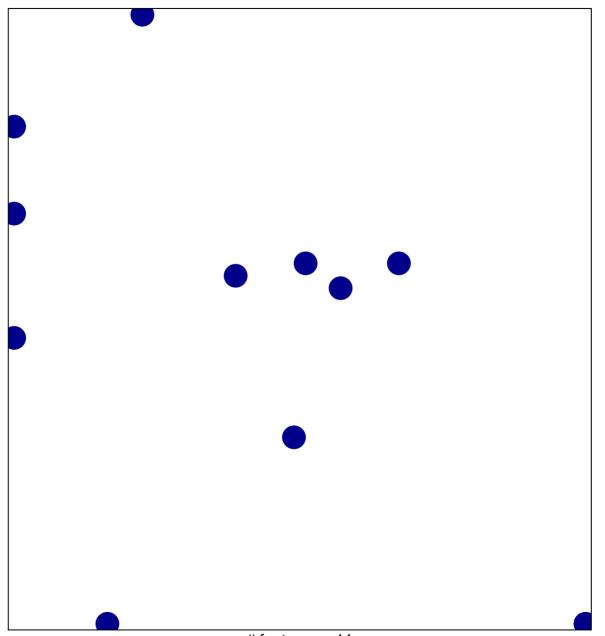
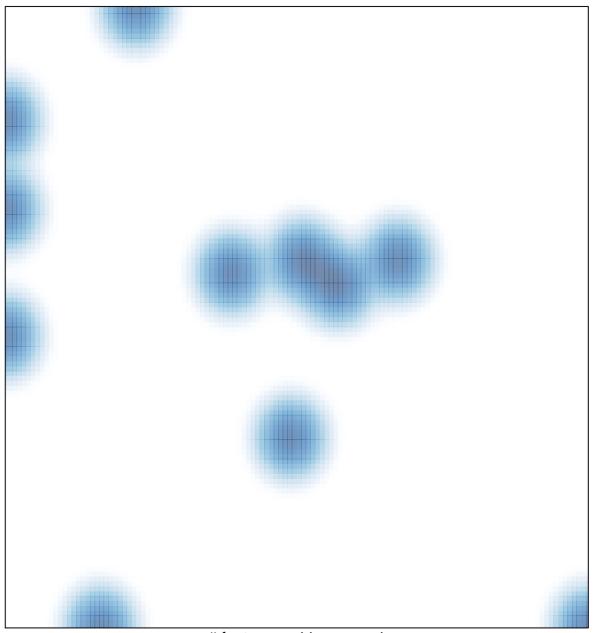
## REACTOME\_CHYLOMICRON\_MEDIATED\_LIPID\_TRANSPORT



# features = 11 chi-square p = 0.82

## CTOME\_CHYLOMICRON\_MEDIATED\_LIPID\_TRANSI



# features = 11 , max = 1