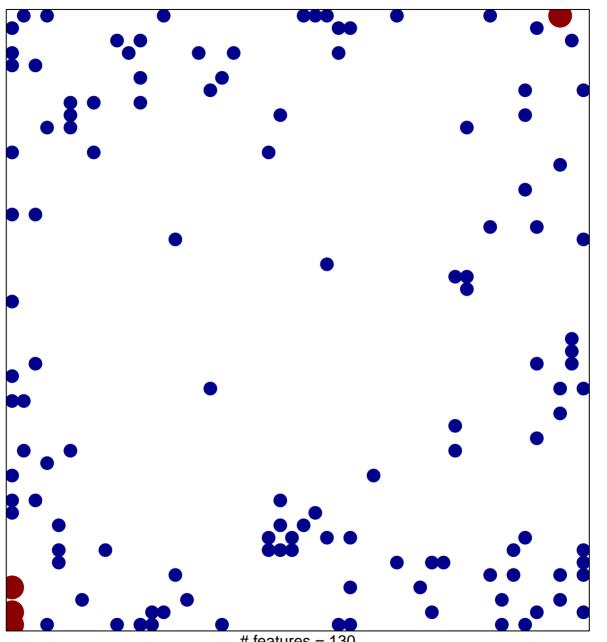
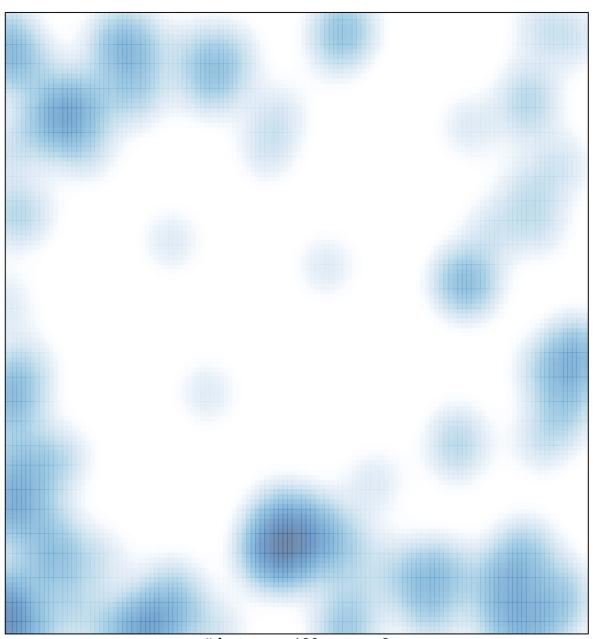
## hsa-miR-145



# features = 130 chi-square p = 0.76

## hsa-miR-145



# features = 130 , max = 2