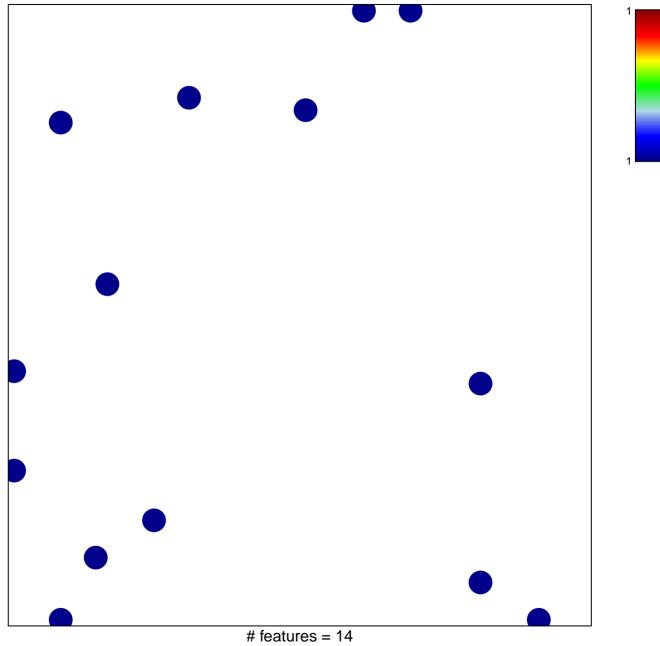
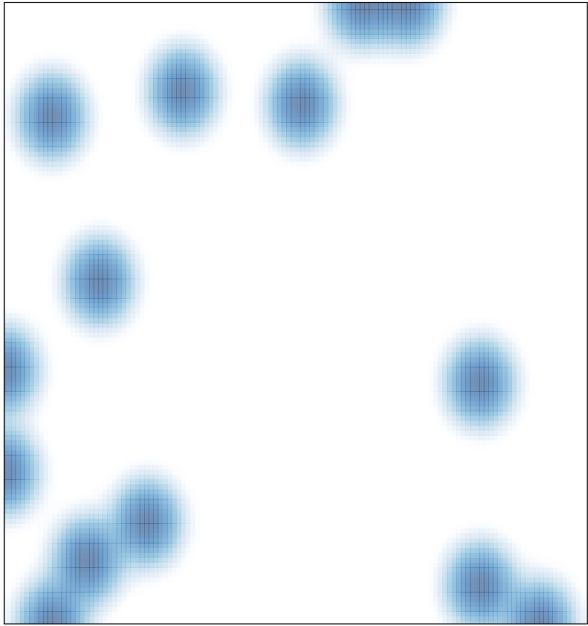
## **REACTOME\_OPIOID\_SIGNALLING**



chi-square p = 0.84

## **REACTOME\_OPIOID\_SIGNALLING**



# features = 14, max = 1