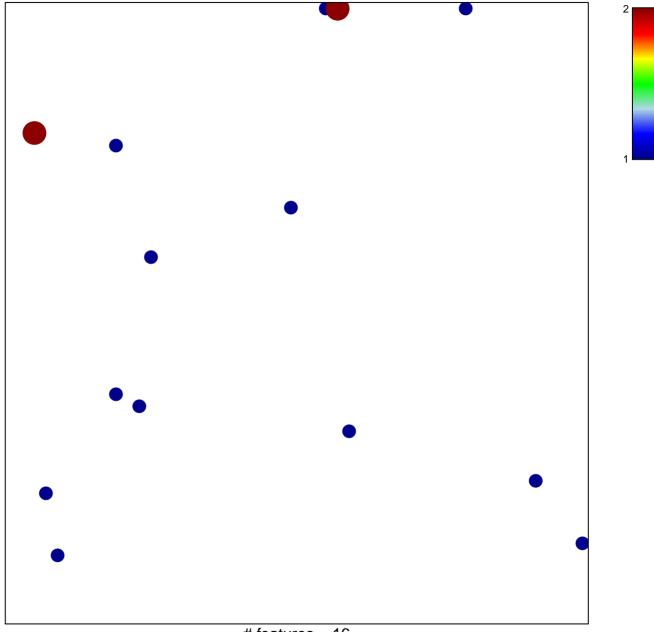
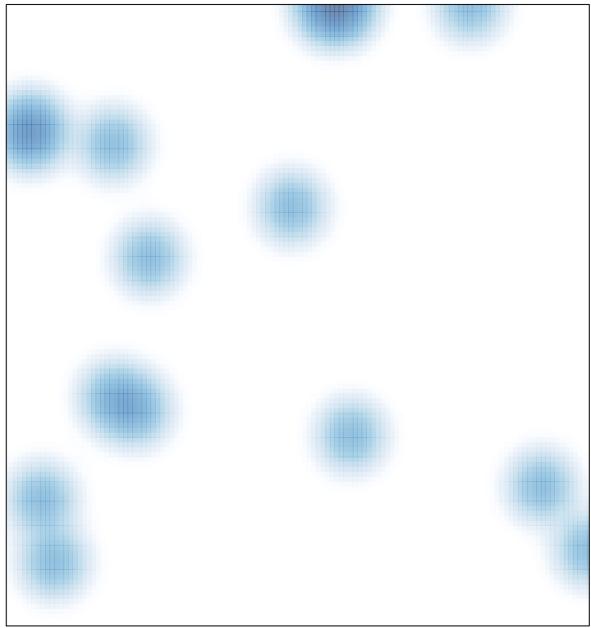
## **REACTOME\_PYRUVATE\_METABOLISM**



# features = 16
chi-square p = 0.82

## **REACTOME\_PYRUVATE\_METABOLISM**



# features = 16, max = 2