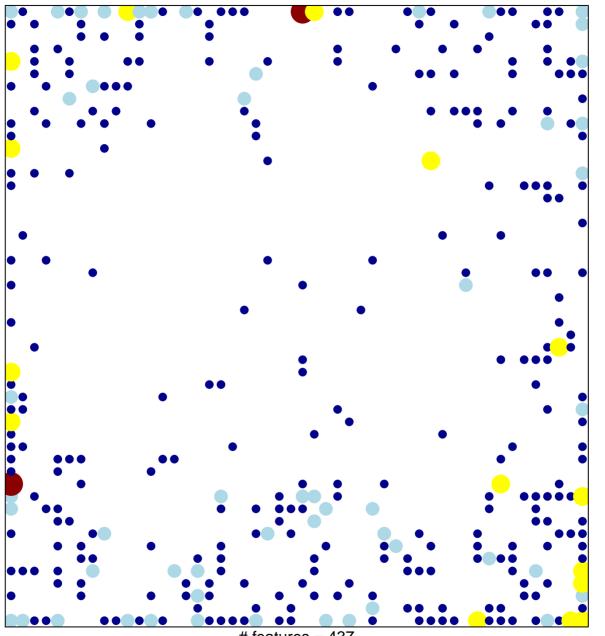
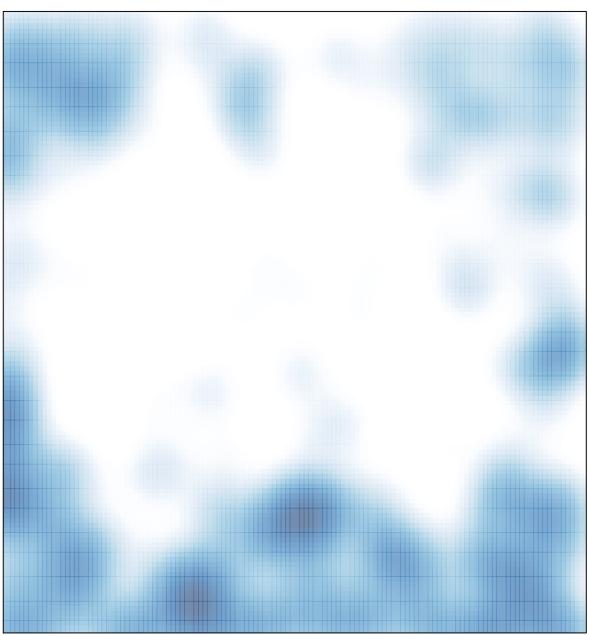
## hsa-miR-19b



# features = 437 chi-square p = 0.56

## hsa-miR-19b



# features = 437 , max = 4