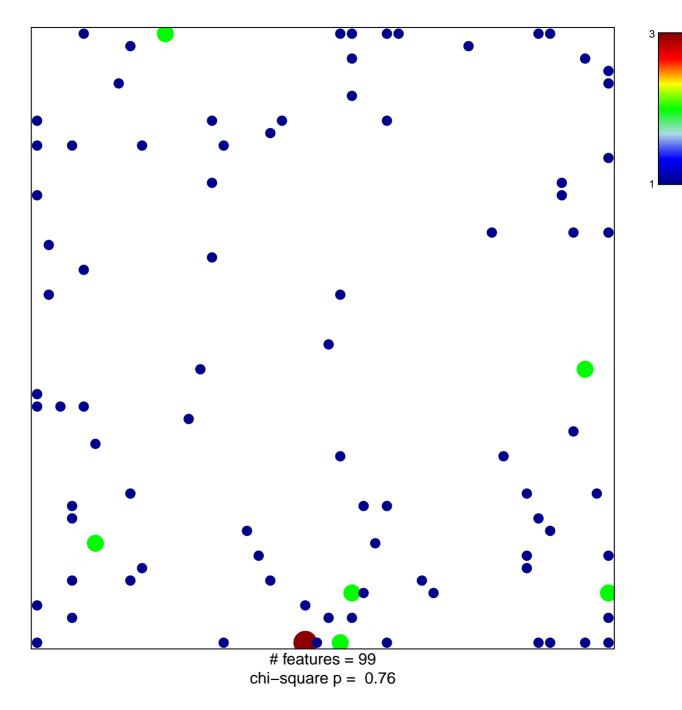
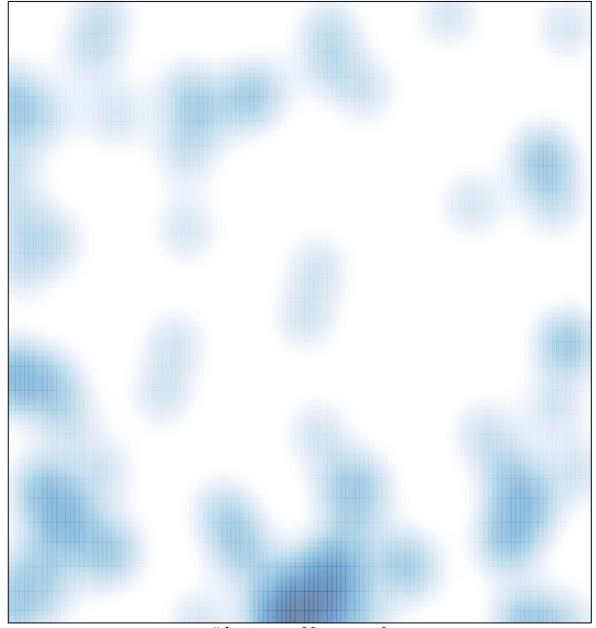
## **REACTOME\_SIGNALING\_BY\_EGFR\_IN\_CANCER**



## **REACTOME\_SIGNALING\_BY\_EGFR\_IN\_CANCER**



# features = 99 , max = 3