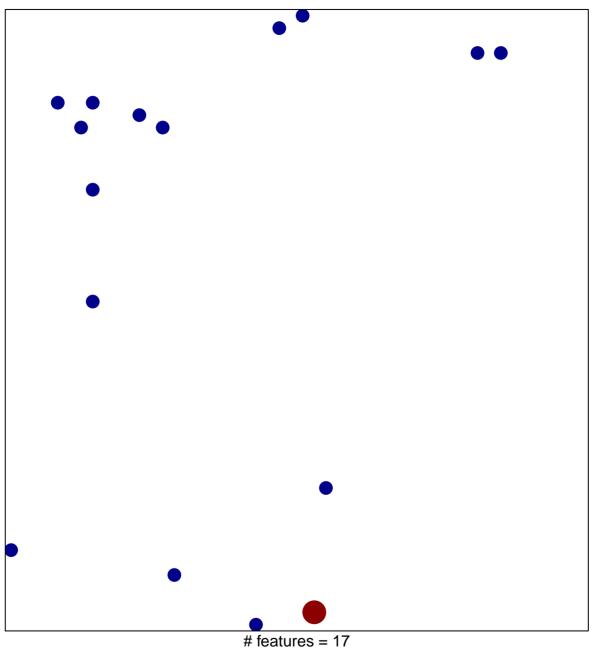
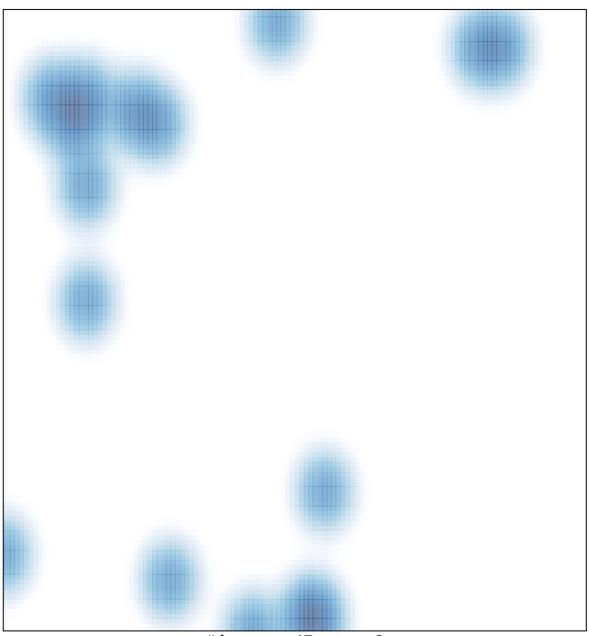
## SMN-Sm protein complex



# features = 17chi-square p = 0.83

## **SMN-Sm** protein complex



# features = 17 , max = 2