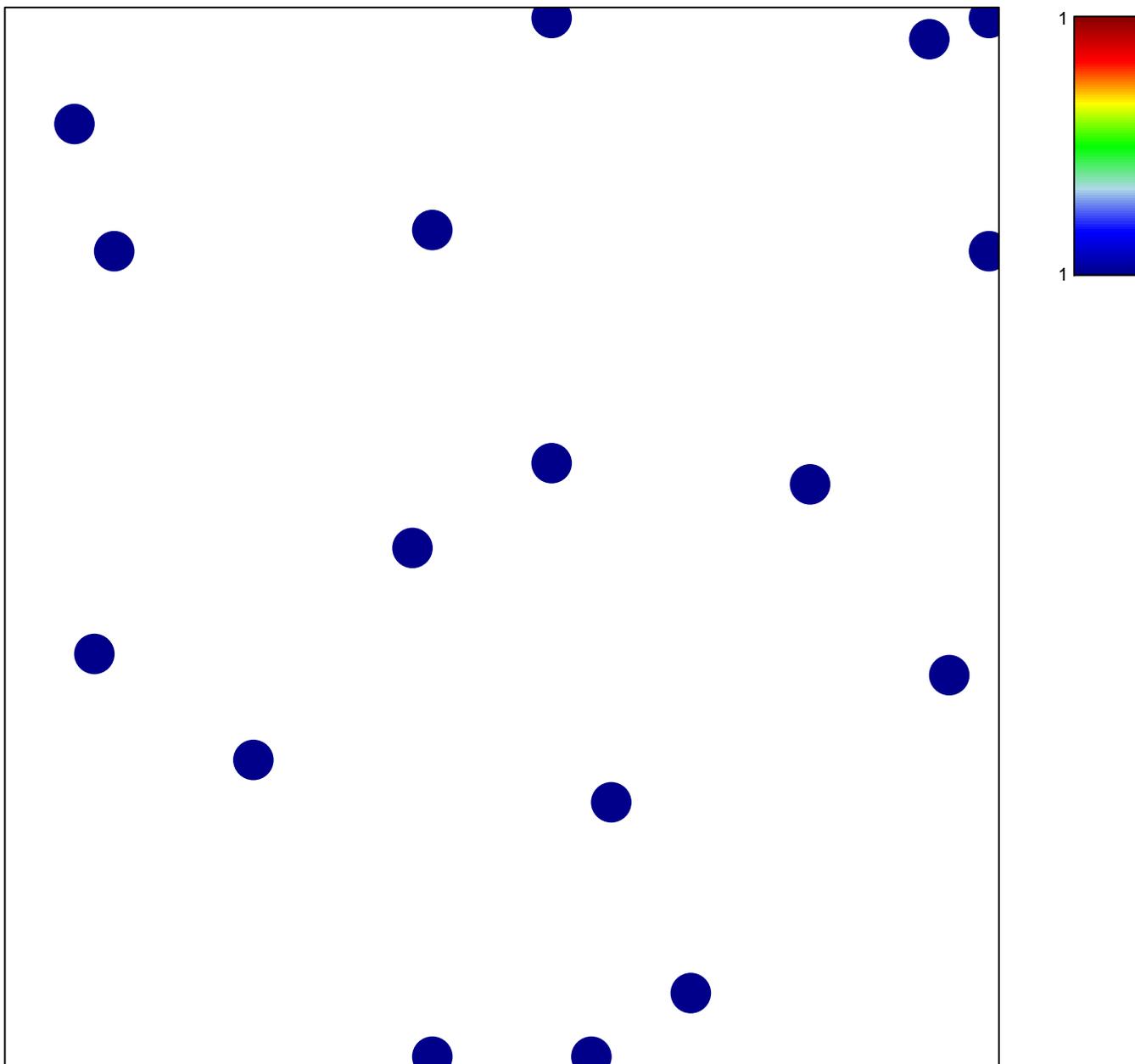
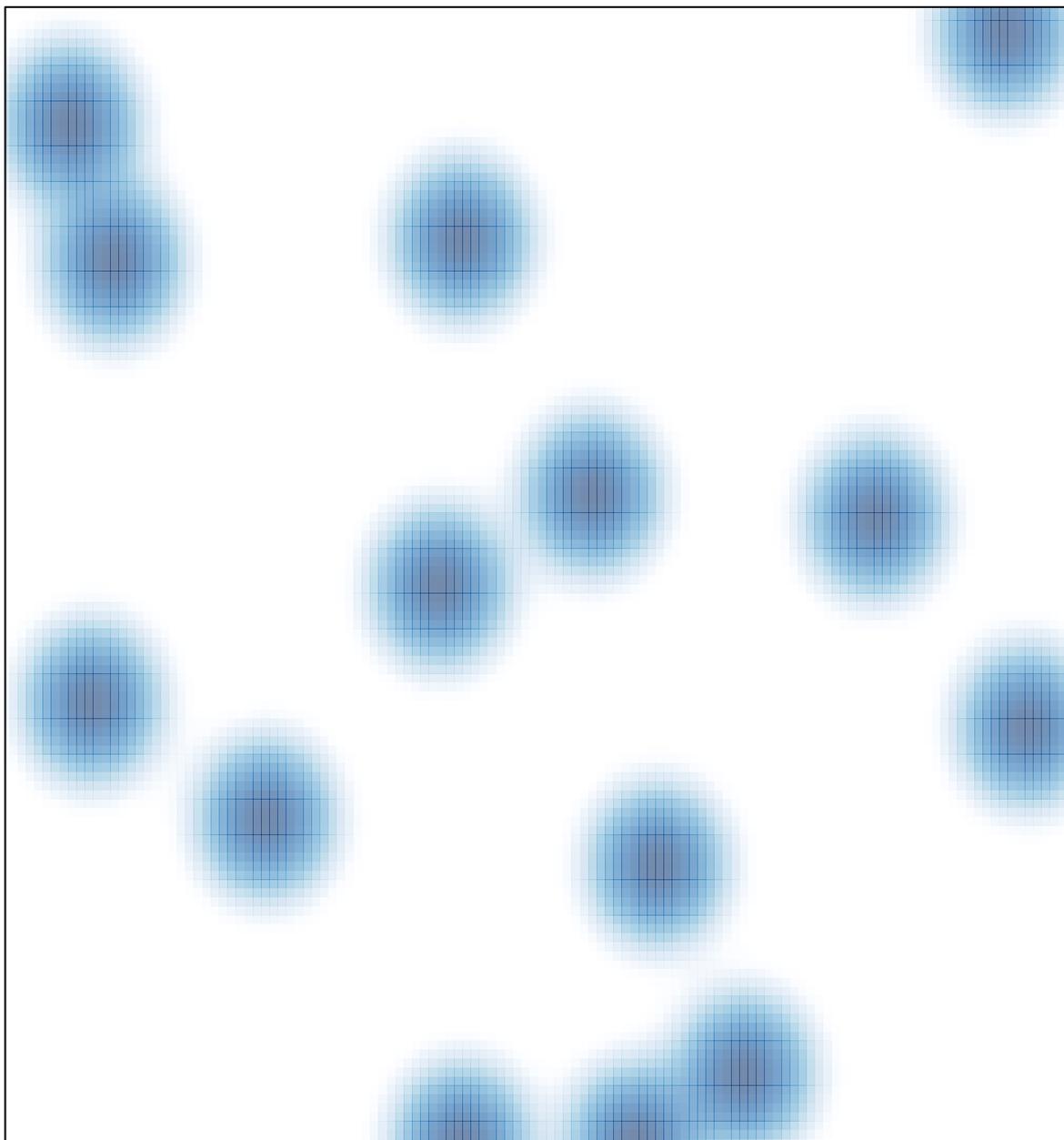


# phosphatidylinositol-3,5-bisphosphate binding



# features = 17  
chi-square p = 0.84

# phosphatidylinositol-3,5-bisphosphate binding



# features = 17 , max = 1