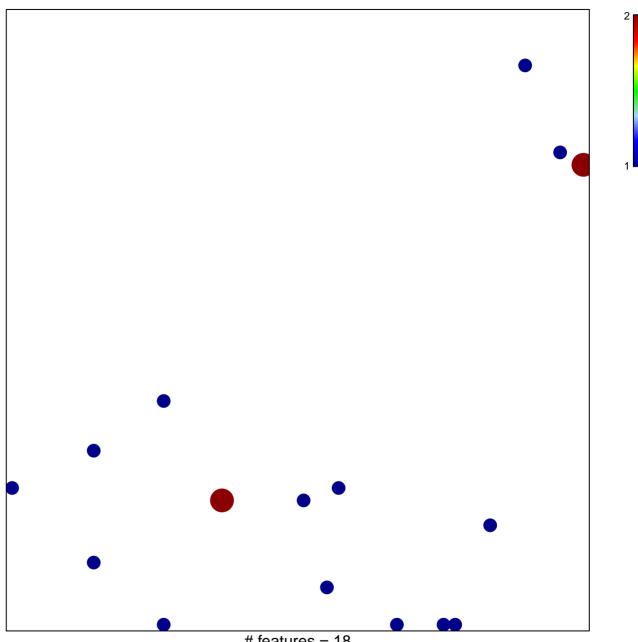
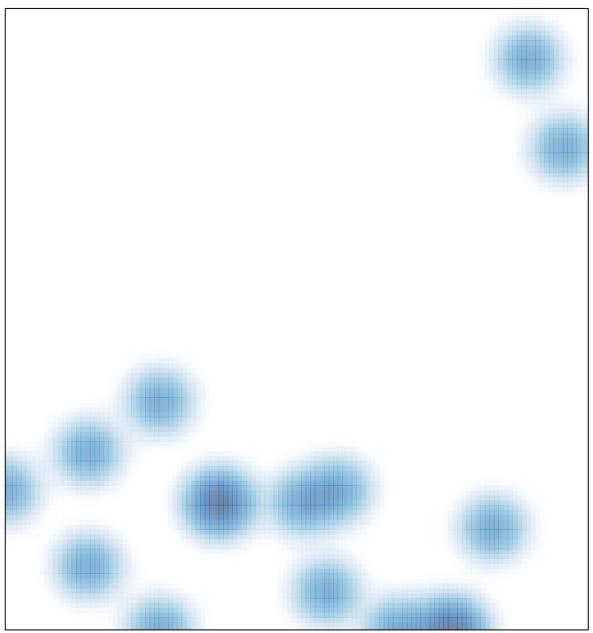
## phospholipid translocation



# features = 18 chi-square p = 0.81

## phospholipid translocation



# features = 18 , max = 2