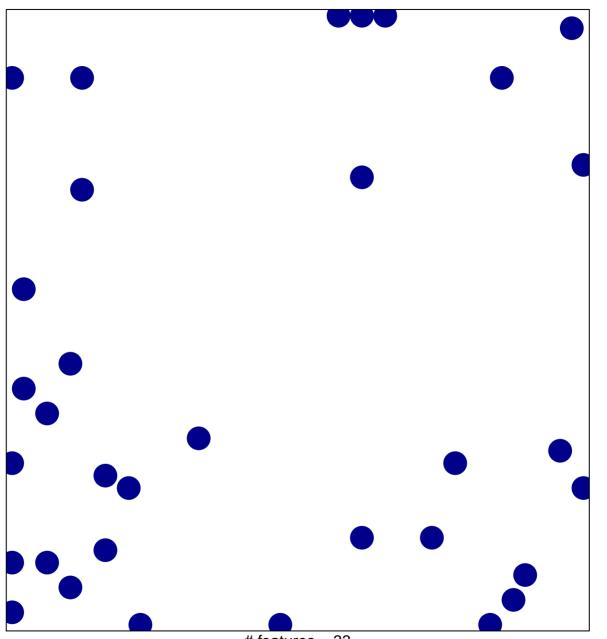
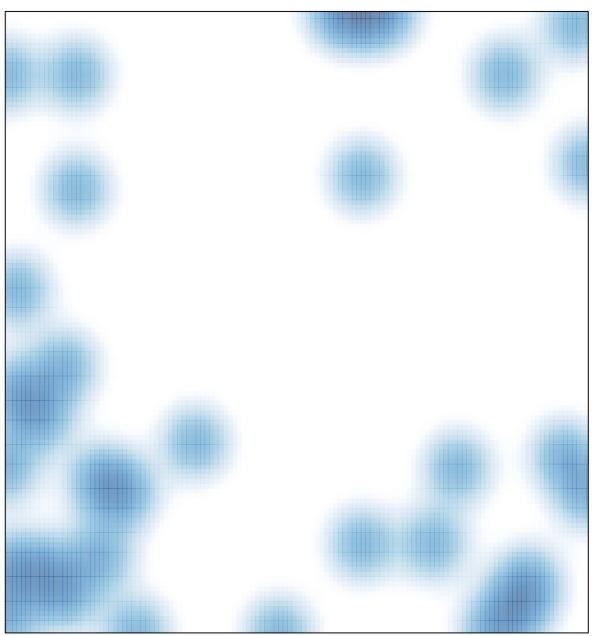
## **COP9** signalosome



# features = 33 chi–square p = 0.83

## **COP9** signalosome



# features = 33 , max = 1