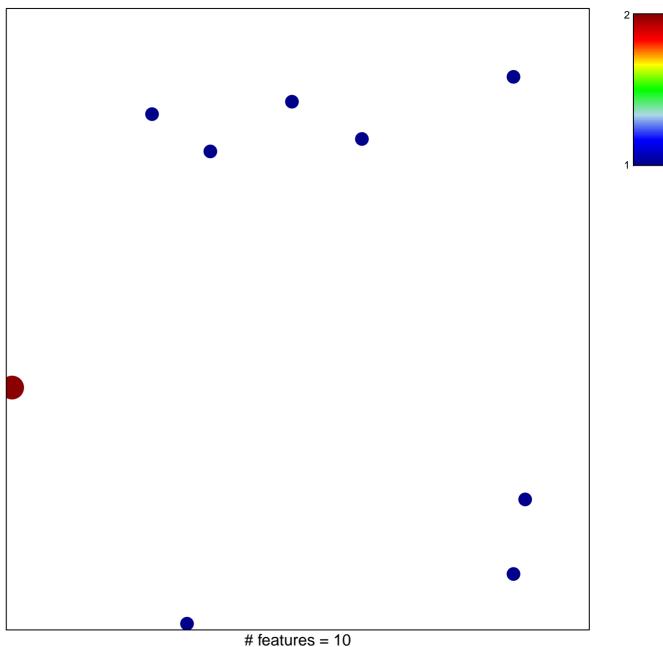
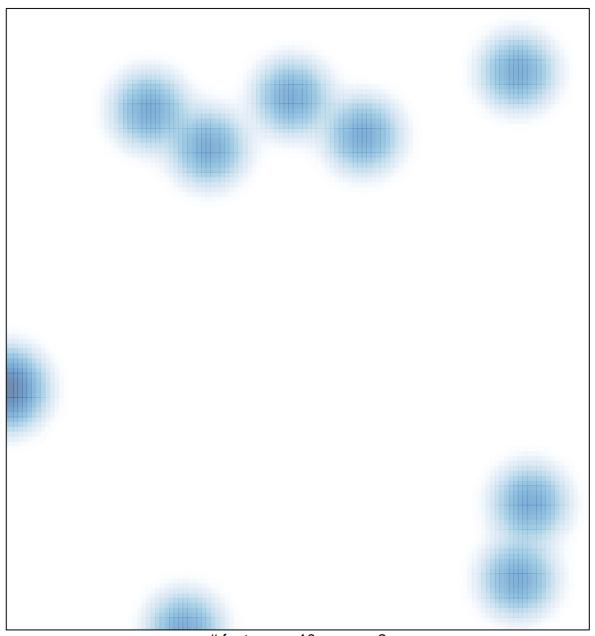
## REACTOME\_CHYLOMICRON\_MEDIATED\_LIPID\_TRANSPORT



# features = 10chi-square p = 0.83

## CTOME\_CHYLOMICRON\_MEDIATED\_LIPID\_TRANSI



# features = 10 , max = 2