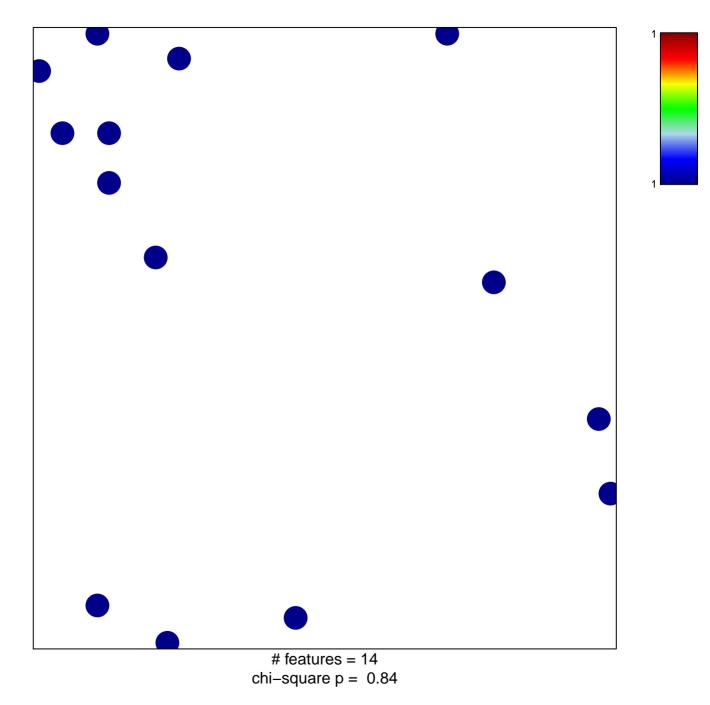
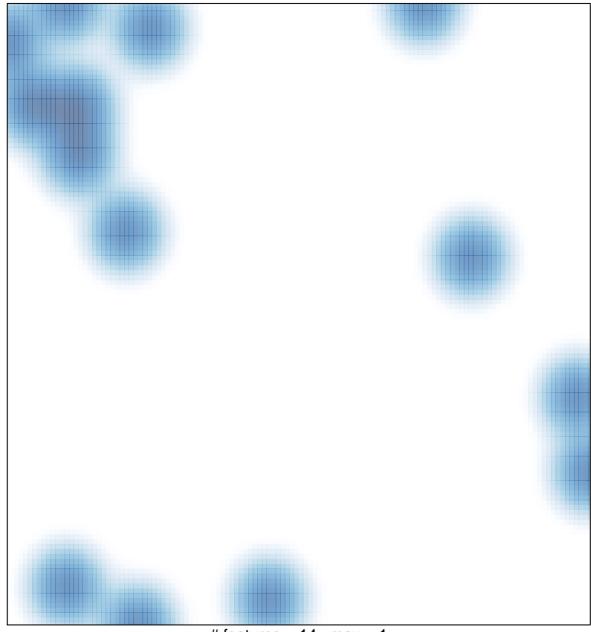
ZHAN\_MULTIPLE\_MYELOMA\_CD2\_DN



## ZHAN\_MULTIPLE\_MYELOMA\_CD2\_DN



# features = 14, max = 1