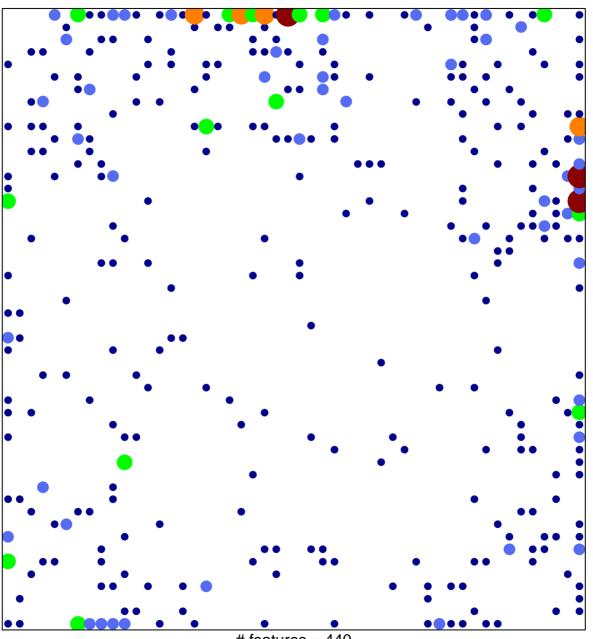
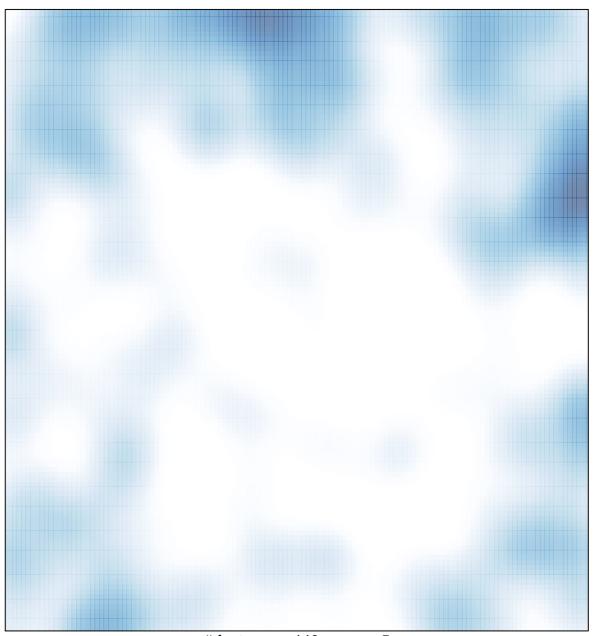
## hsa-miR-19a



# features = 440 chi-square p = 0.57

## hsa-miR-19a



# features = 440 , max = 5