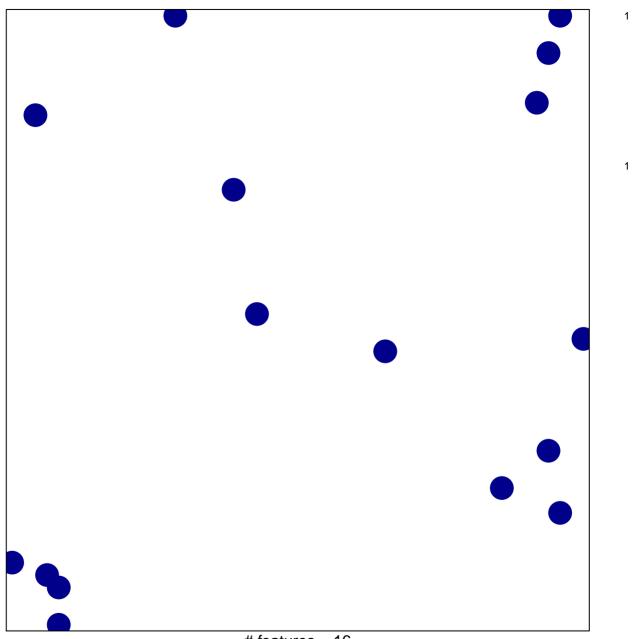
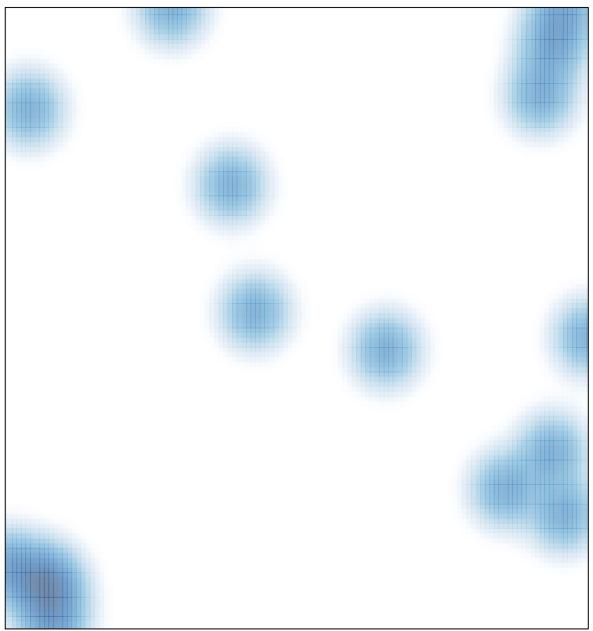
## ZHAN\_MULTIPLE\_MYELOMA\_MS\_UP



# features = 16 chi-square p = 0.85

## ZHAN\_MULTIPLE\_MYELOMA\_MS\_UP



# features = 16 , max = 1