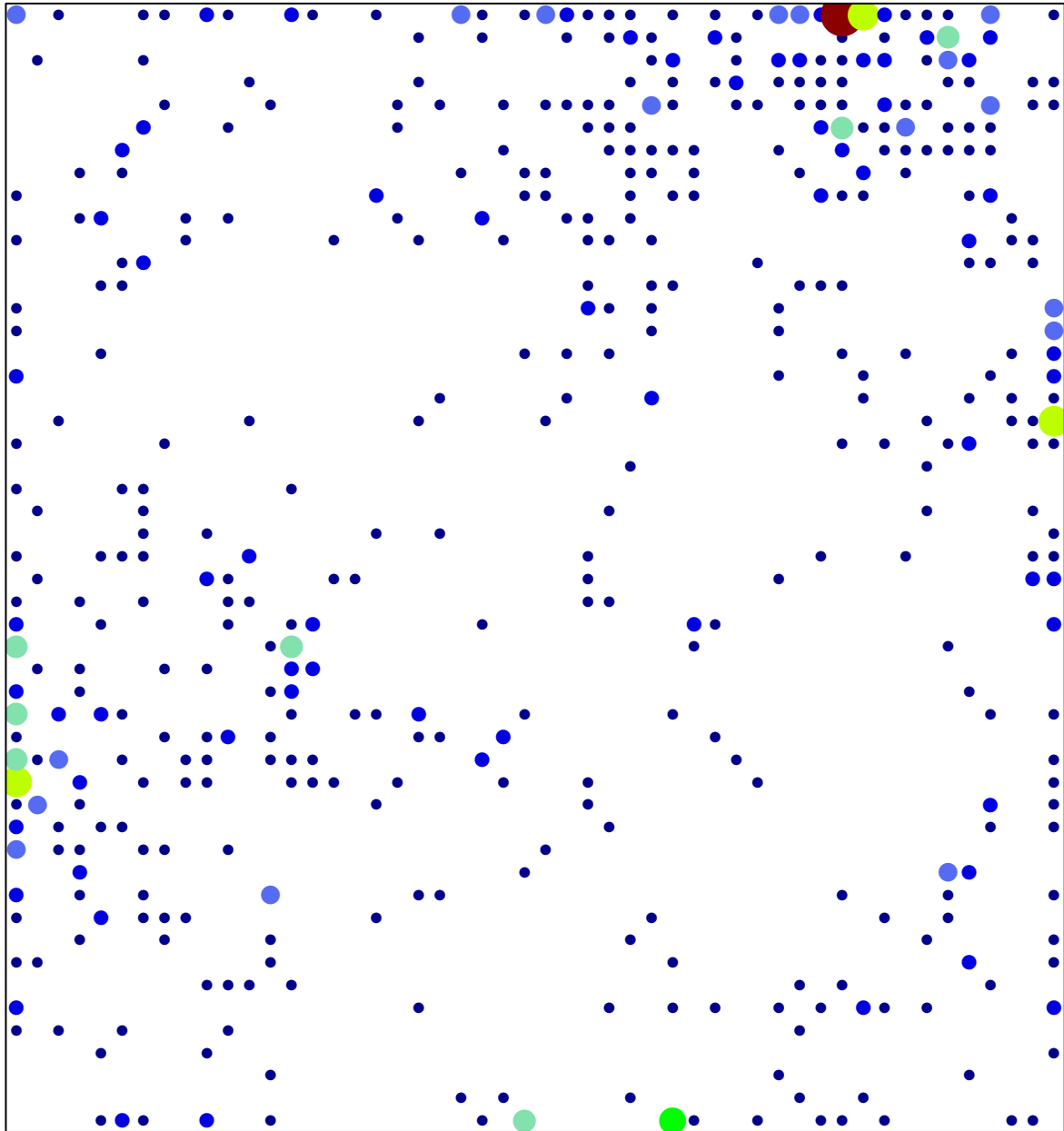
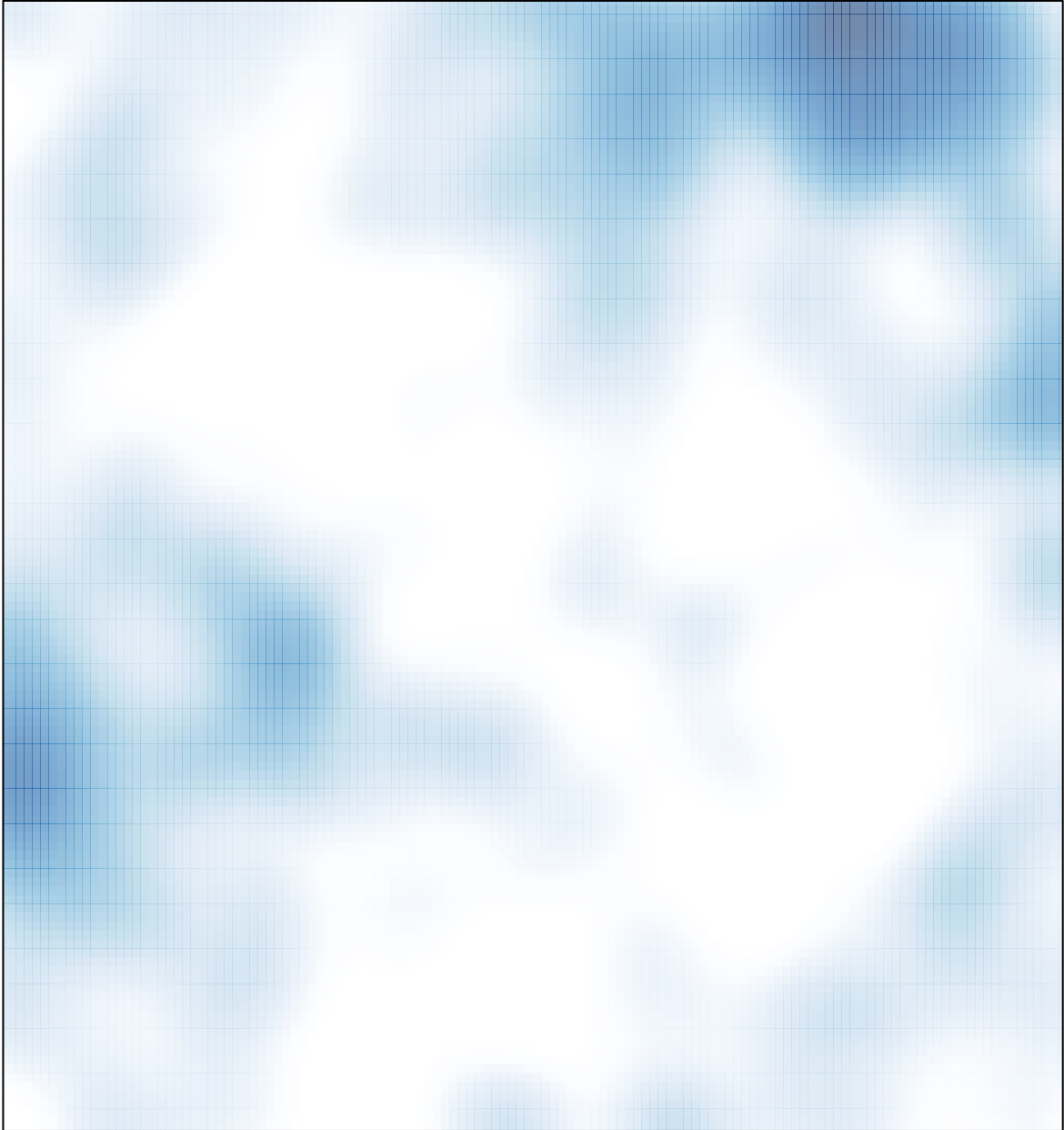


# nucleolus



# features = 592  
chi-square p = 0.62

# nucleolus



# features = 592 , max = 9