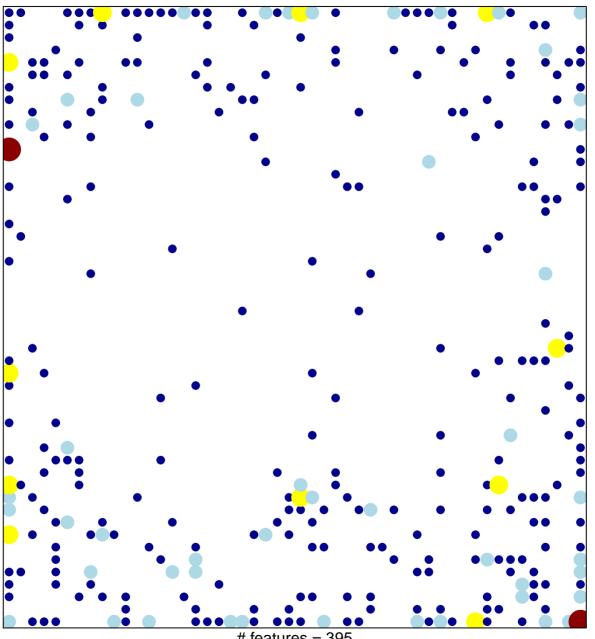
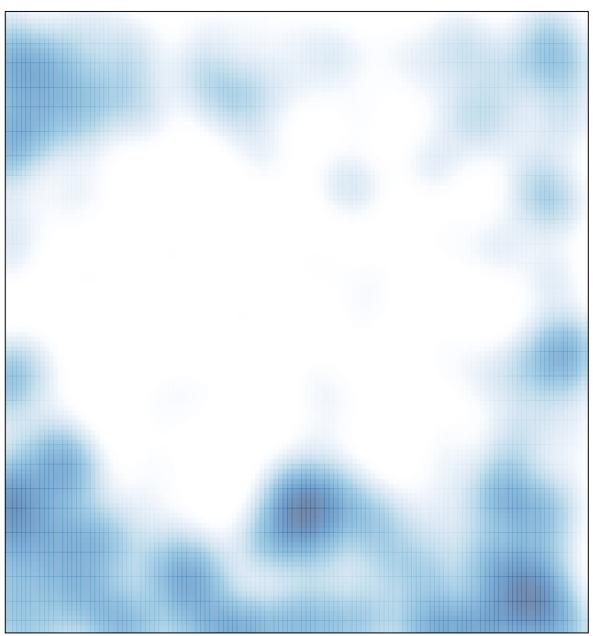
## hsa-miR-301b



# features = 395 chi-square p = 0.58

## hsa-miR-301b



# features = 395 , max = 4