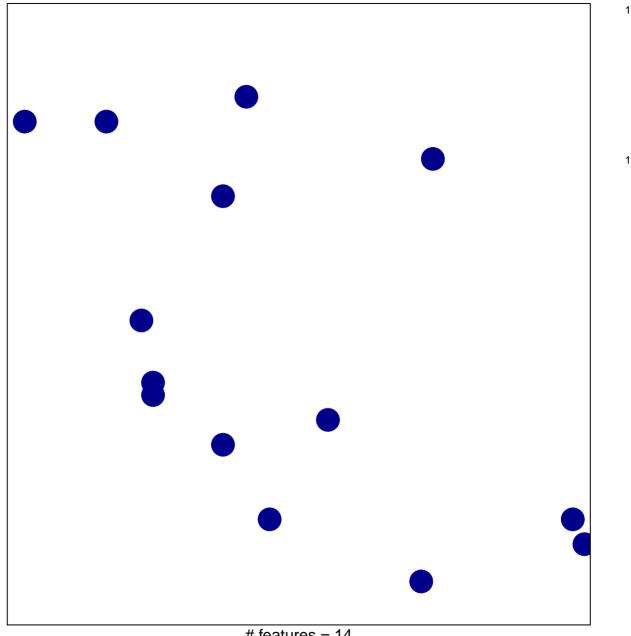
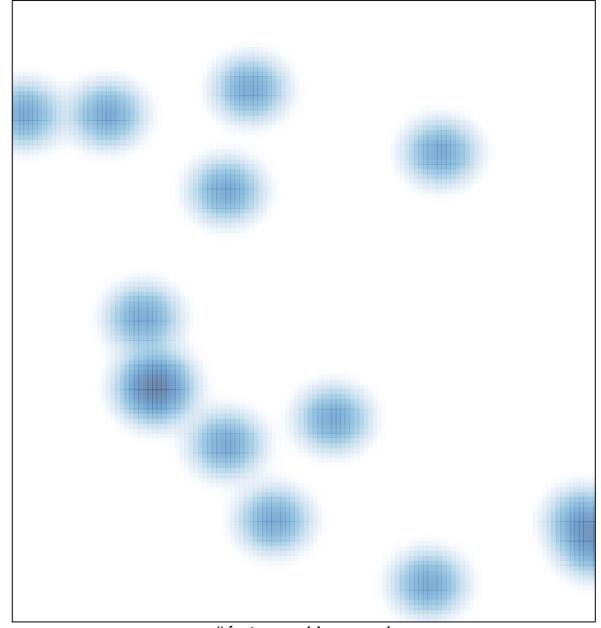
## **REACTOME\_METABOLISM\_OF\_BILE\_ACIDS\_AND\_BILE\_SALTS**



# features = 14chi-square p = 0.84

## TOME\_METABOLISM\_OF\_BILE\_ACIDS\_AND\_BILE\_



# features = 14 , max = 1