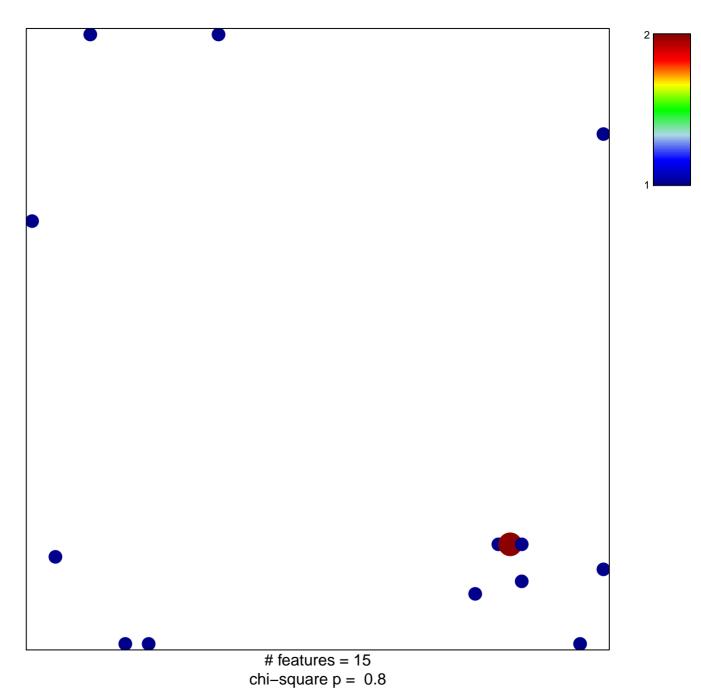
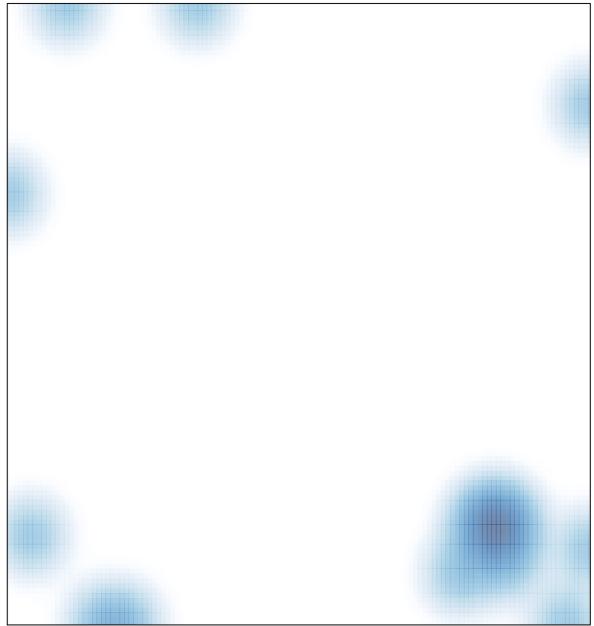
ZHAN\_MULTIPLE\_MYELOMA\_CD2\_UP



## ZHAN\_MULTIPLE\_MYELOMA\_CD2\_UP



# features = 15, max = 2