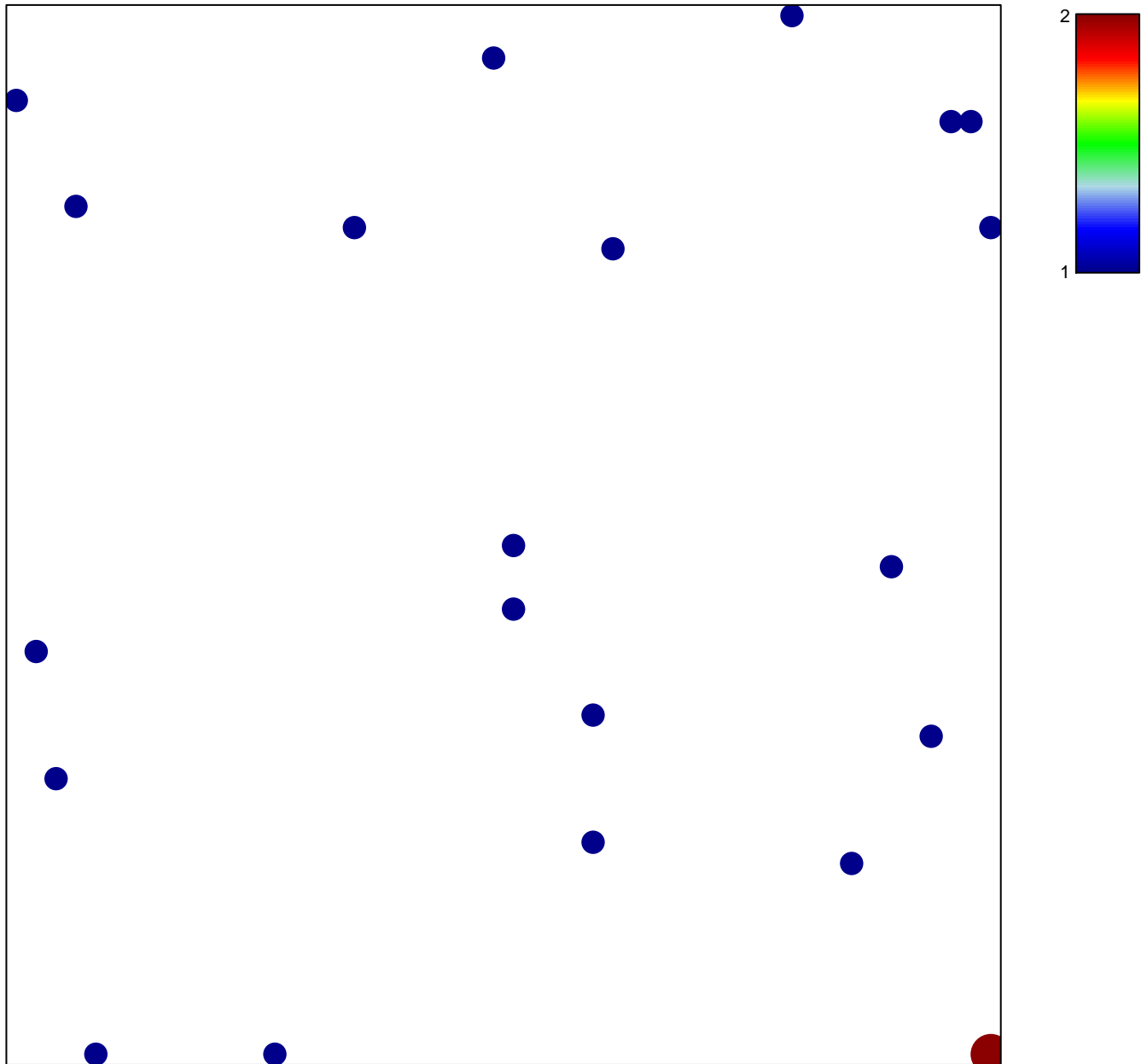
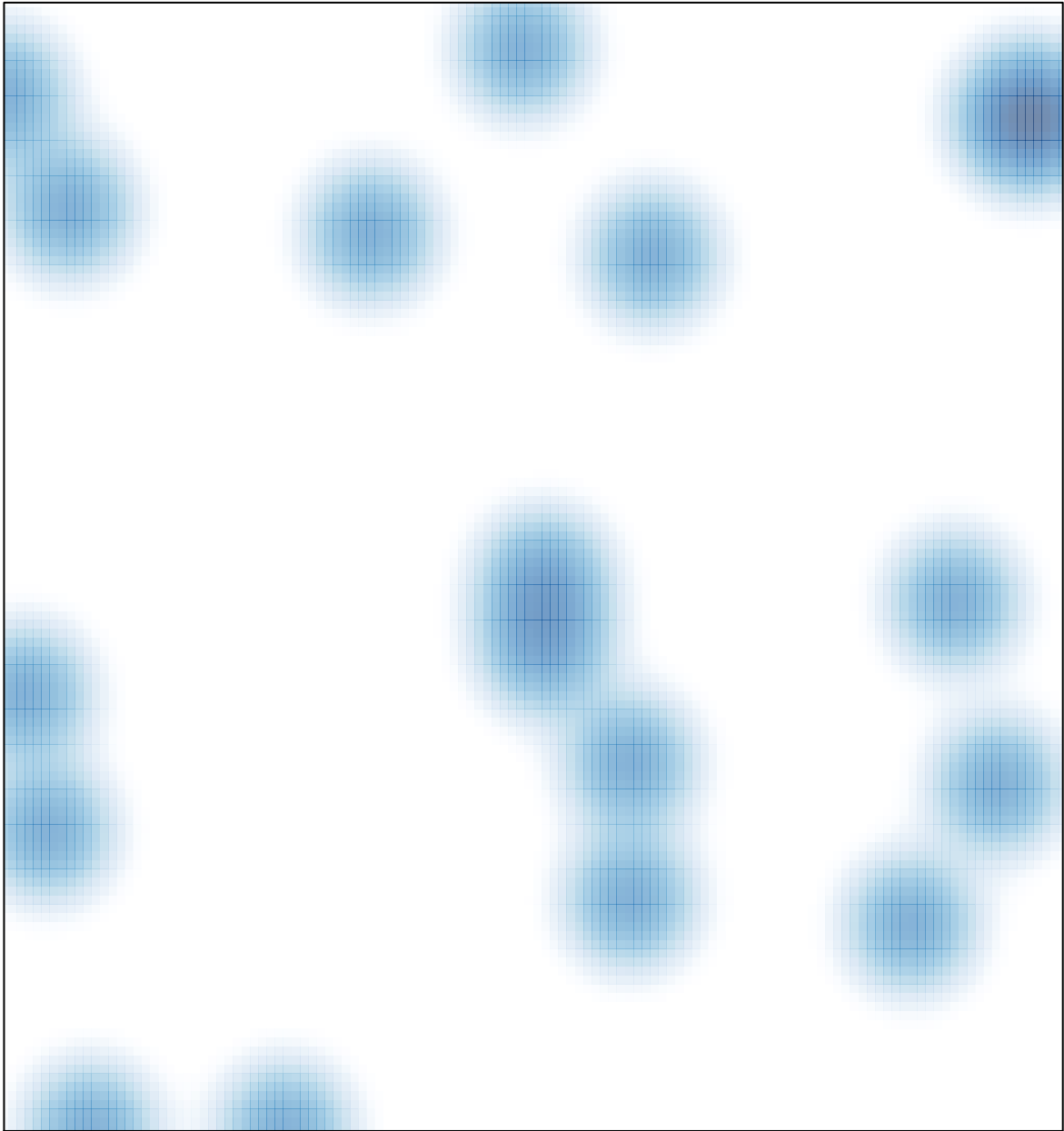


# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotC



# features = 22  
chi-square p = 0.83

# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotC



# features = 22 , max = 2