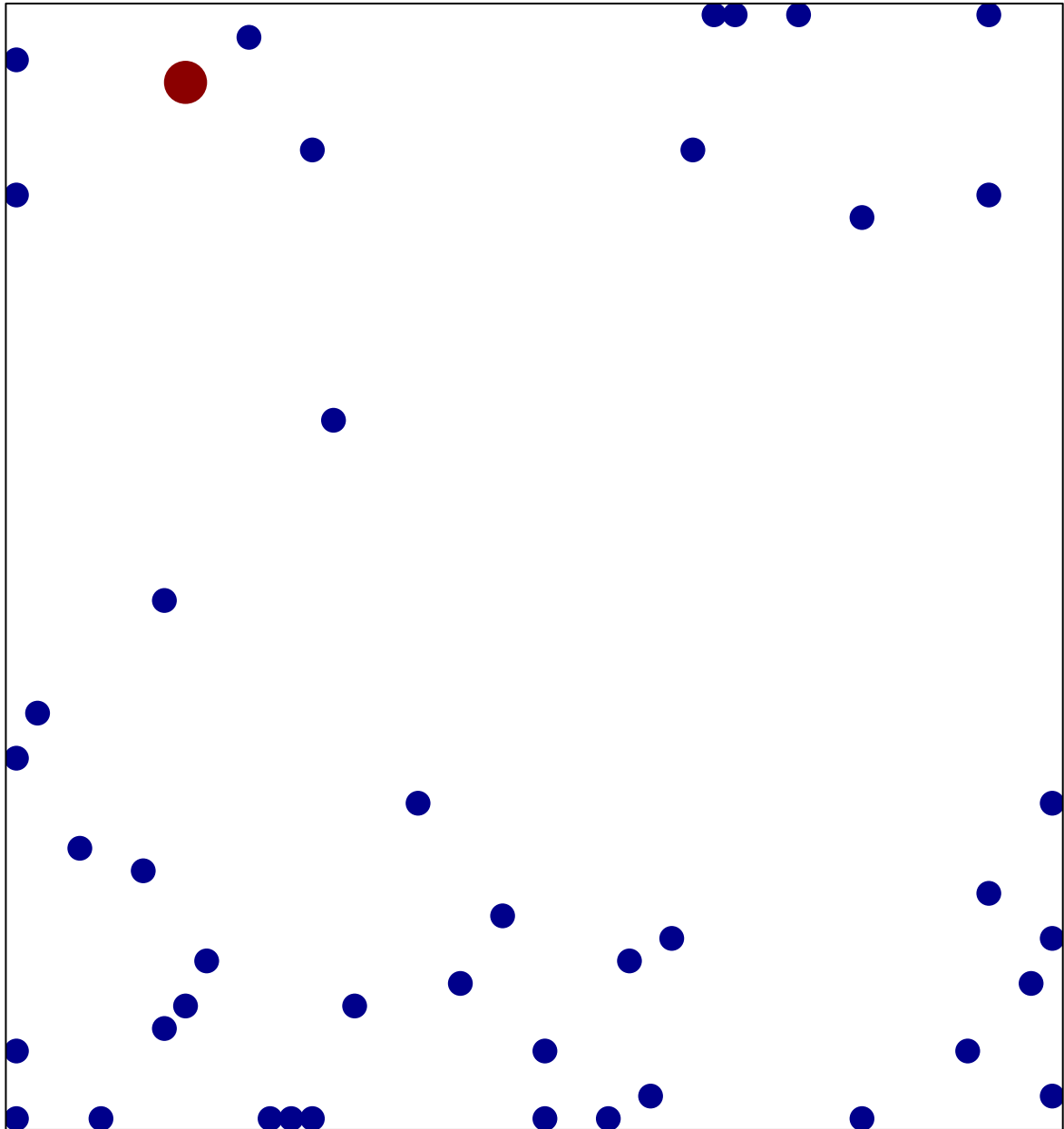
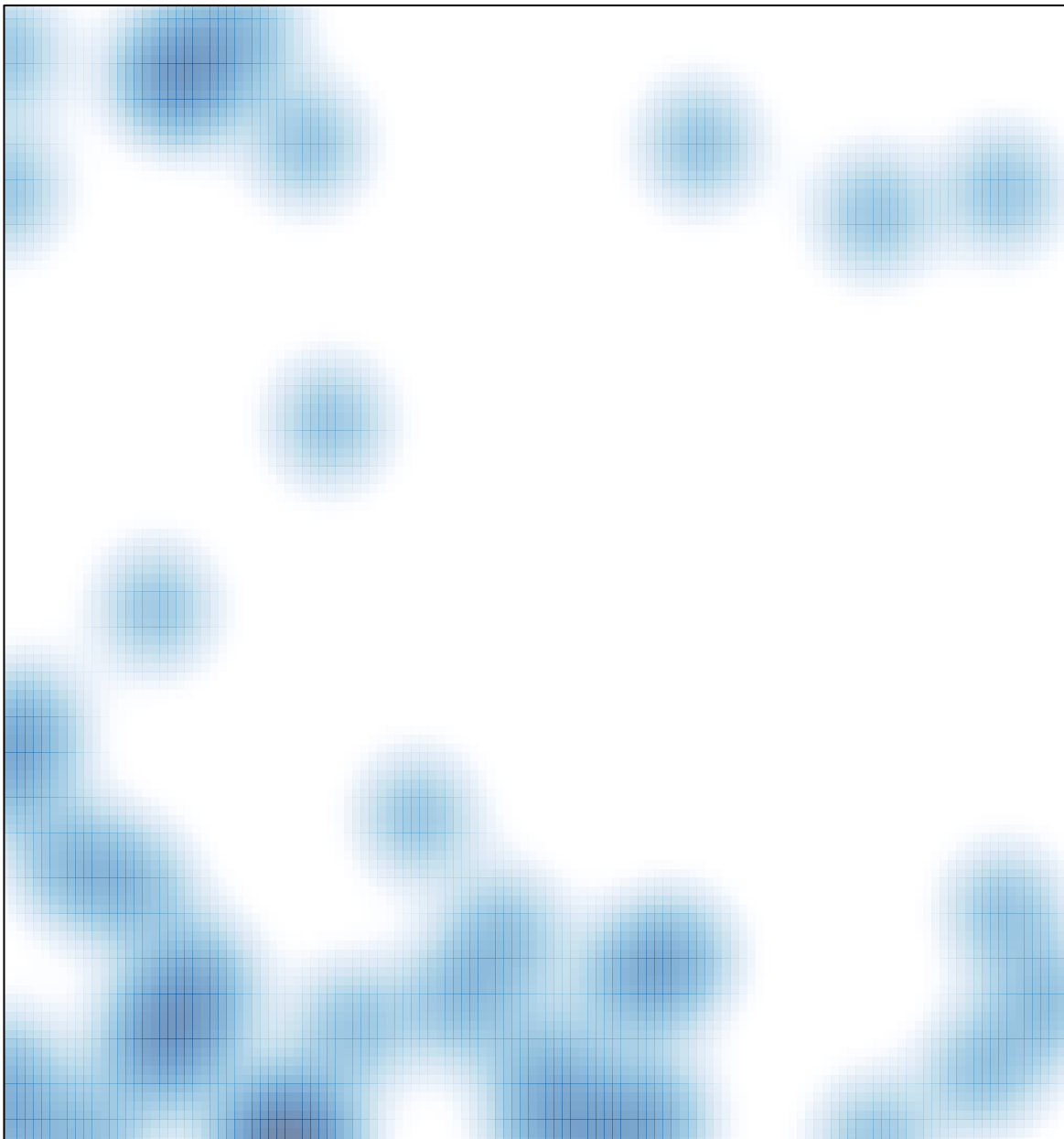


# hsa-miR-411



# features = 45  
chi-square p = 0.8

# hsa-miR-411



# features = 45 , max = 2