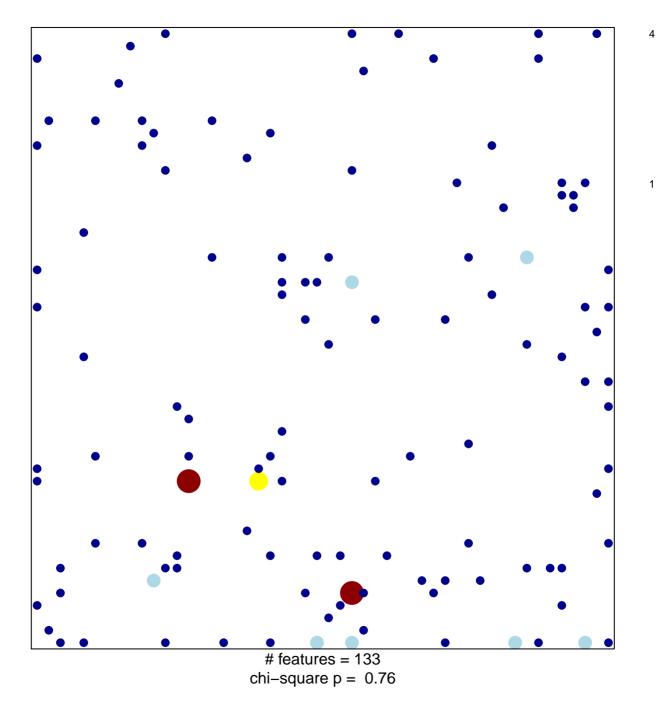
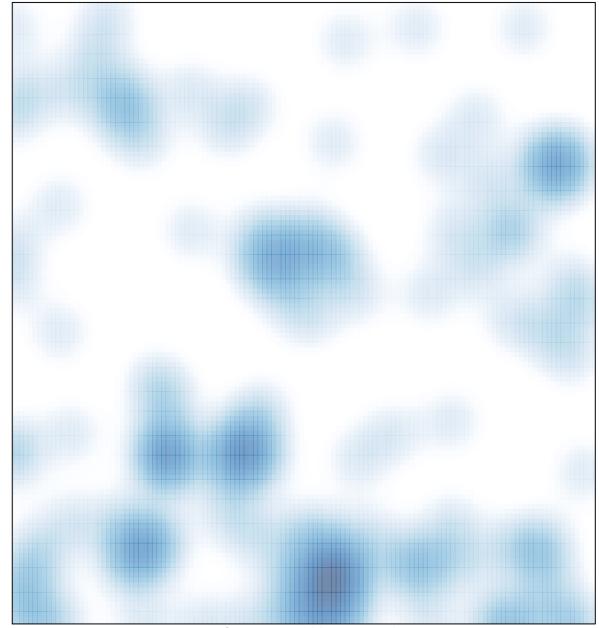
## **REACTOME\_TRANSMISSION\_ACROSS\_CHEMICAL\_SYNAPSES**



## CTOME\_TRANSMISSION\_ACROSS\_CHEMICAL\_SYN/



# features = 133 , max = 4