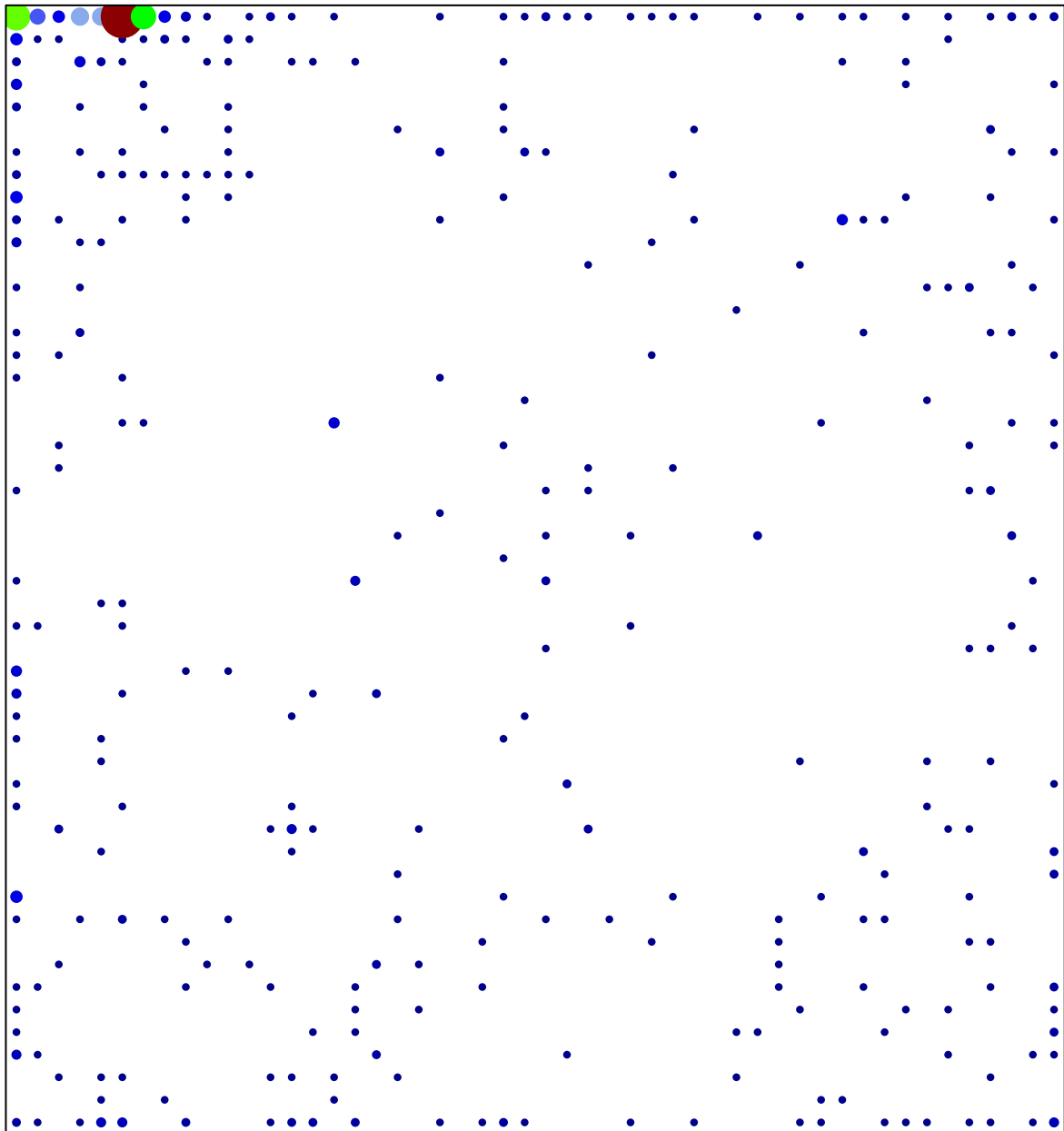
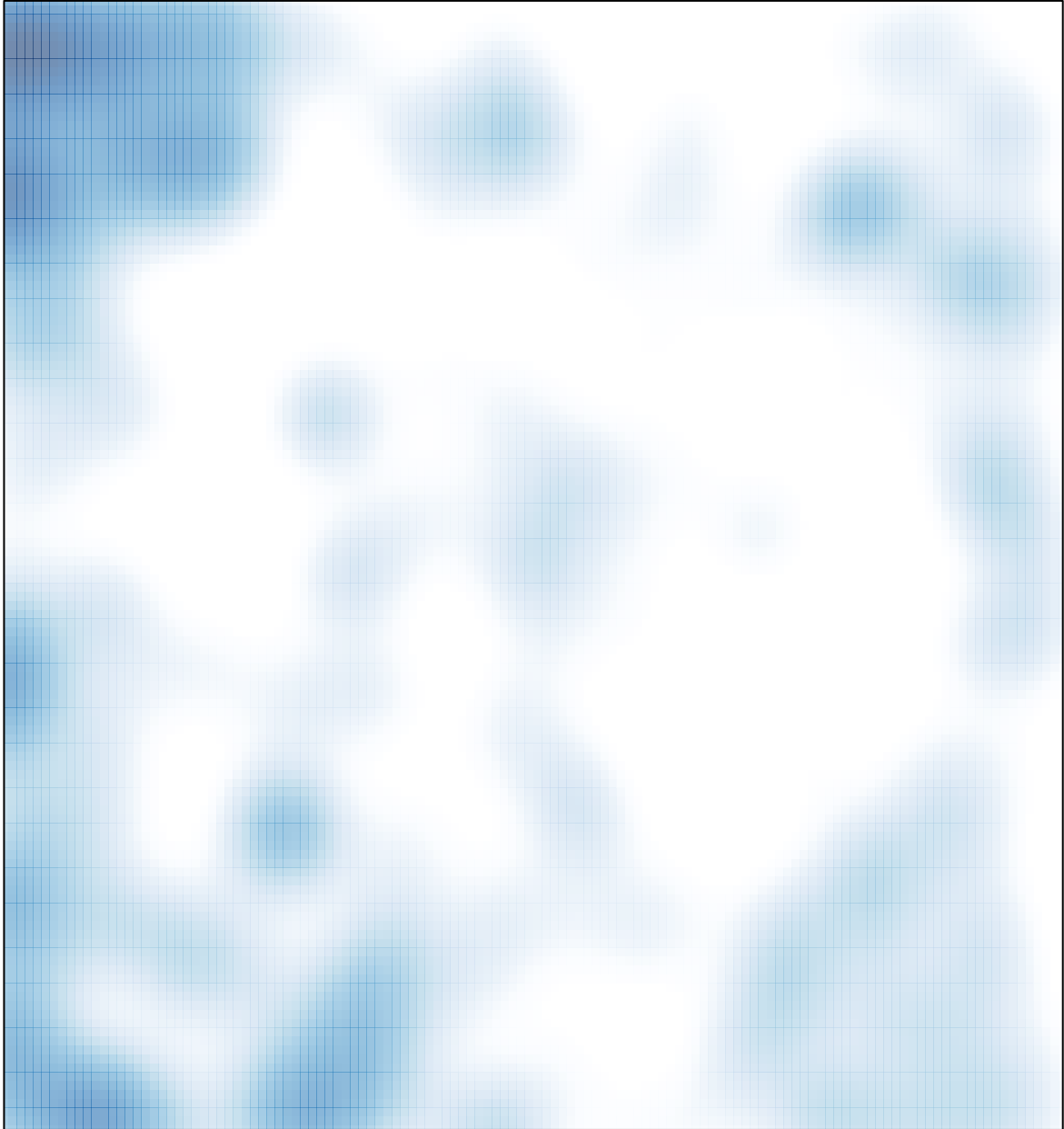


# Lembcke\_Normal vs Adenoma



# features = 489  
chi-square p = 0.31

# Lembcke\_Normal vs Adenoma



# features = 489 , max = 31