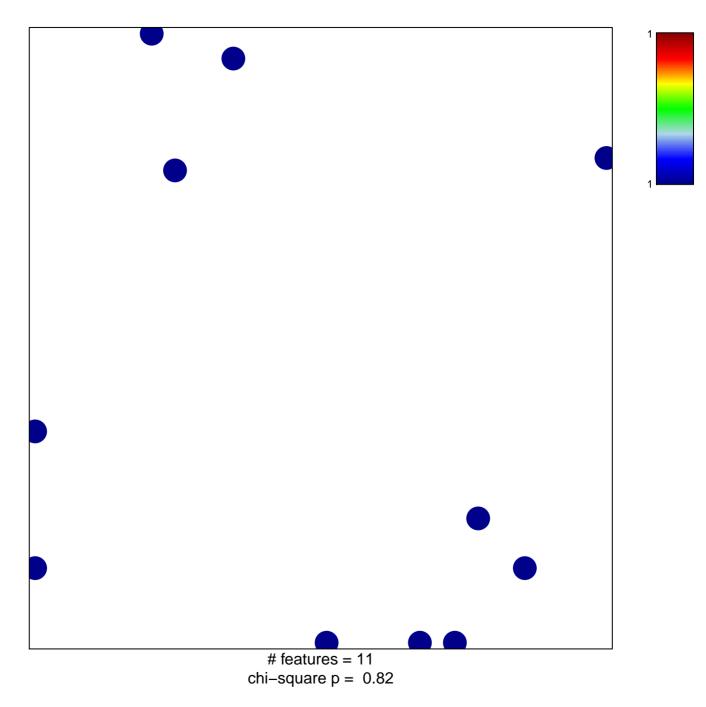
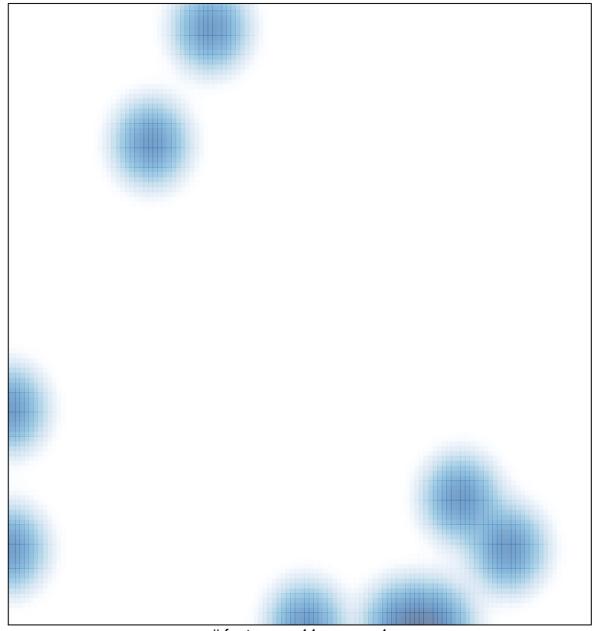
## REACTOME\_ALPHA\_LINOLENIC\_ACID\_ALA\_METABOLISM



## ACTOME\_ALPHA\_LINOLENIC\_ACID\_ALA\_METABOI



# features = 11, max = 1