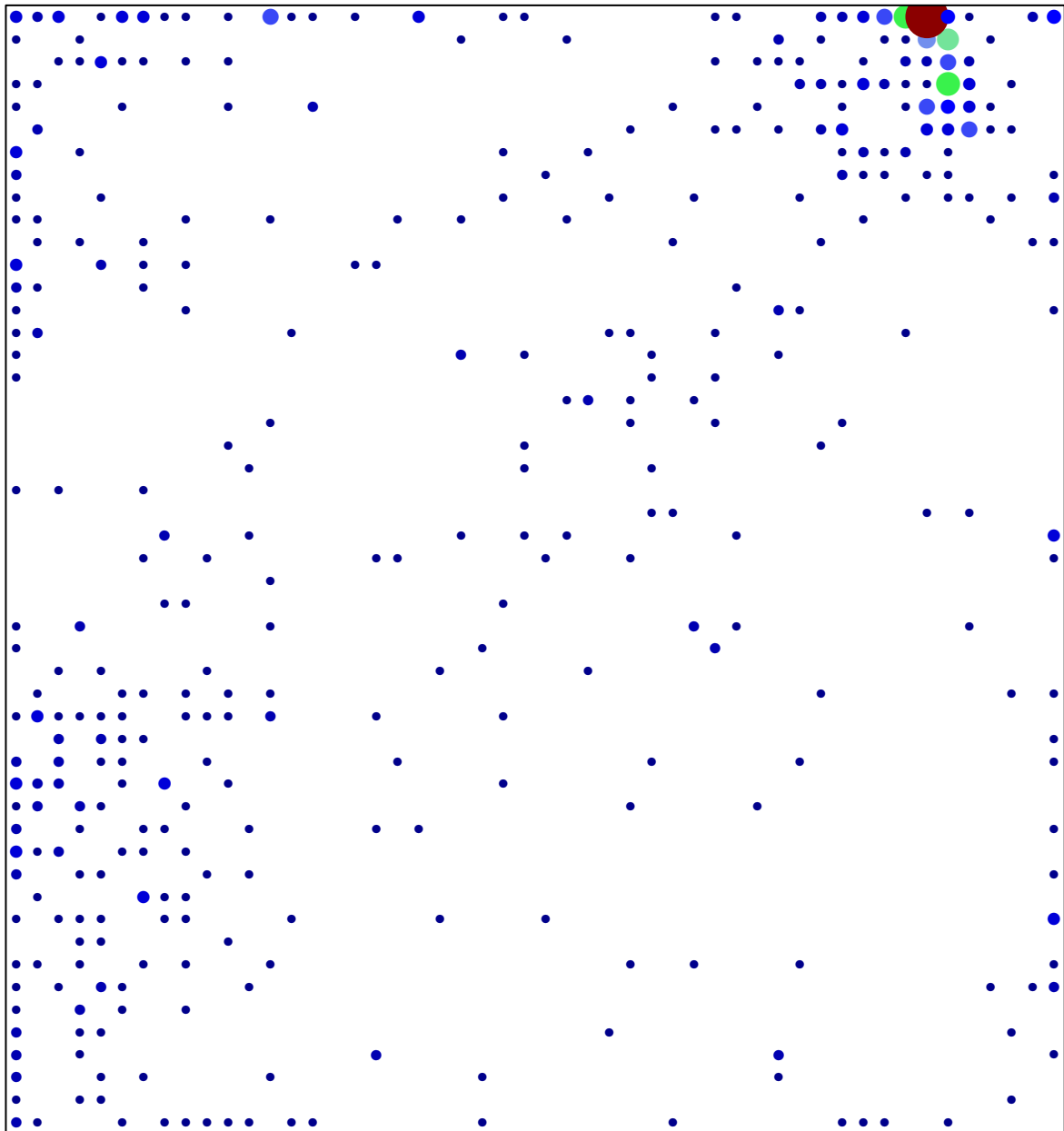
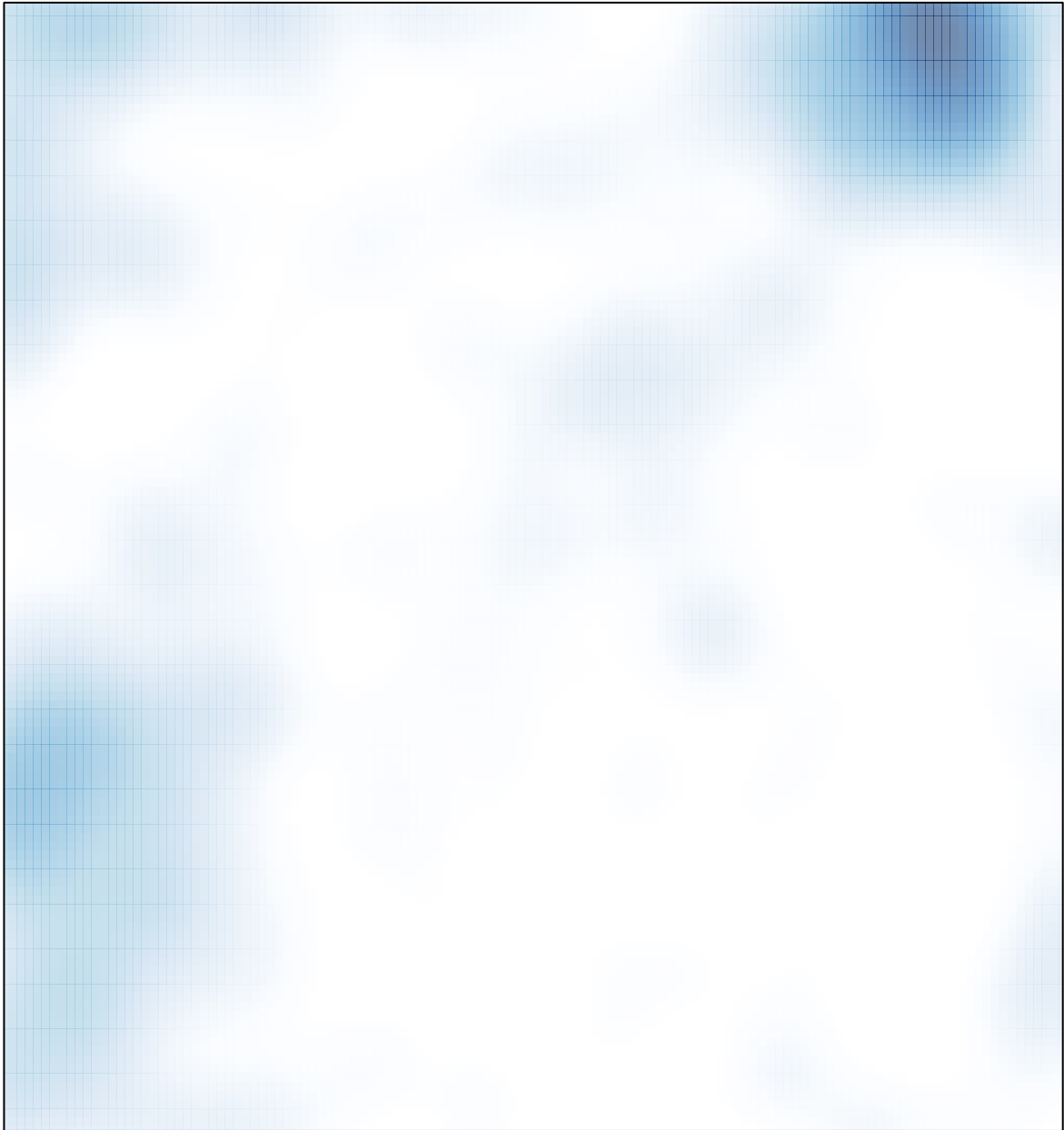


# Lembcke\_Normal vs Adenoma



# features = 531  
chi-square p = 0.49

# Lembcke\_Normal vs Adenoma



# features = 531 , max = 19