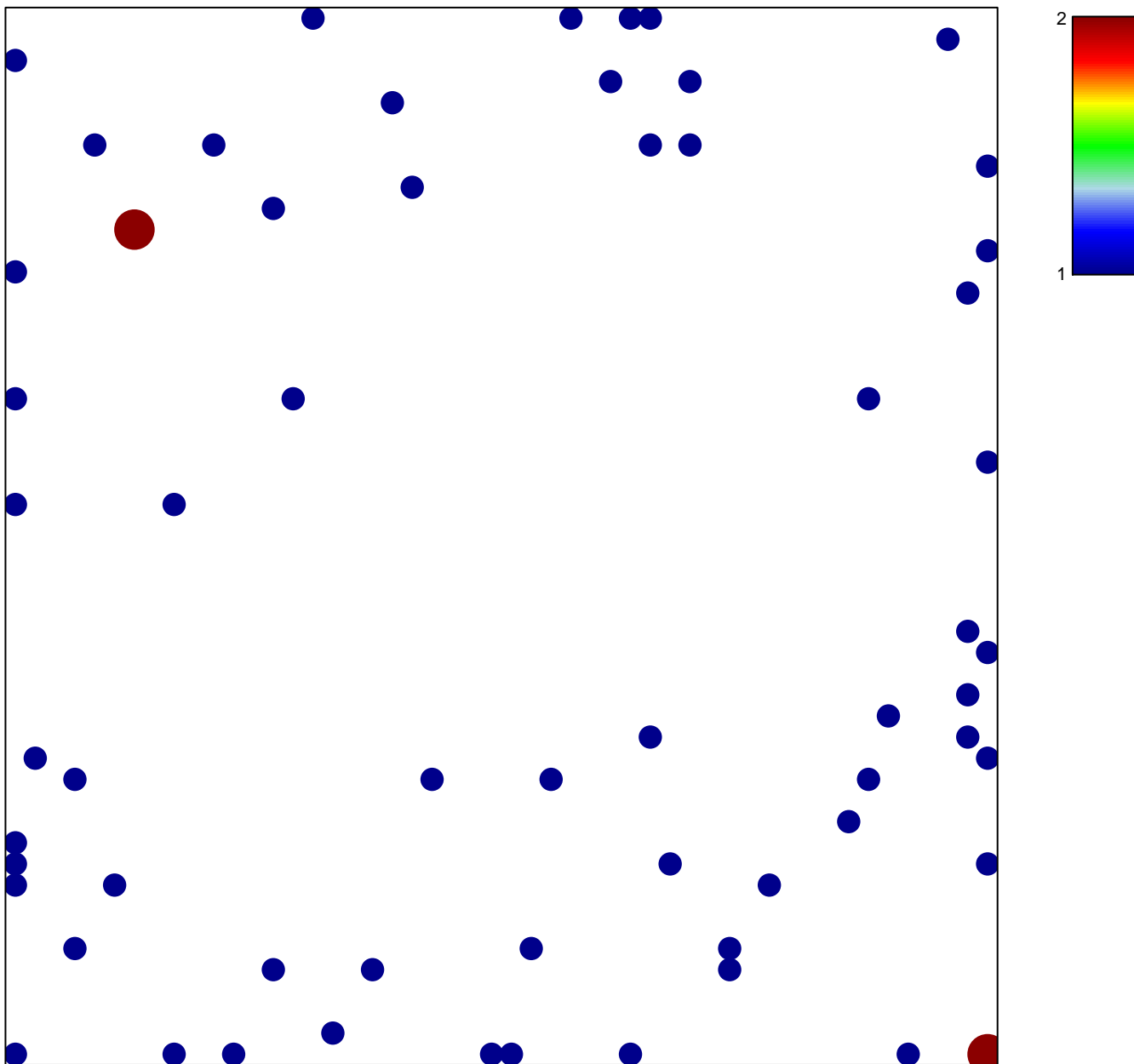
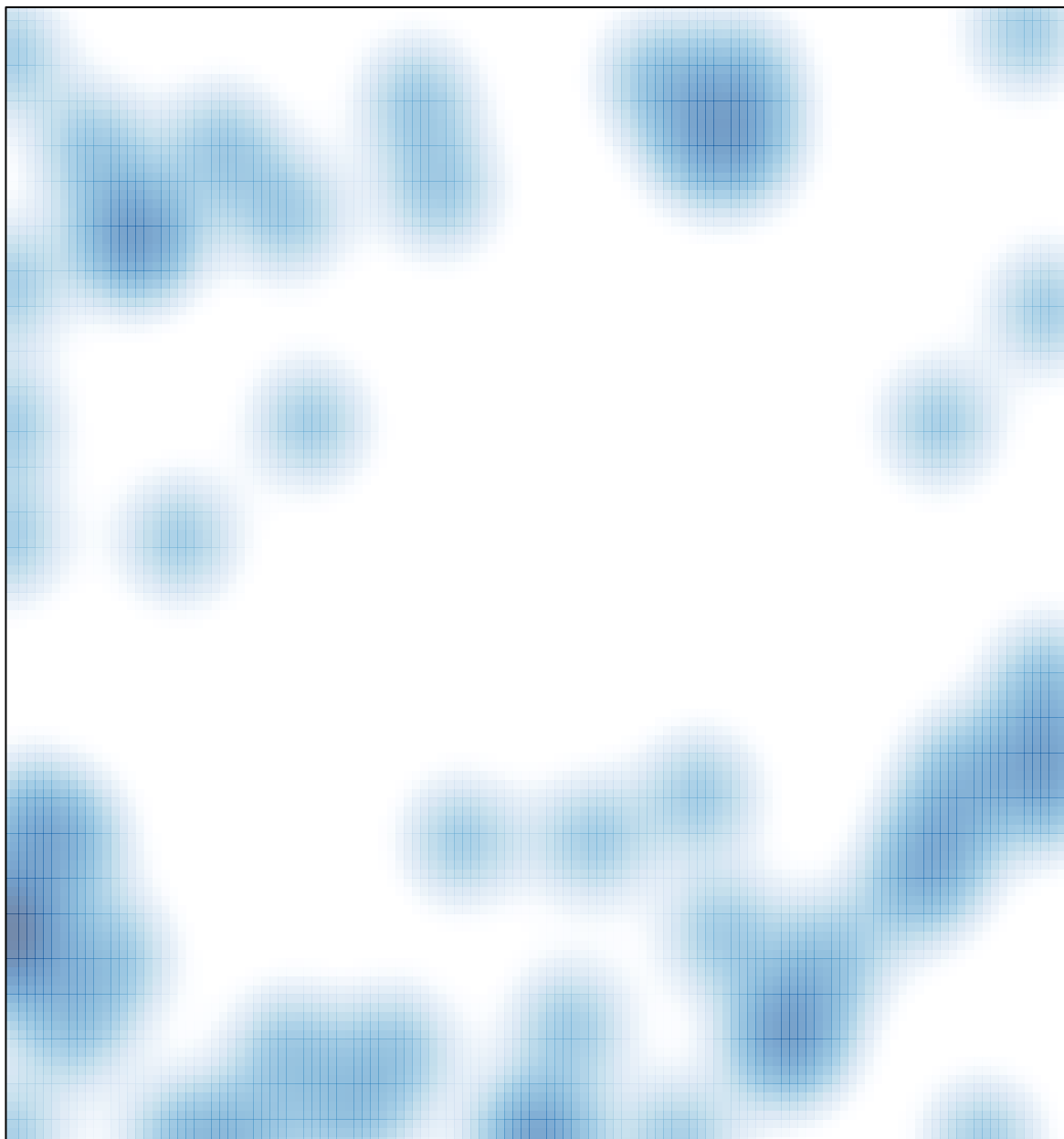


# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotJ



# features = 63  
chi-square p = 0.77

# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotJ



# features = 63 , max = 2