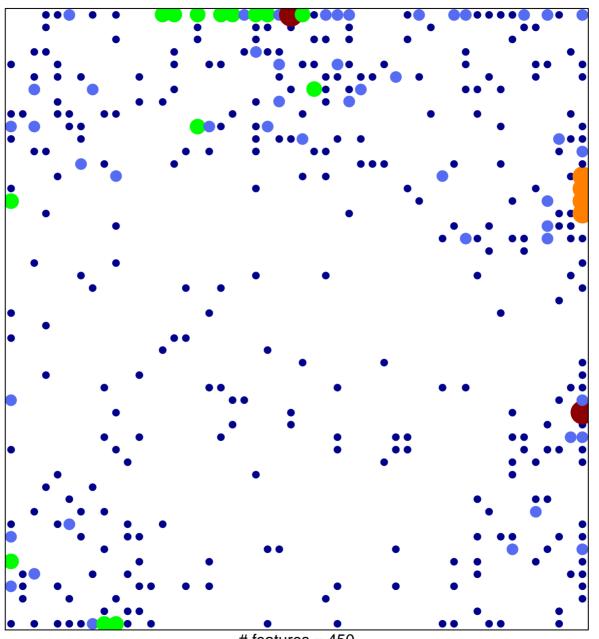
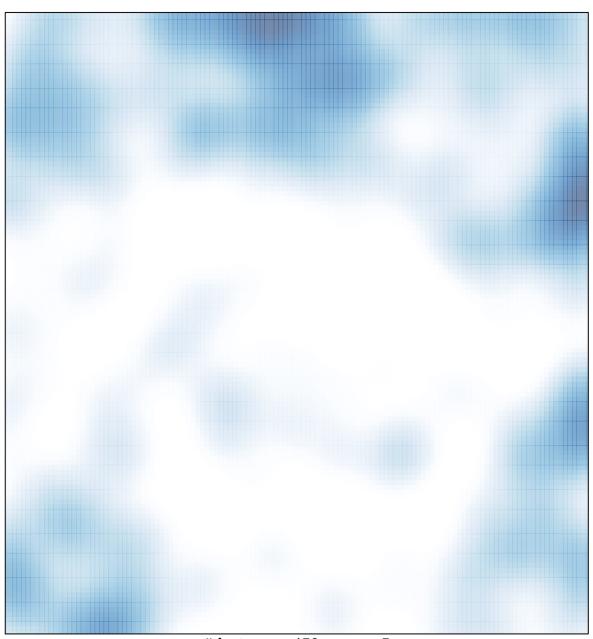
## hsa-miR-130b



# features = 450 chi-square p = 0.59

## hsa-miR-130b



# features = 450 , max = 5