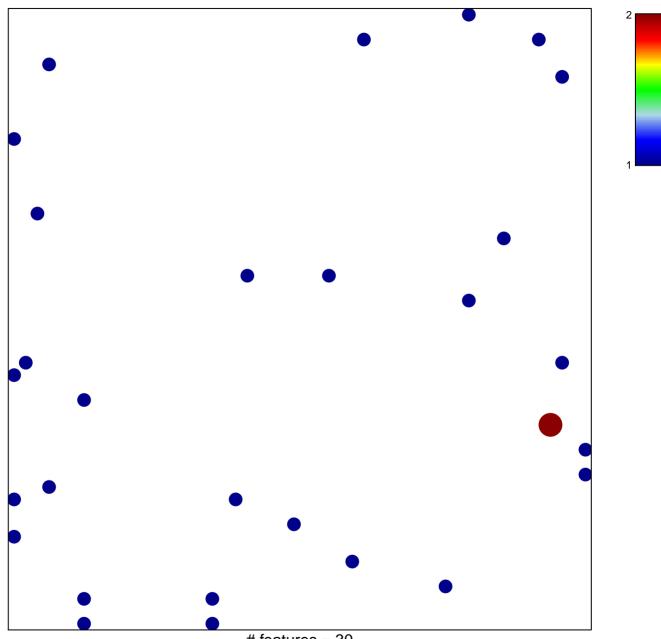
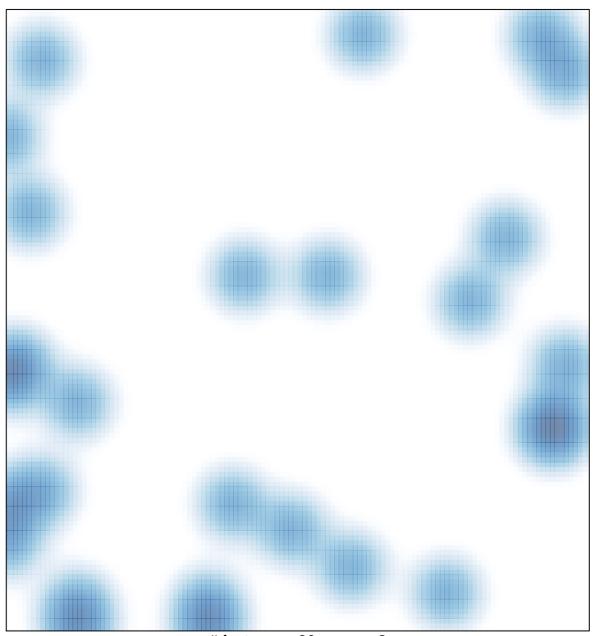
## REACTOME\_GLYCOSPHINGOLIPID\_METABOLISM



# features = 30 chi-square p = 0.83

## REACTOME\_GLYCOSPHINGOLIPID\_METABOLISM



# features = 30, max = 2