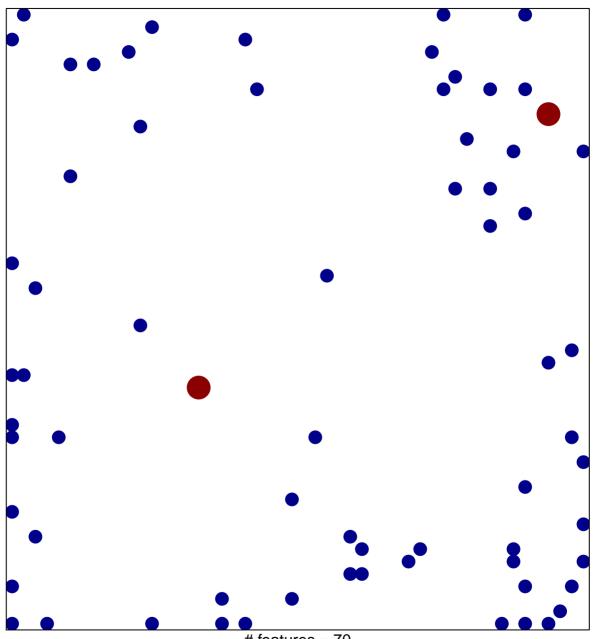
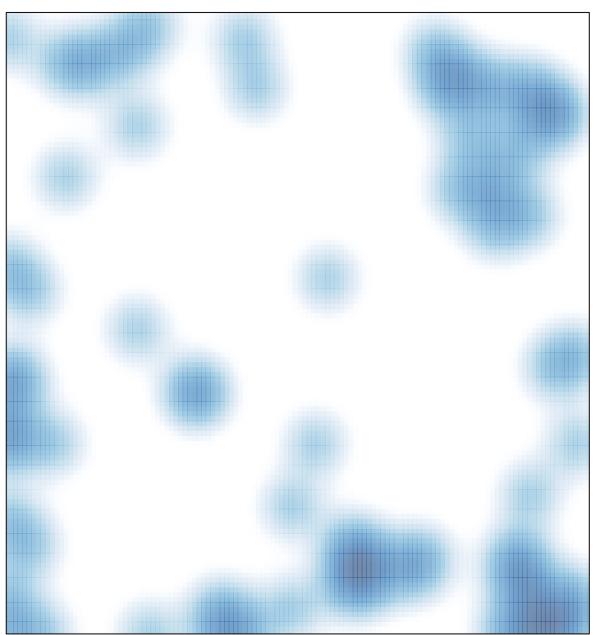
## hsa-miR-375



# features = 70 chi-square p = 0.8

## hsa-miR-375



# features = 70 , max = 2