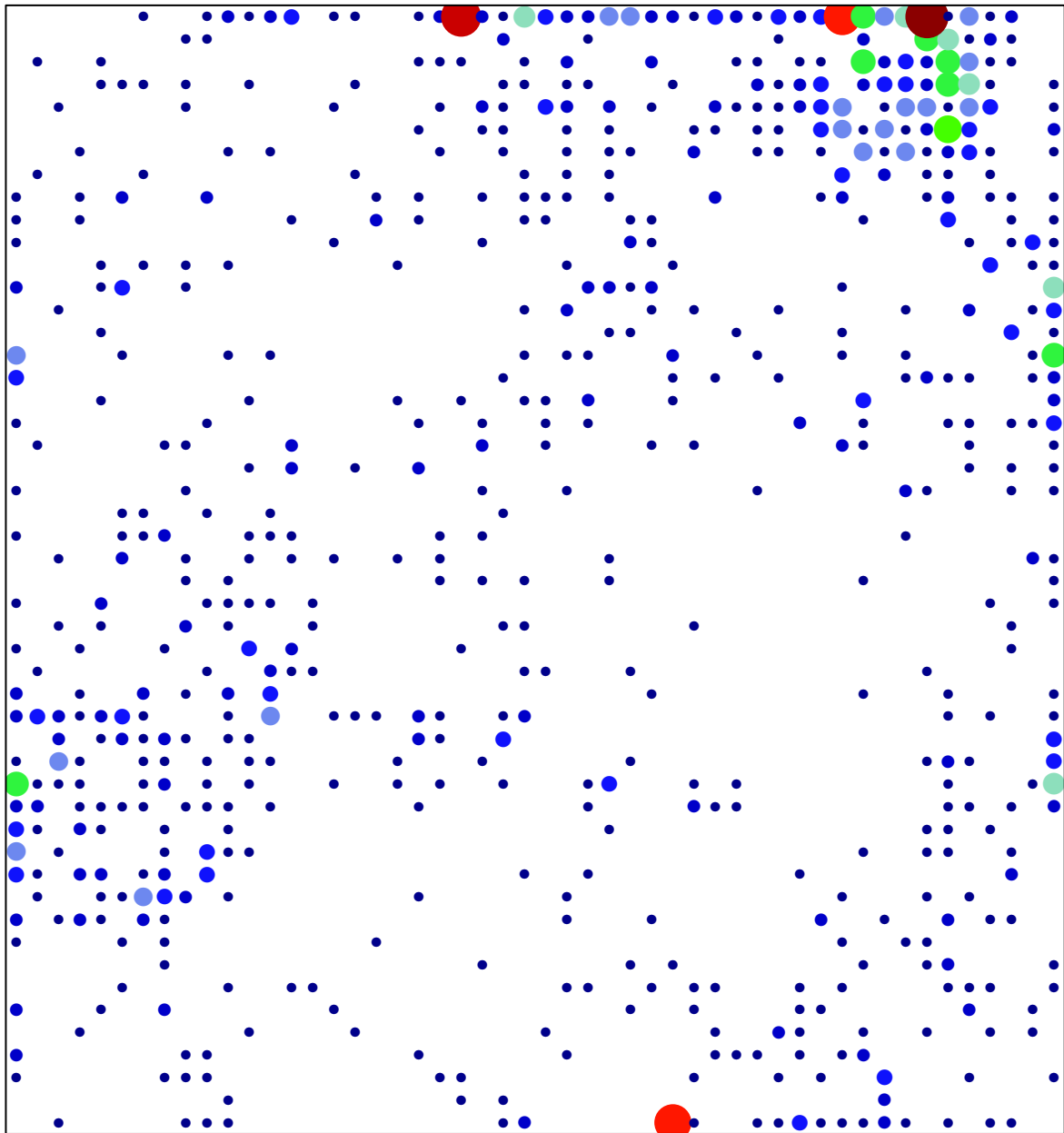


# nucleoplasm

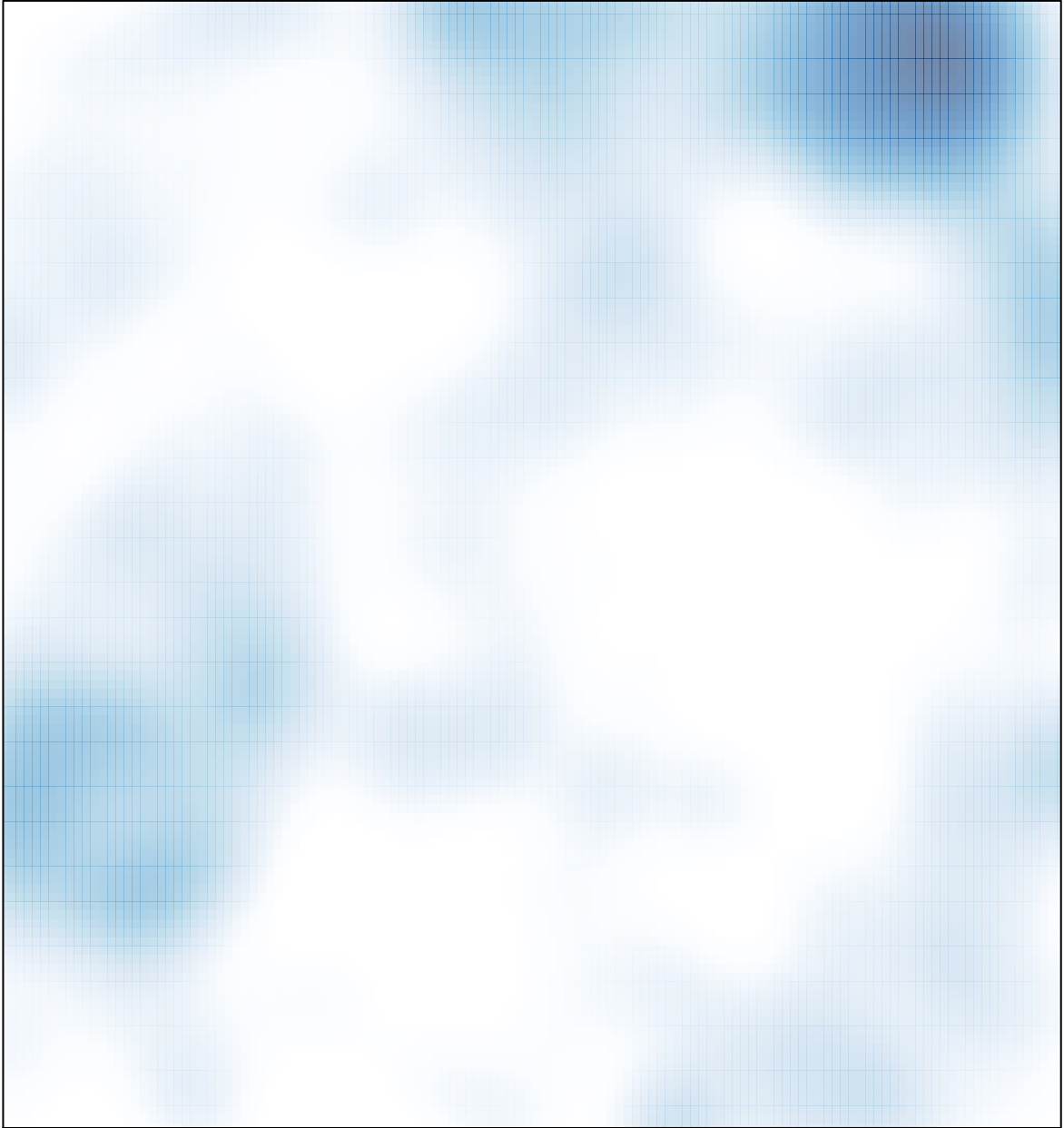


12

1

# features = 973  
chi-square p = 0.53

# nucleoplasm



# features = 973 , max = 12