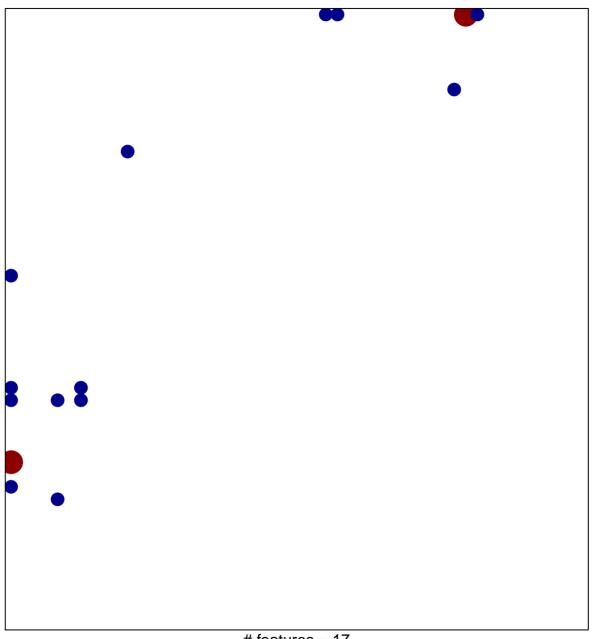
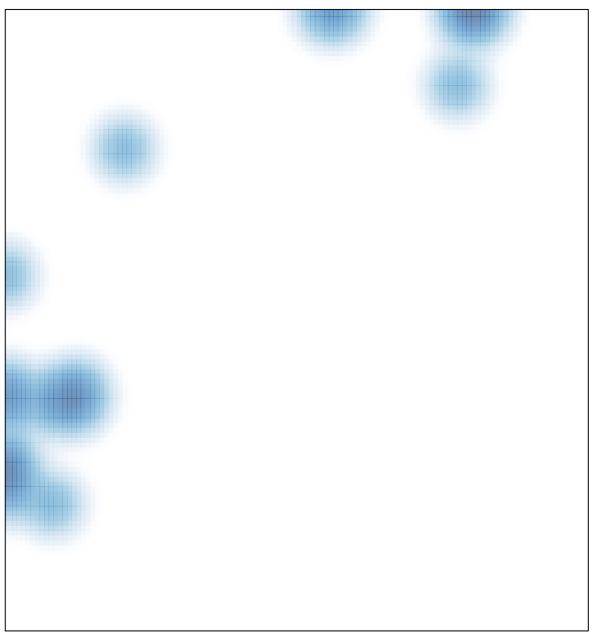
## proteasome accessory complex



# features = 17 chi-square p = 0.75

## proteasome accessory complex



# features = 17 , max = 2