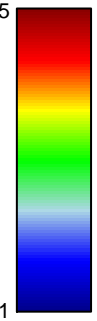
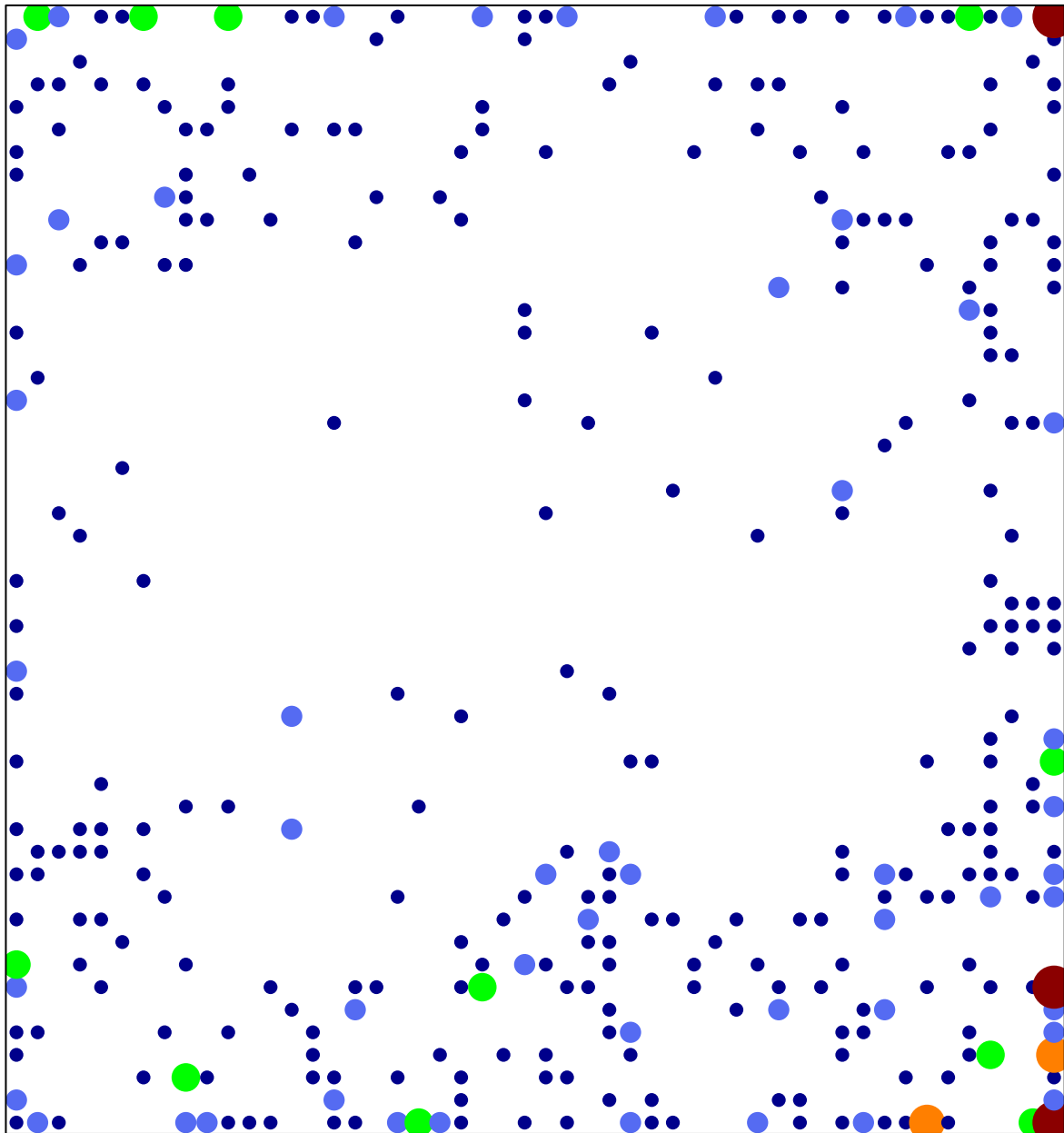
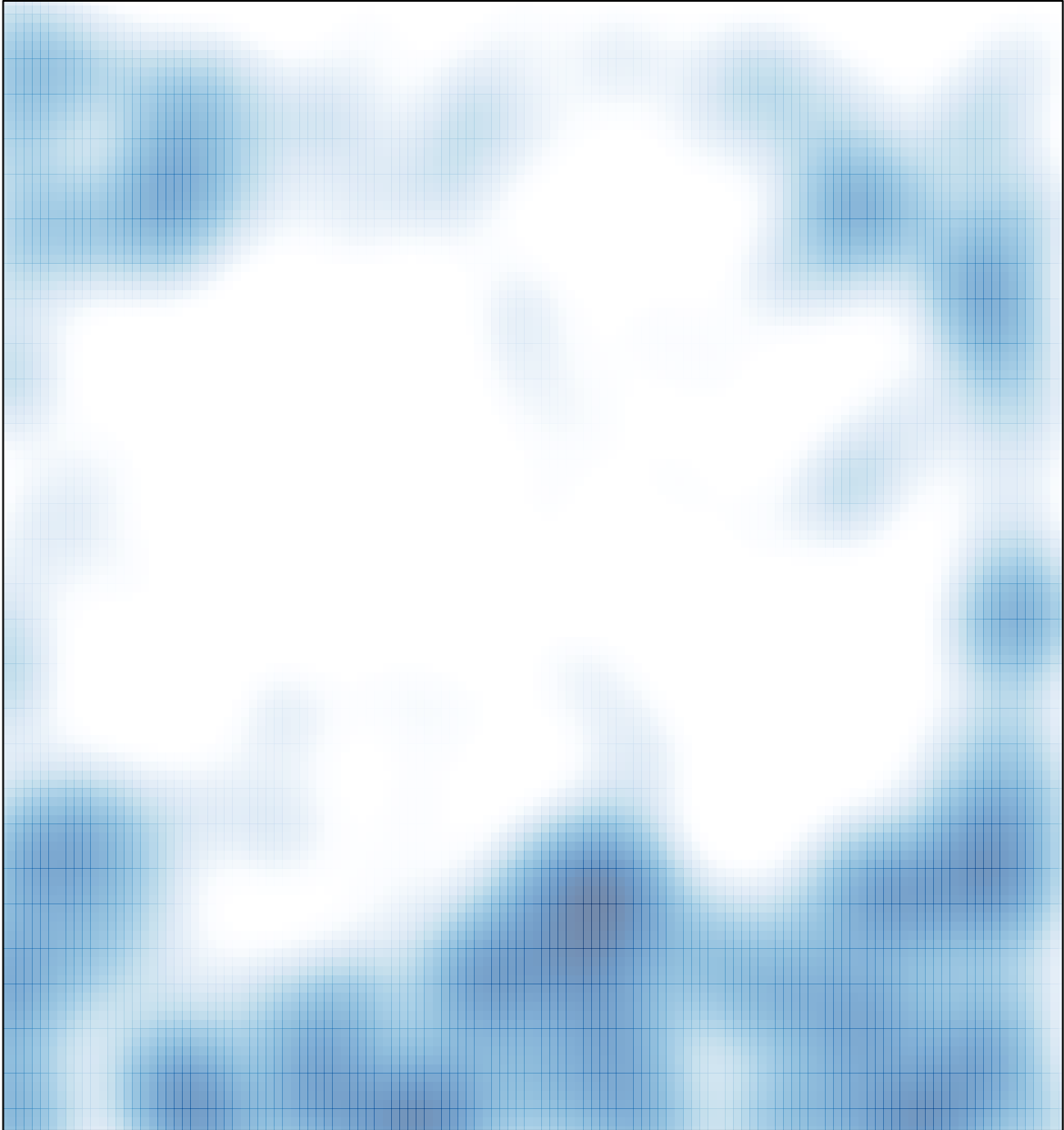


# hsa-miR-519d



# features = 426  
chi-square p = 0.56

# hsa-miR-519d



# features = 426 , max = 5