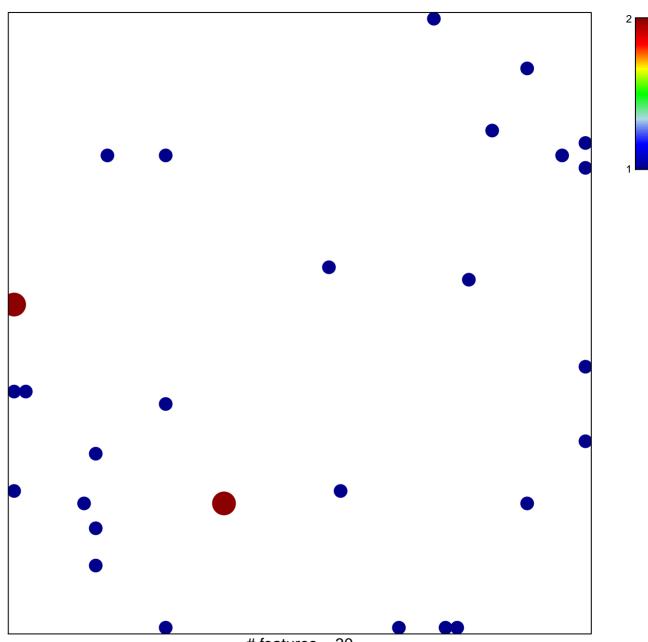
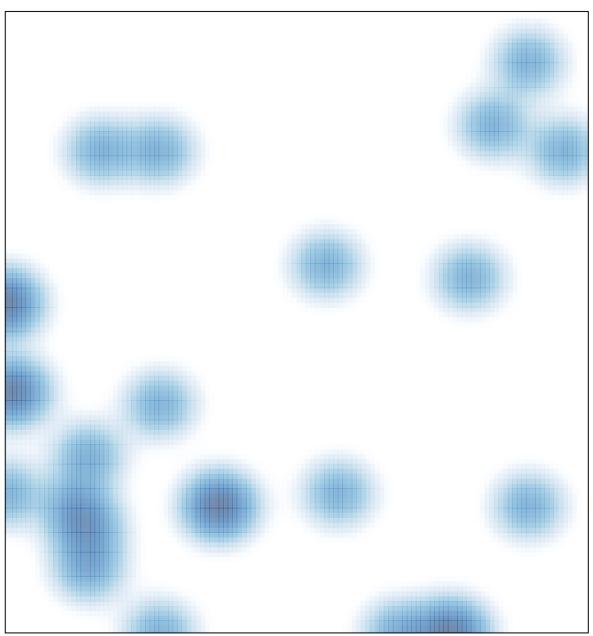
## phospholipid transport



# features = 30 chi-square p = 0.8

## phospholipid transport



# features = 30 , max = 2