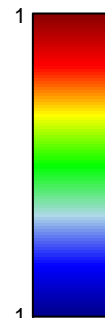
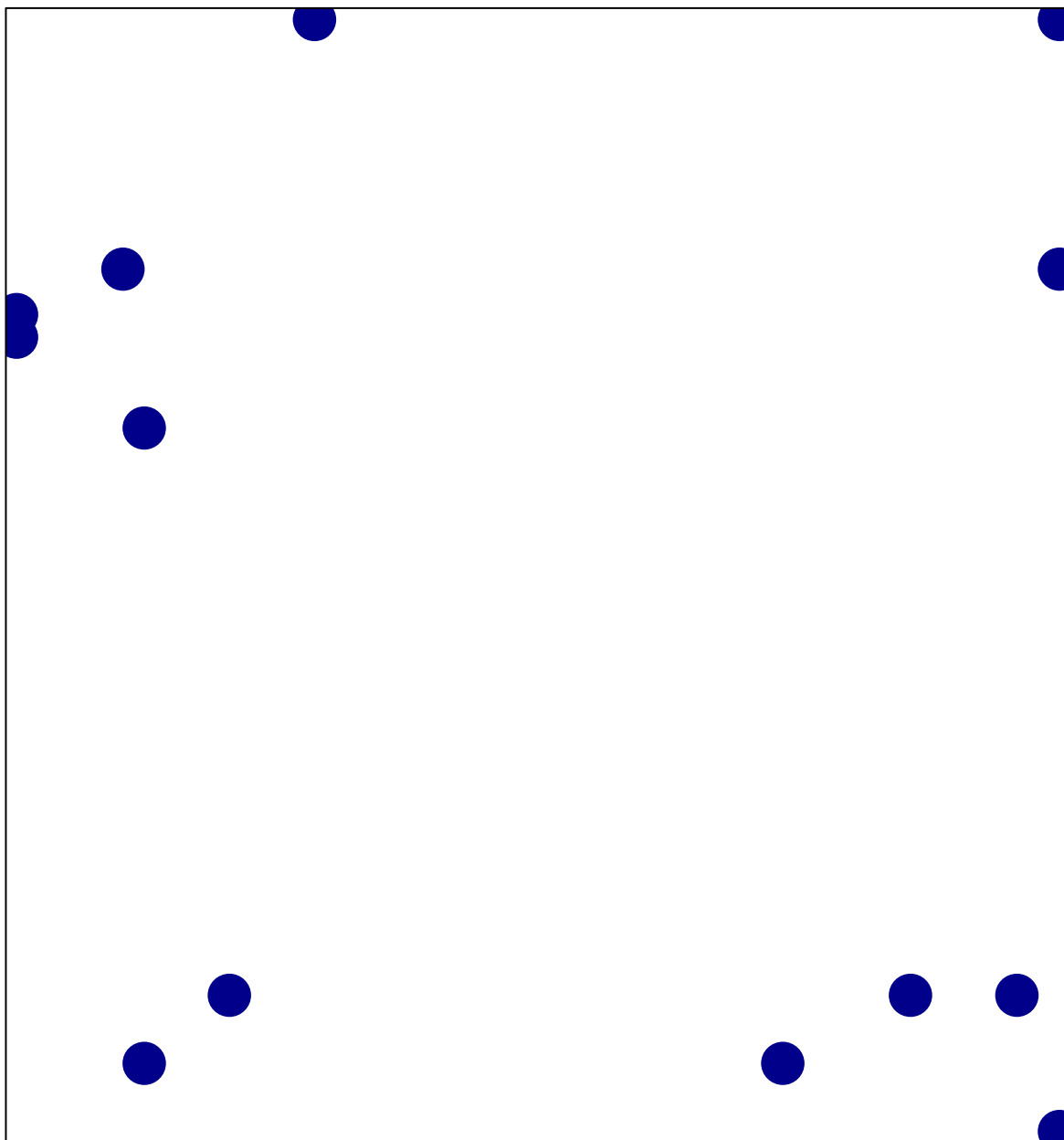
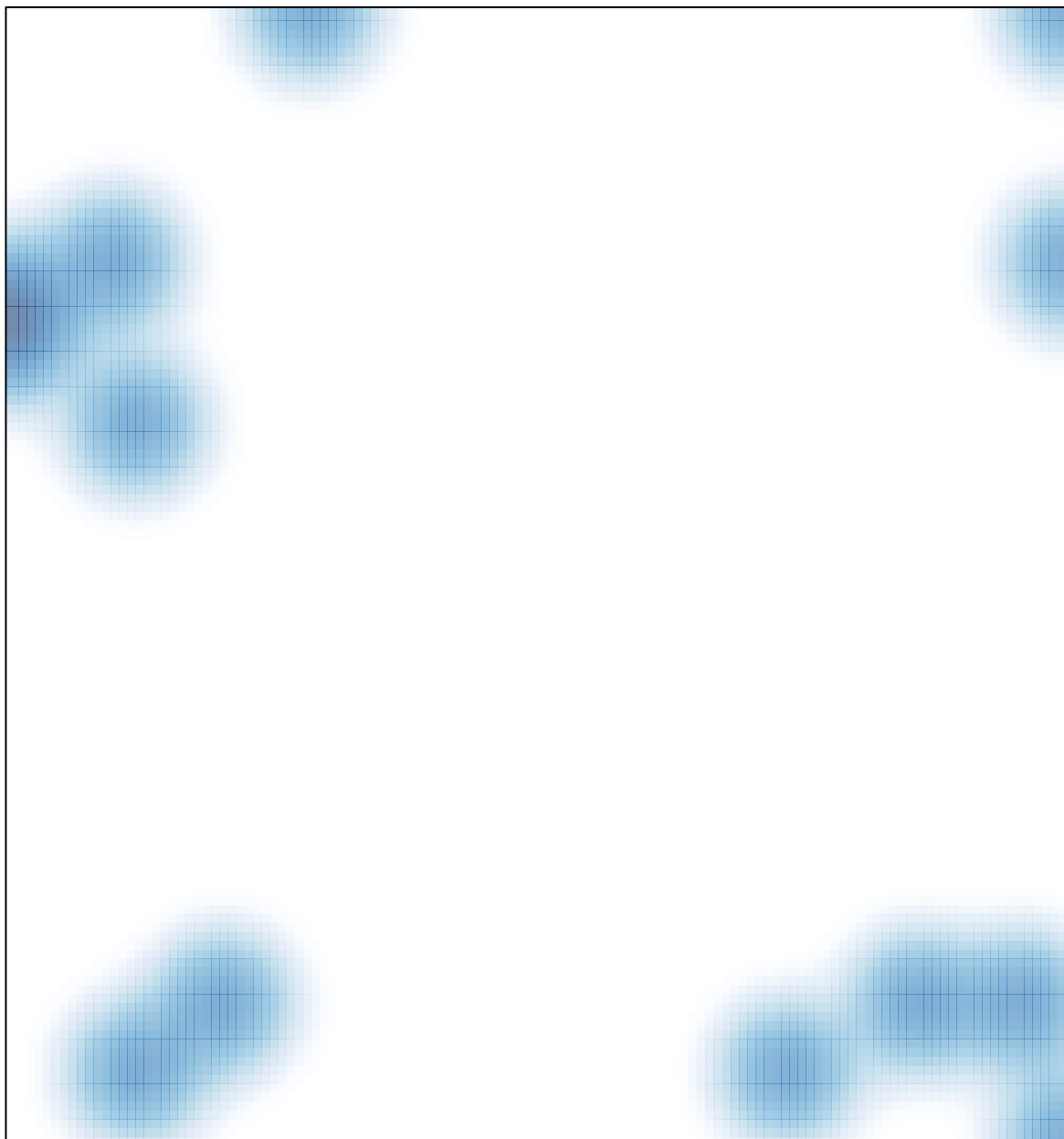


# phosphatidylinositol-3,4-bisphosphate binding



# features = 13  
chi-square p = 0.8

# phosphatidylinositol-3,4-bisphosphate binding



# features = 13 , max = 1