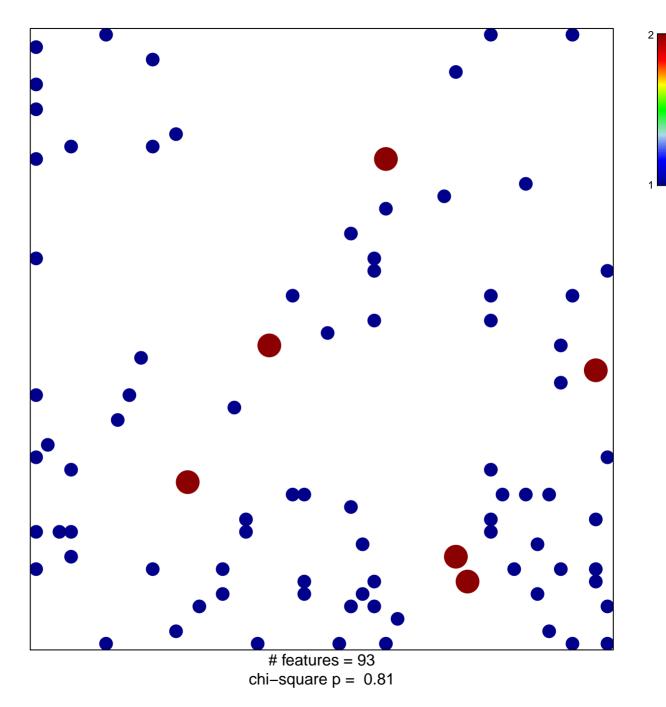
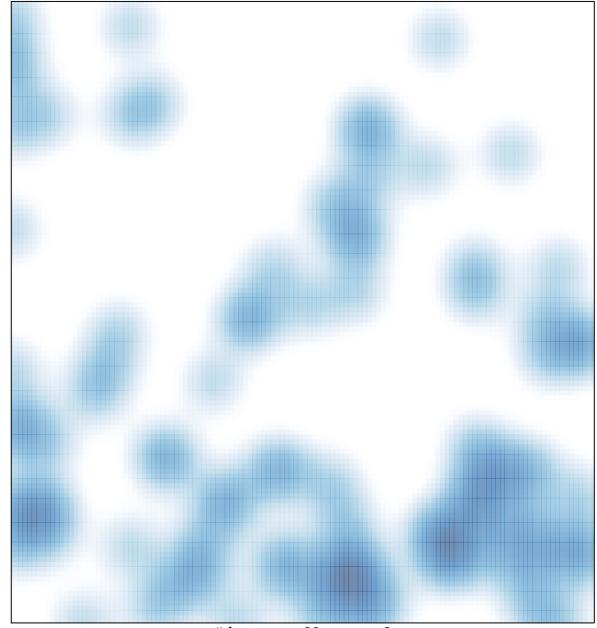
## REACTOME\_GLYCOSAMINOGLYCAN\_METABOLISM



## **REACTOME\_GLYCOSAMINOGLYCAN\_METABOLISM**



# features = 93 , max = 2