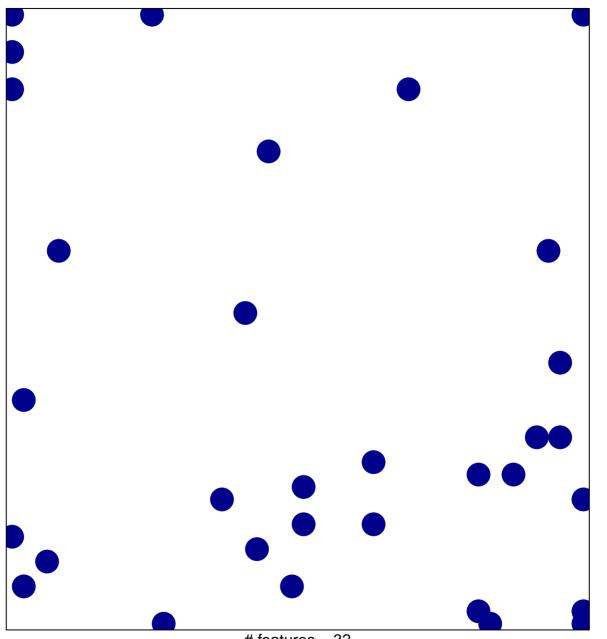
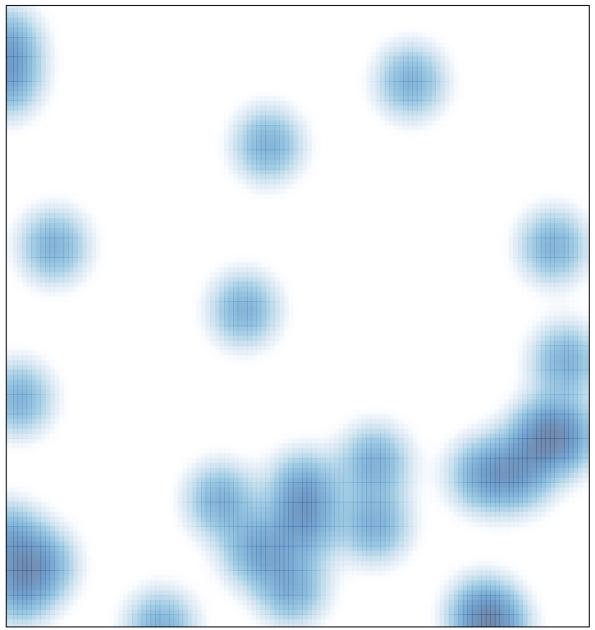
## REACTOME\_APOPTOTIC\_CLEAVAGE\_OF\_CELLULAR\_PROTEINS



# features = 32 chi–square p = 0.82

## TOME\_APOPTOTIC\_CLEAVAGE\_OF\_CELLULAR\_PR



# features = 32, max = 1