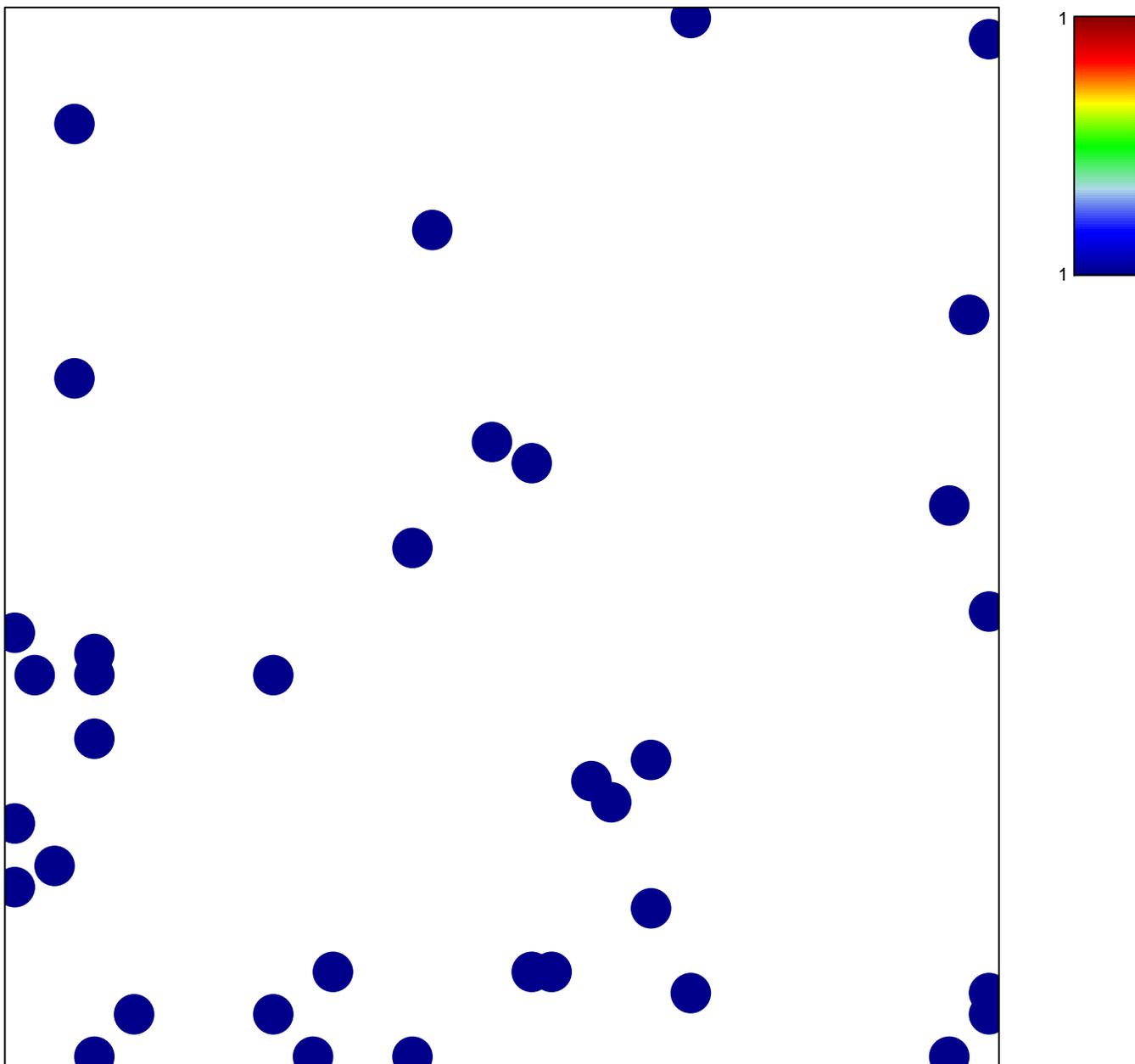
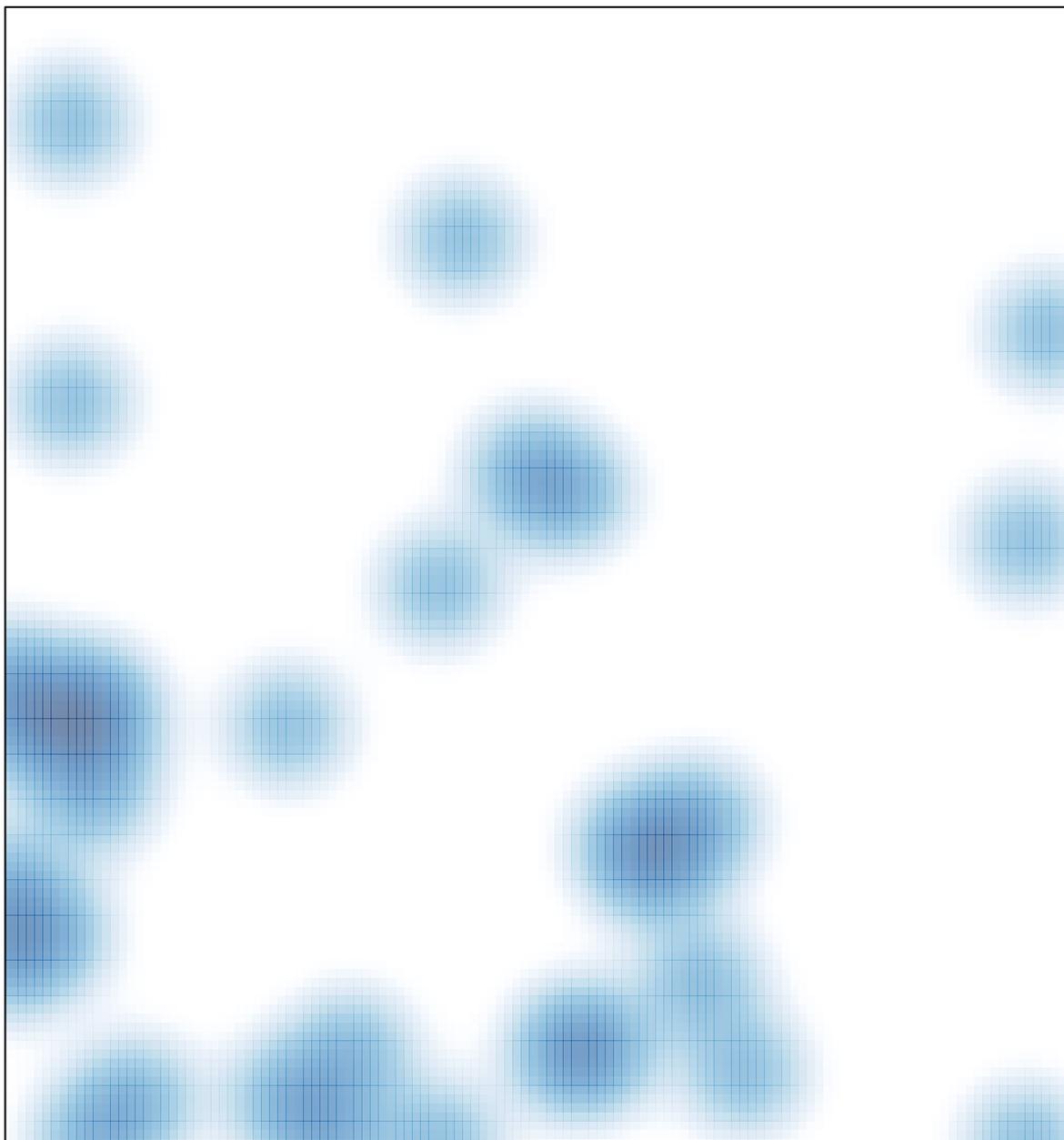


# phosphatidylinositol-4,5-bisphosphate binding



# features = 36  
chi-square p = 0.82

# phosphatidylinositol-4,5-bisphosphate binding



# features = 36 , max = 1