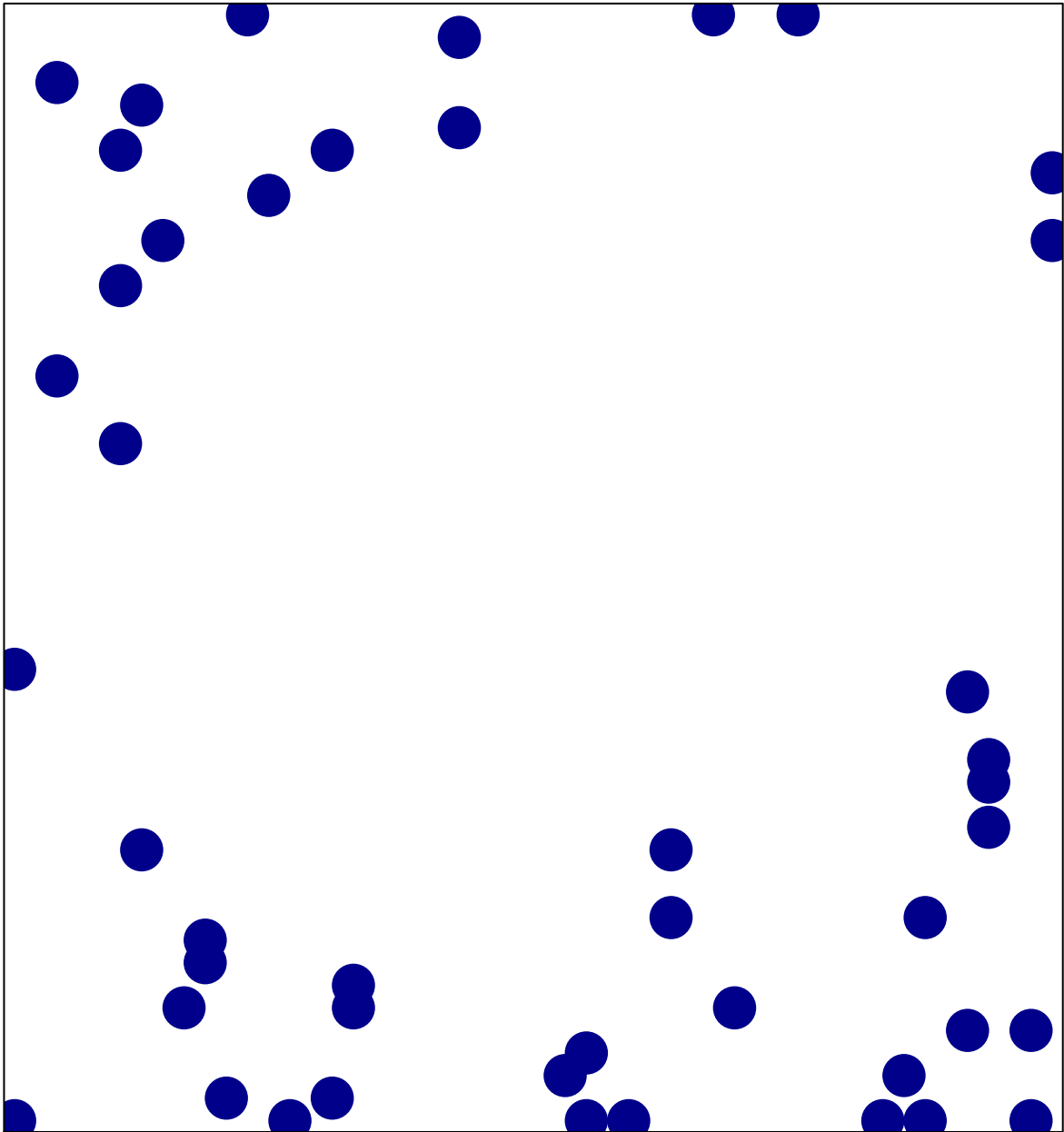
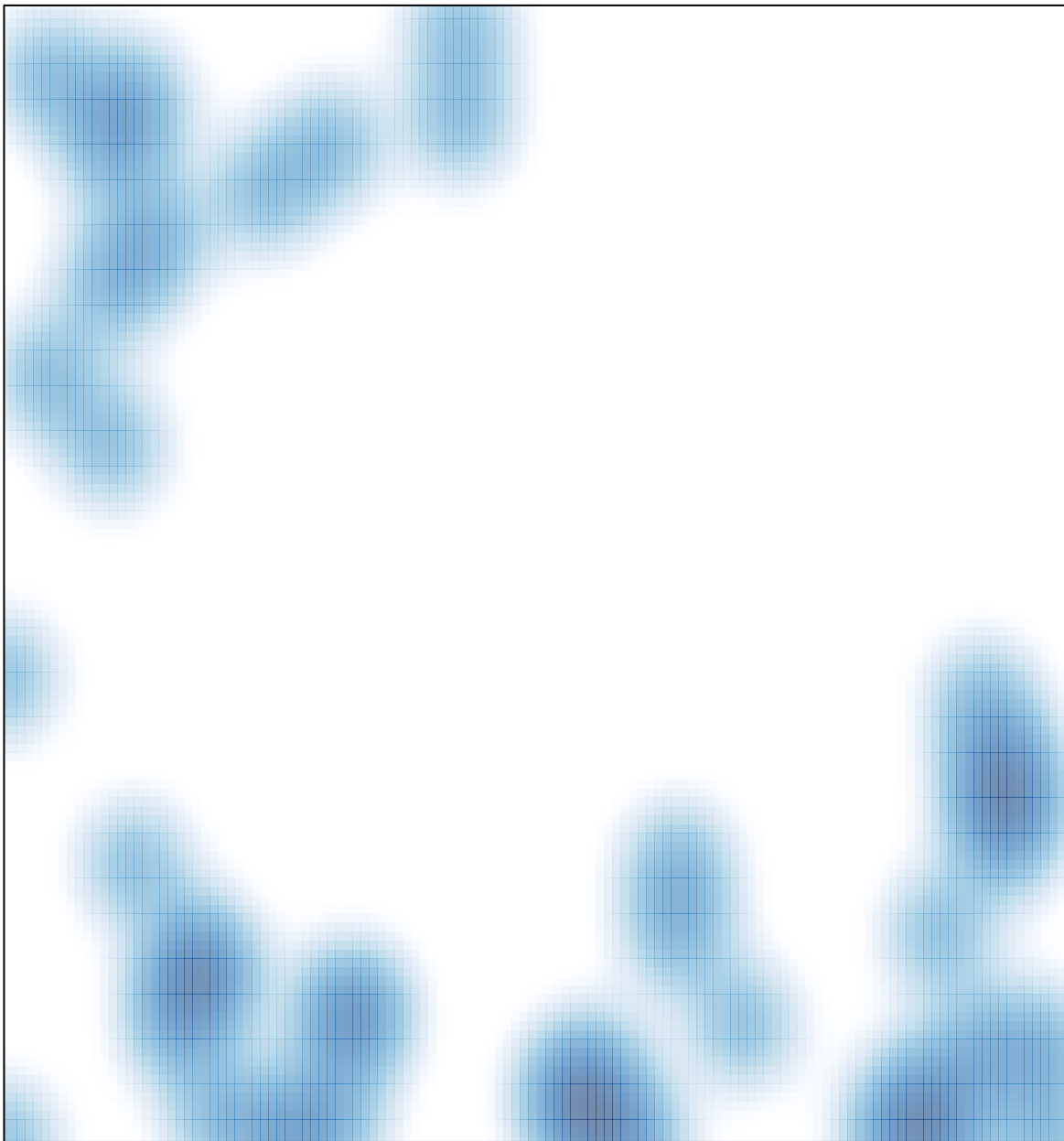


# hsa-miR-660



# features = 45  
chi-square p = 0.82

# hsa-miR-660



# features = 45 , max = 1