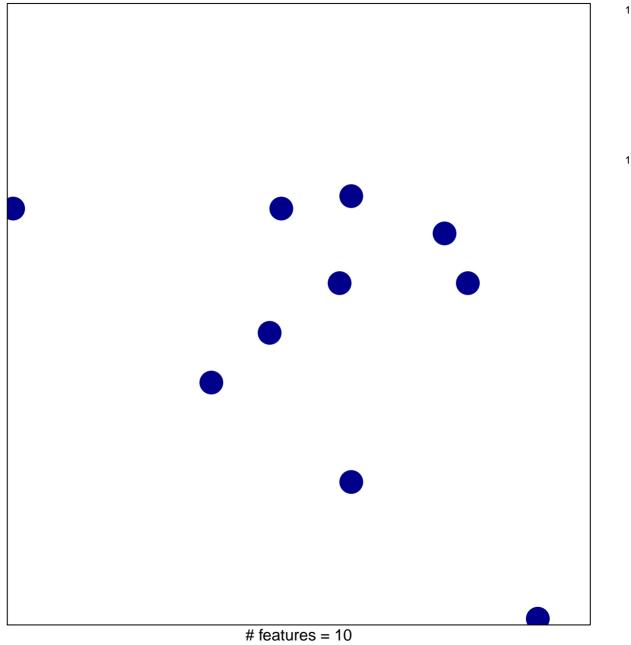
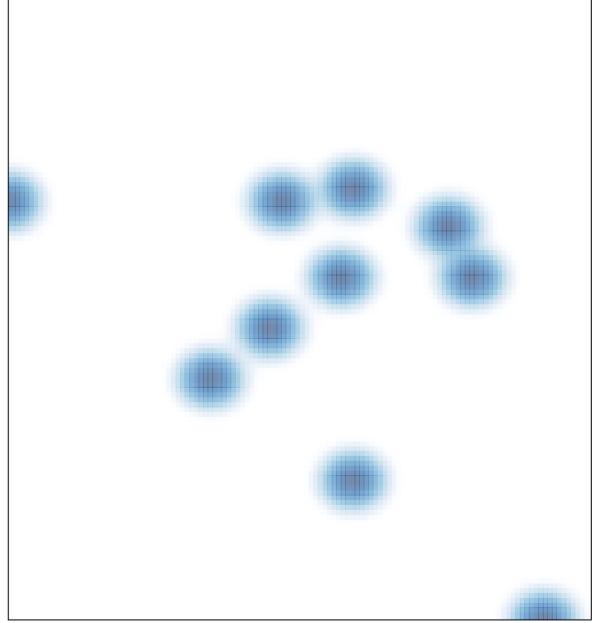
## **REACTOME\_GAP\_JUNCTION\_ASSEMBLY**



chi-square p = 0.83

## **REACTOME\_GAP\_JUNCTION\_ASSEMBLY**



# features = 10, max = 1