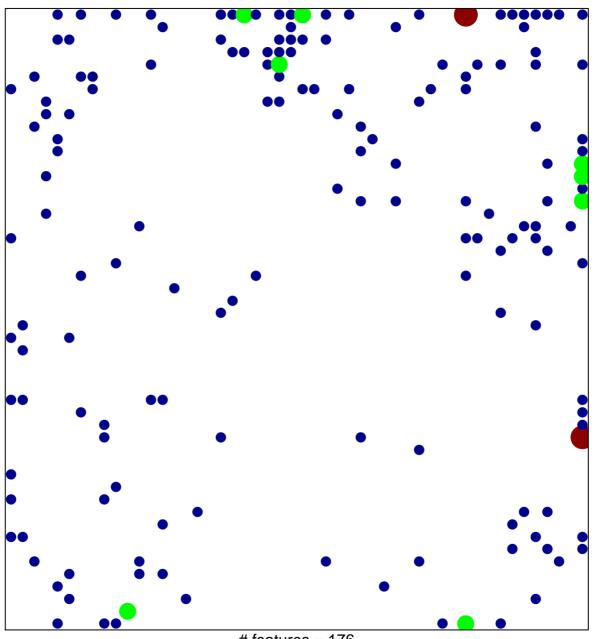
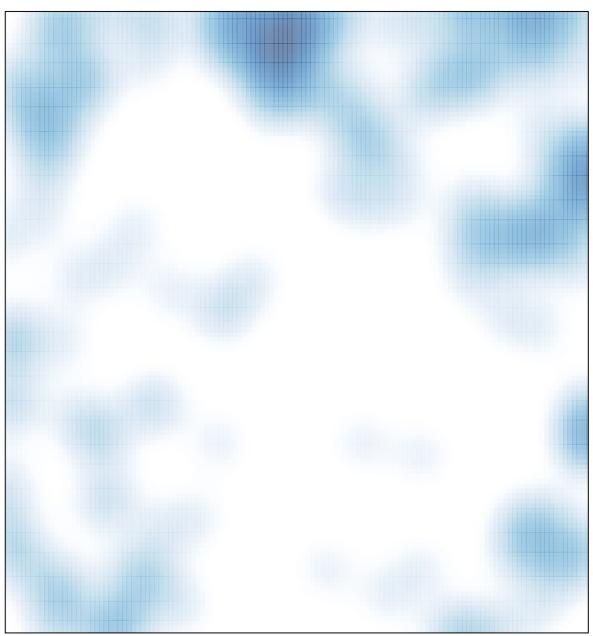
## hsa-miR-23b



# features = 176 chi-square p = 0.71

## hsa-miR-23b



# features = 176 , max = 3