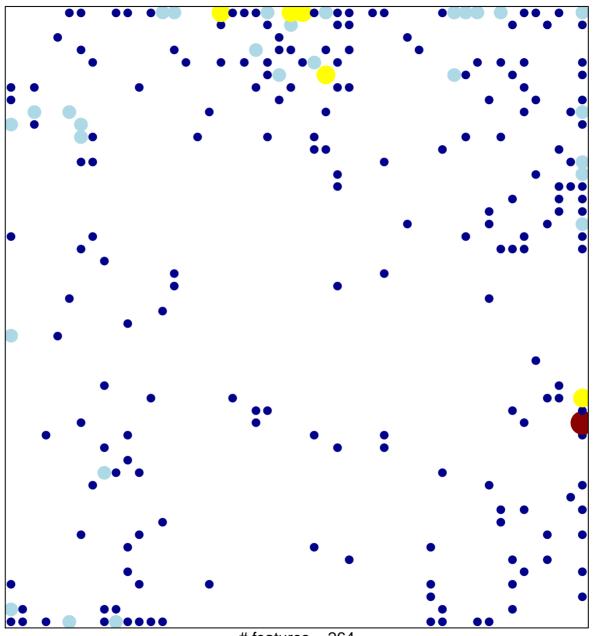
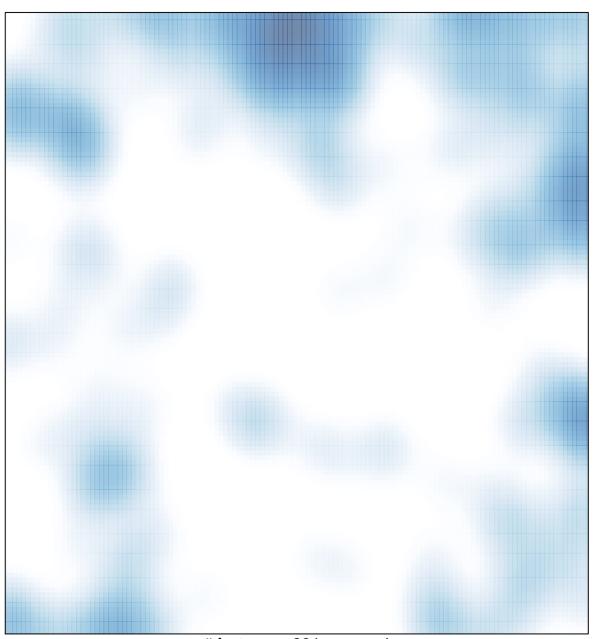
## hsa-miR-30a



# features = 264 chi-square p = 0.62

## hsa-miR-30a



# features = 264 , max = 4