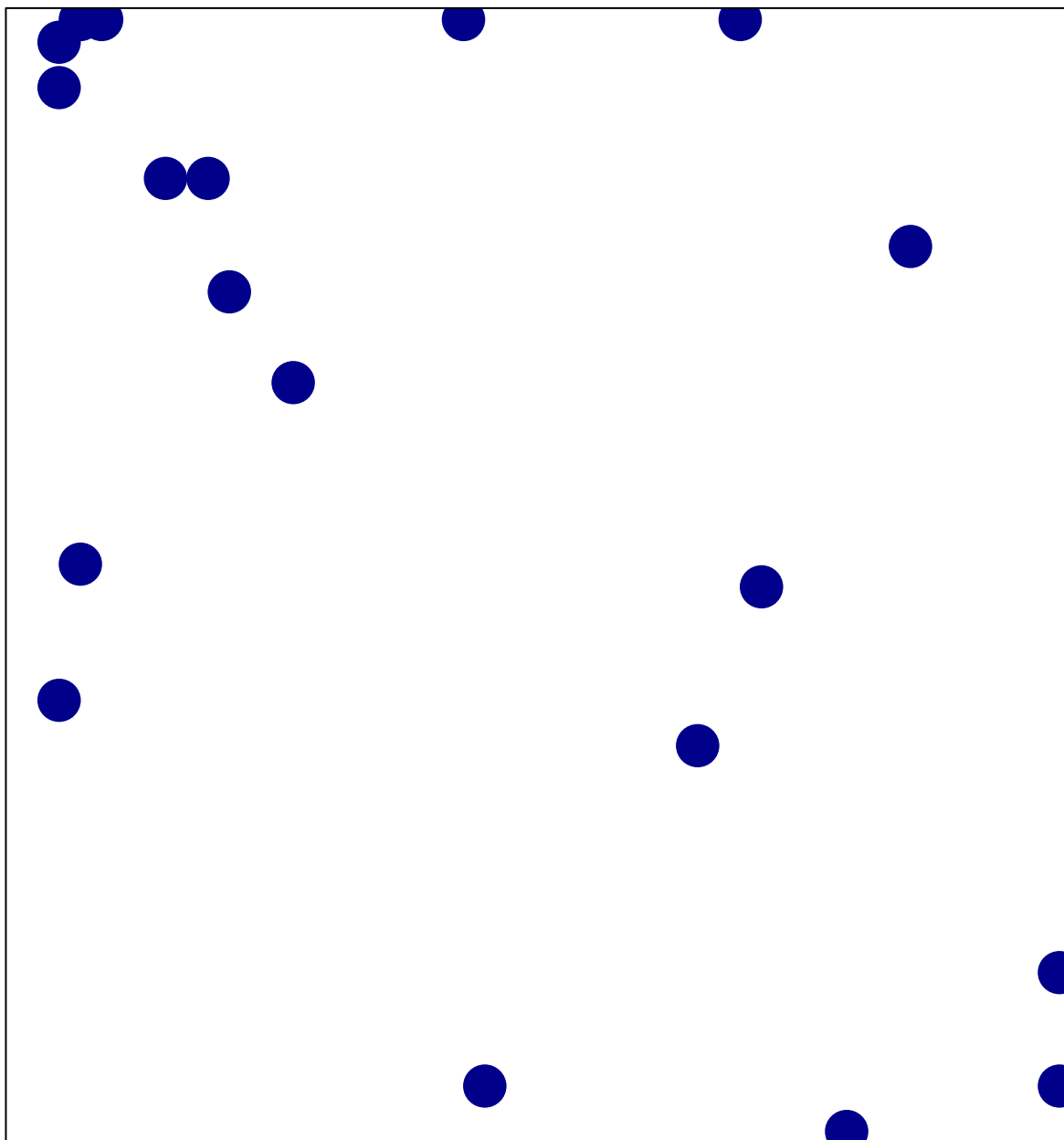
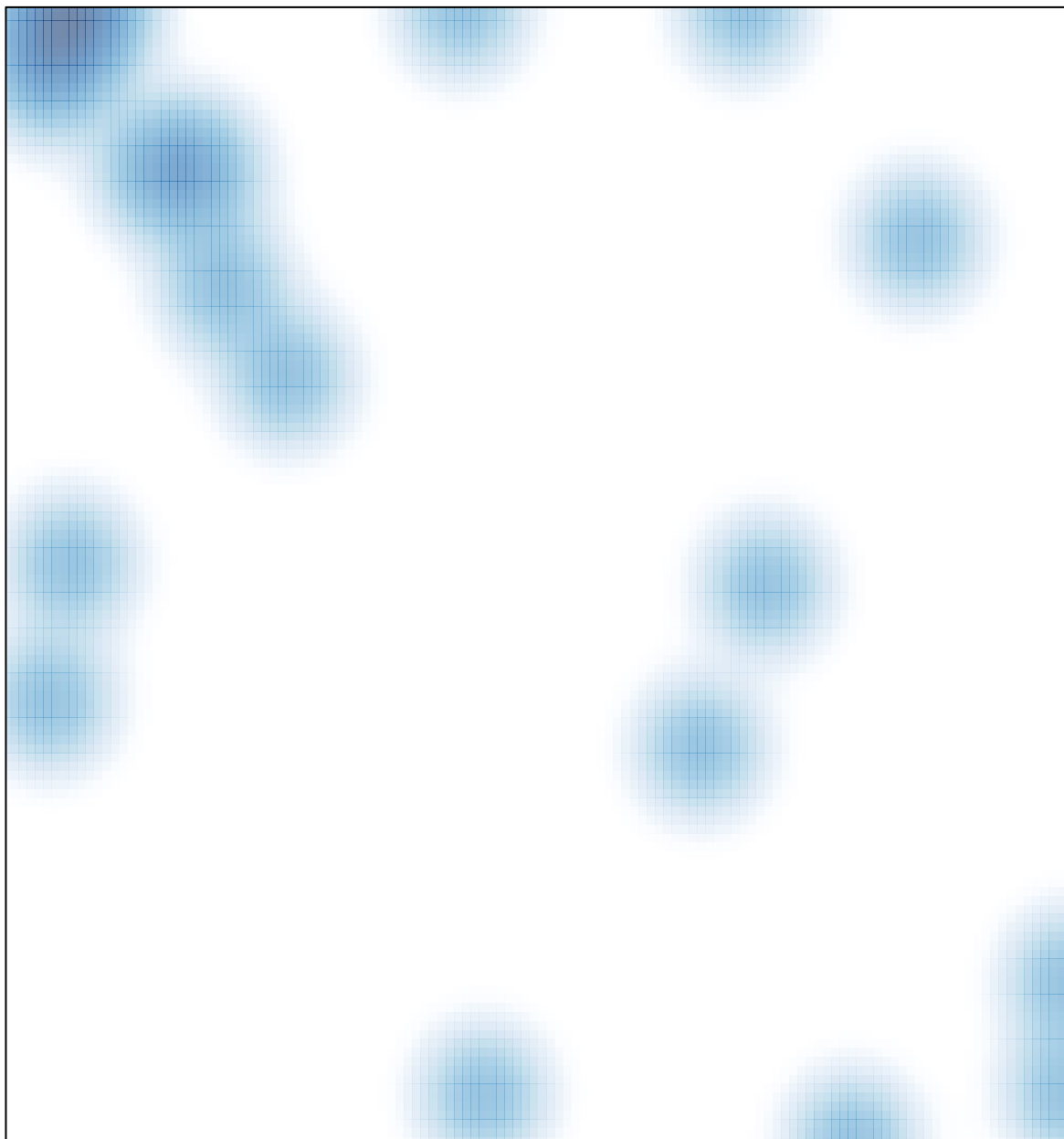


phosphatidylethanolamine acyl-chain remodeling



features = 19
chi-square p = 0.82

phosphatidylethanolamine acyl-chain remodeling



features = 19 , max = 1