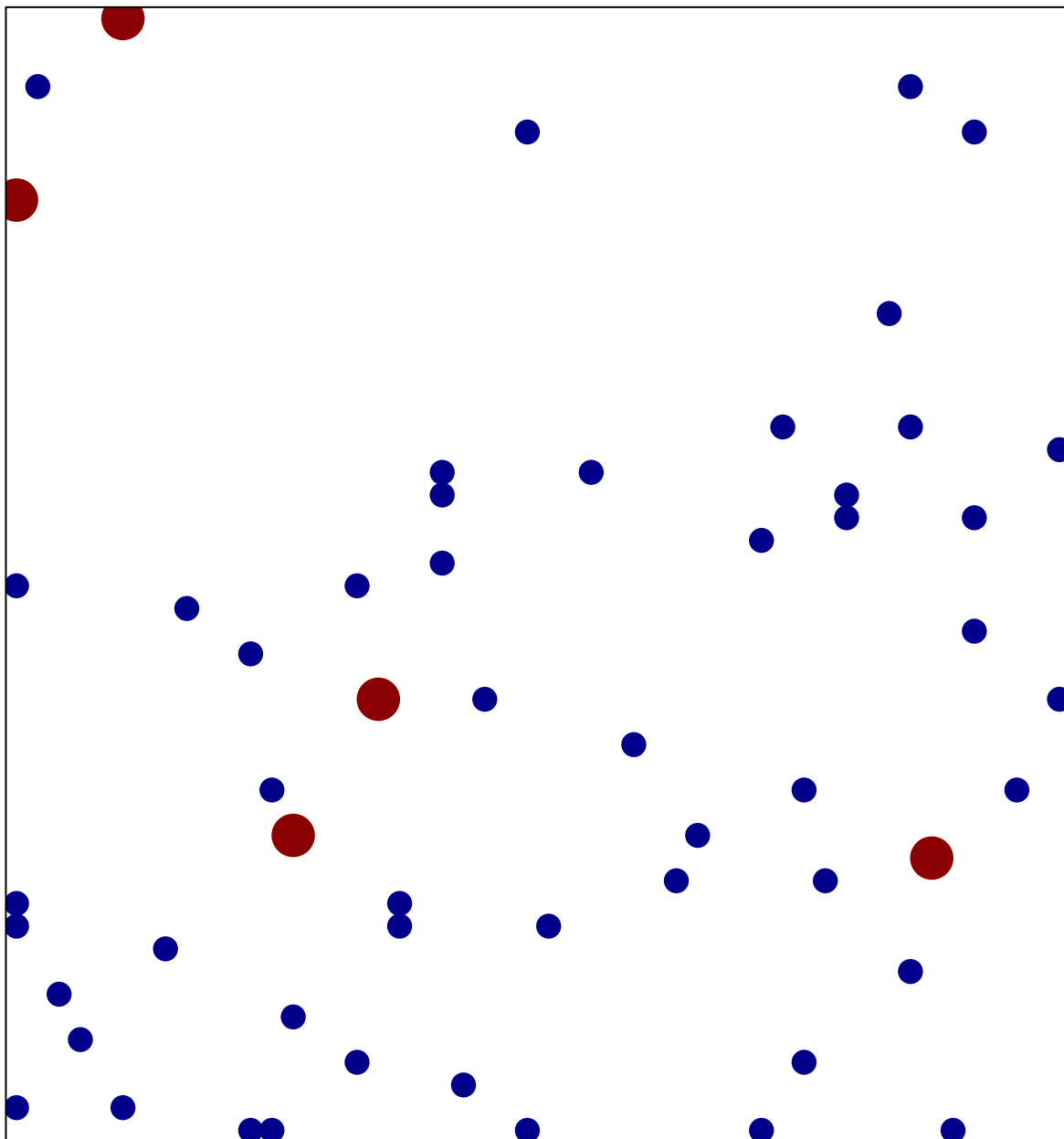
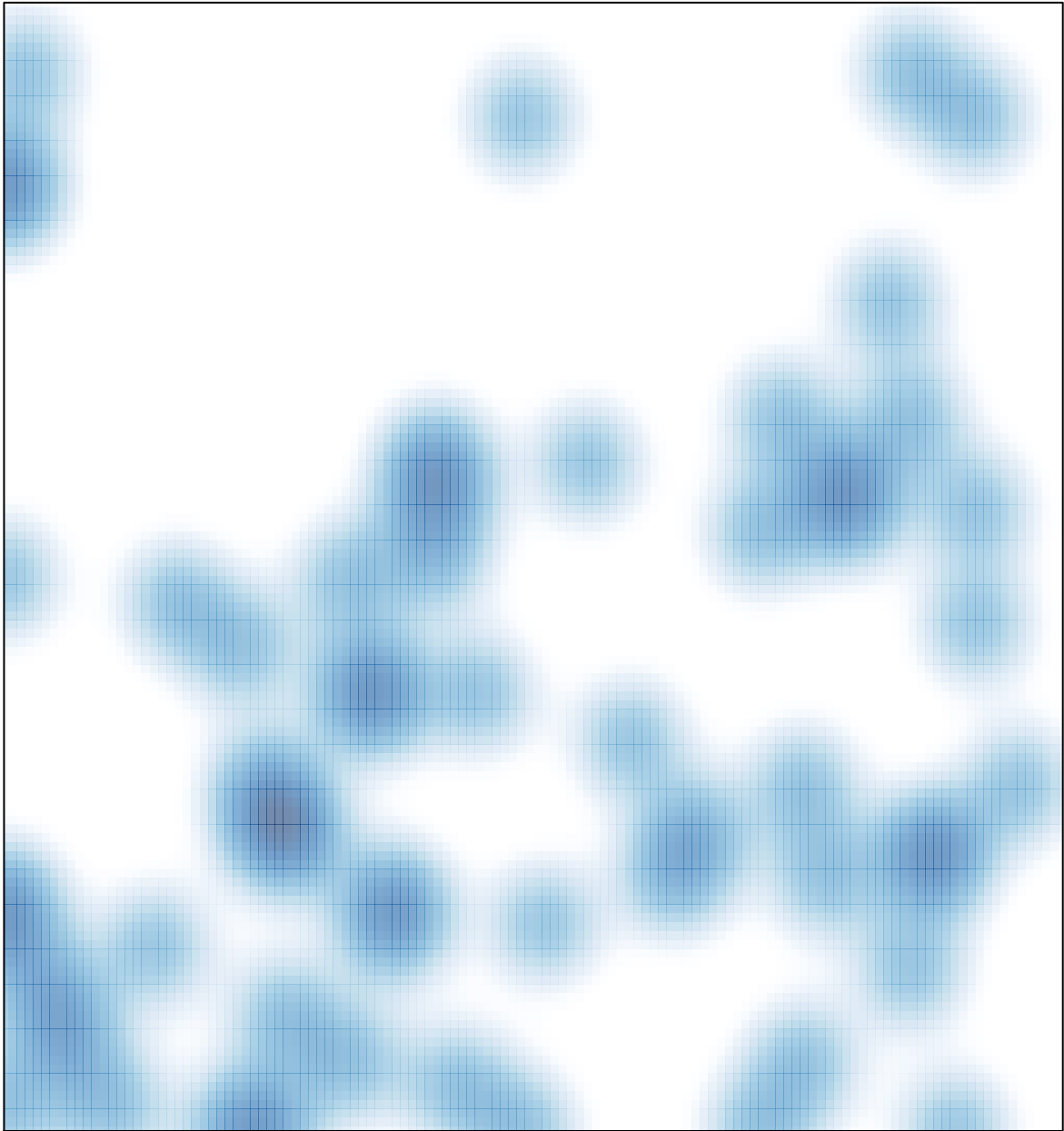


# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_UP



# features = 60  
chi-square p = 0.83

# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_UP



# features = 60 , max = 2