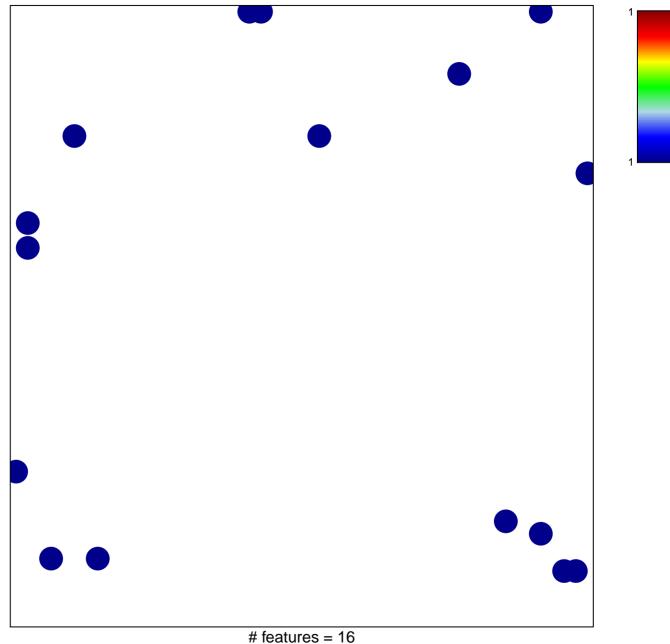
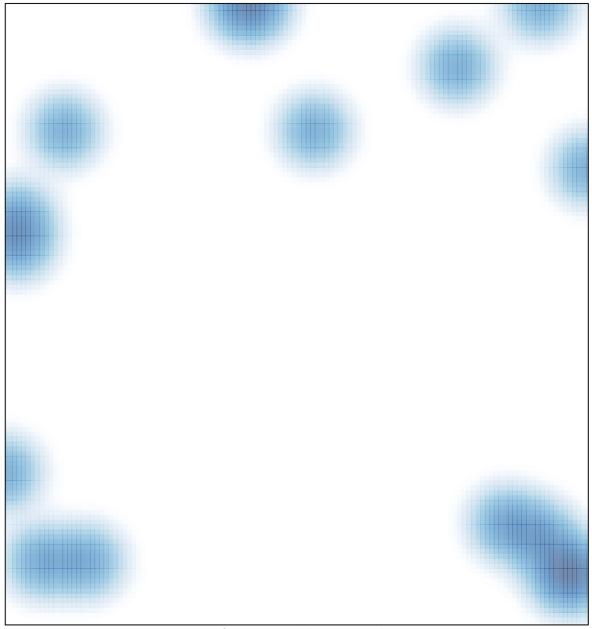
## **REACTOME\_UNFOLDED\_PROTEIN\_RESPONSE**



chi-square p = 0.83

## **REACTOME\_UNFOLDED\_PROTEIN\_RESPONSE**



# features = 16 , max = 1