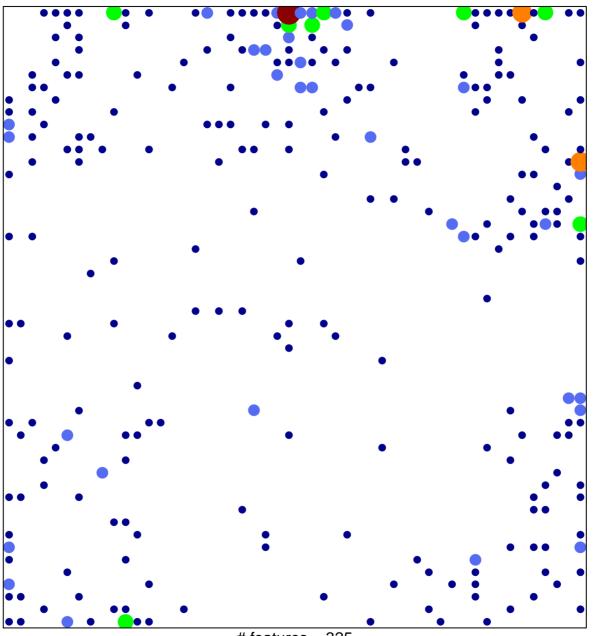
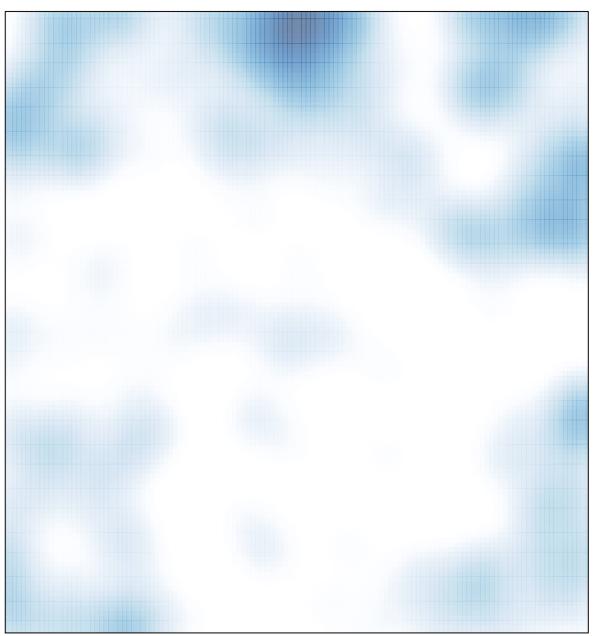
## hsa-miR-18a



# features = 325 chi-square p = 0.63

## hsa-miR-18a



# features = 325 , max = 5