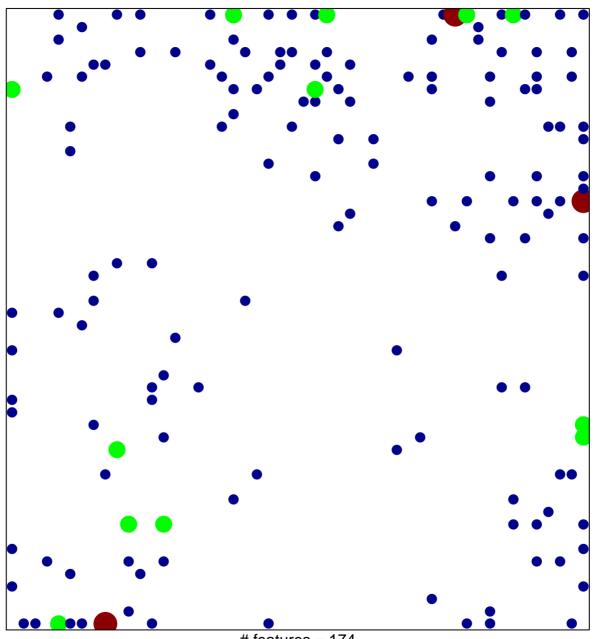
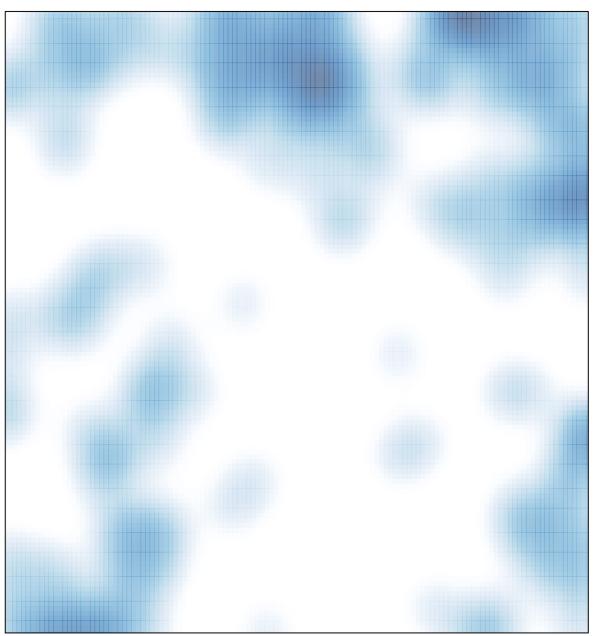
## hsa-miR-196b



# features = 174 chi-square p = 0.71

## hsa-miR-196b



# features = 174 , max = 3