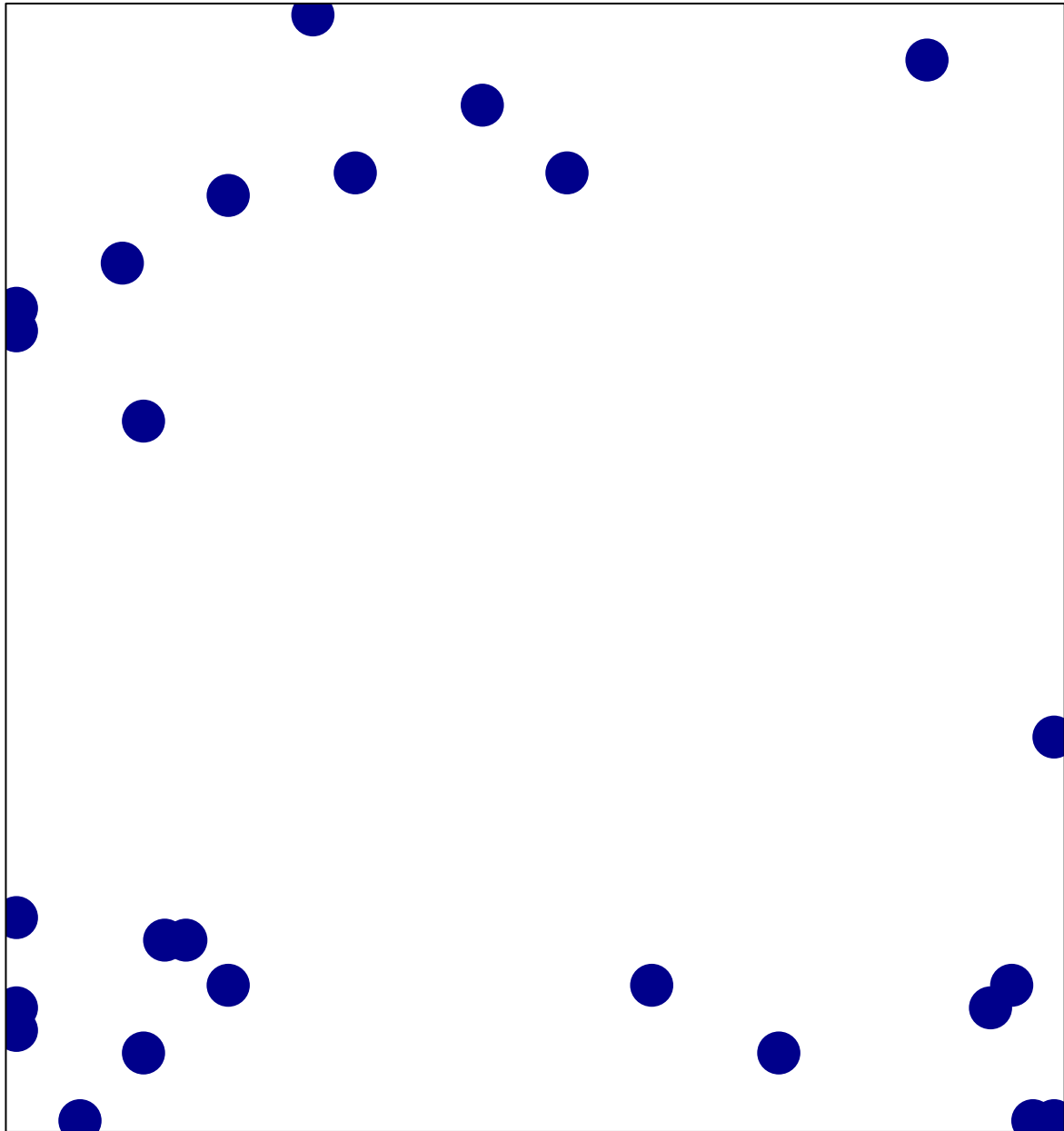
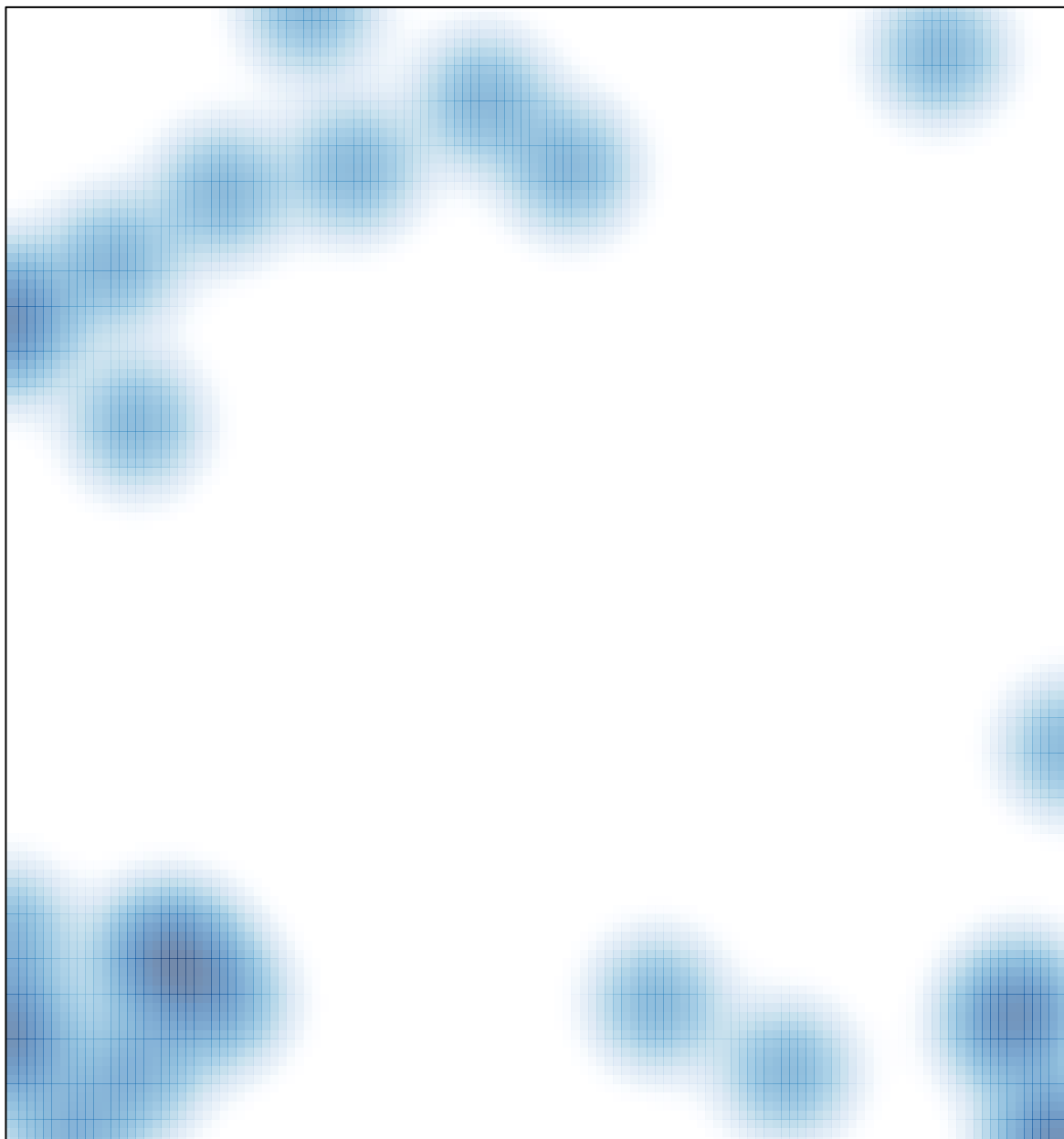


# phosphatidylinositol-3,4,5-trisphosphate binding



# features = 25  
chi-square p = 0.81

# phosphatidylinositol-3,4,5-trisphosphate binding



# features = 25 , max = 1