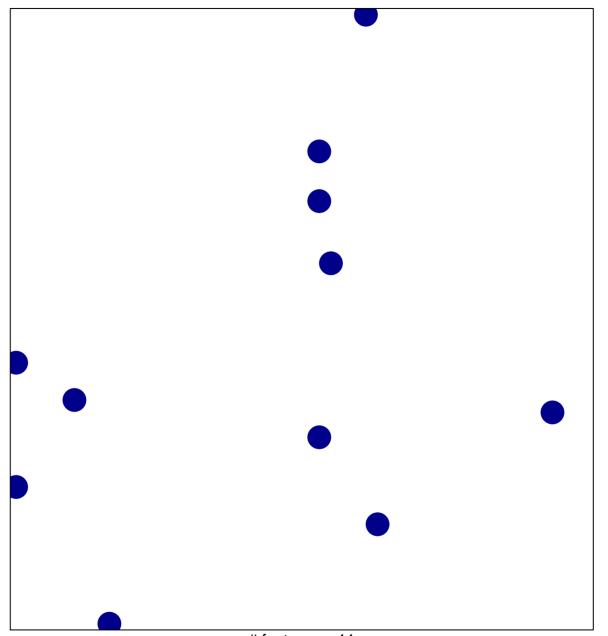
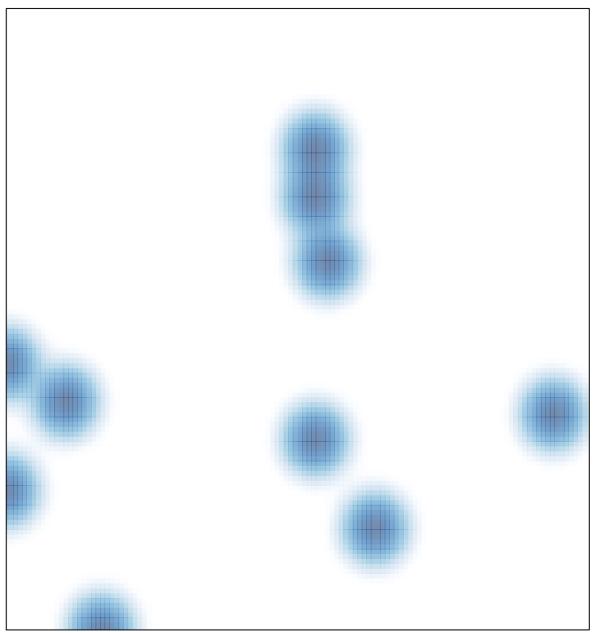
## REACTOME\_SIGNAL\_REGULATORY\_PROTEIN\_SIRP\_FAMILY\_INTERACTIONS



# features = 11 chi-square p = 0.83

## \_SIGNAL\_REGULATORY\_PROTEIN\_SIRP\_FAMILY\_IN



# features = 11 , max = 1