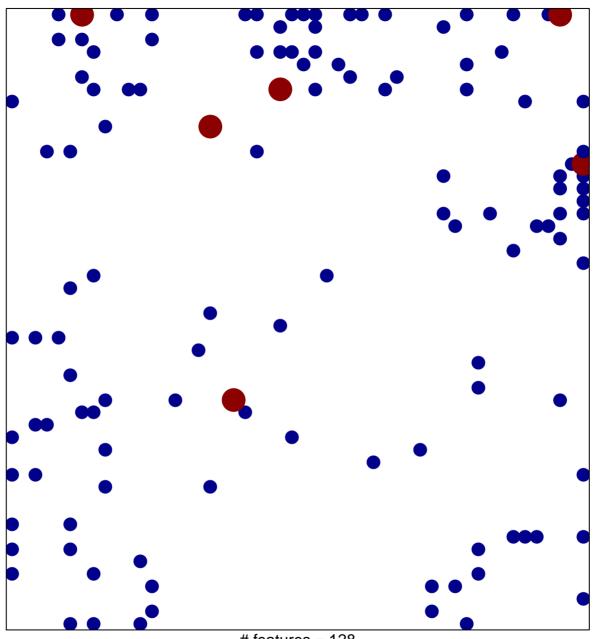
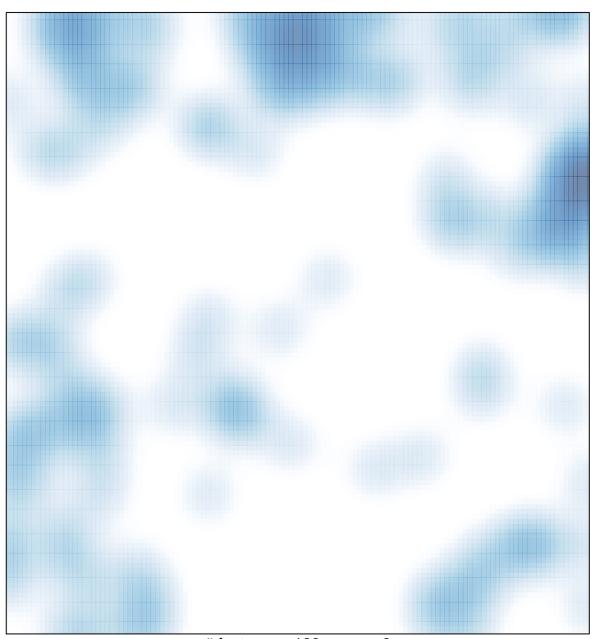
## hsa-miR-223



# features = 128 chi-square p = 0.76

## hsa-miR-223



# features = 128 , max = 2