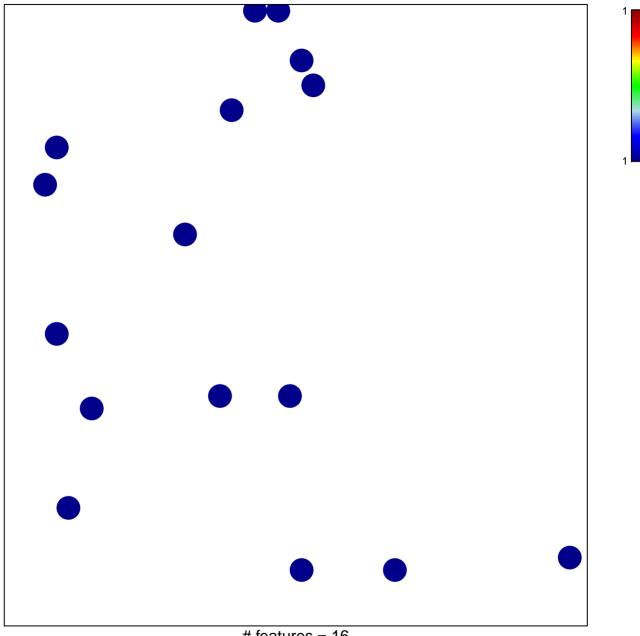
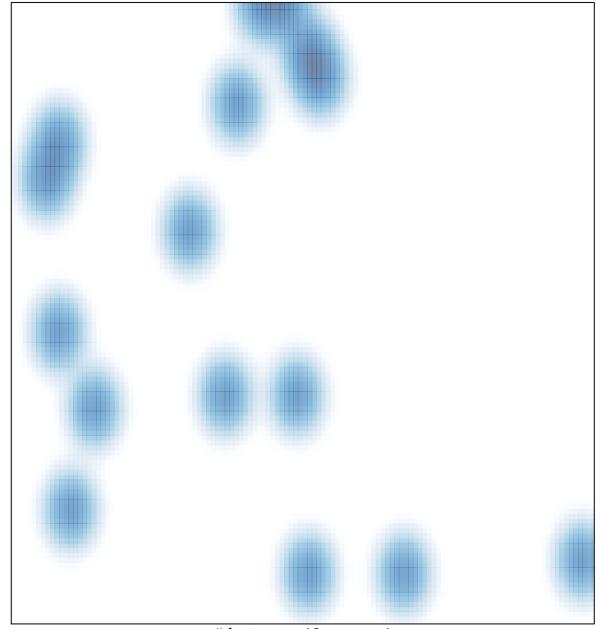
## REACTOME\_MAP\_KINASES\_ACTIVATION\_IN\_TLR\_CASCADE



# features = 16chi-square p = 0.84

## CTOME\_MAP\_KINASES\_ACTIVATION\_IN\_TLR\_CAS



# features = 16, max = 1