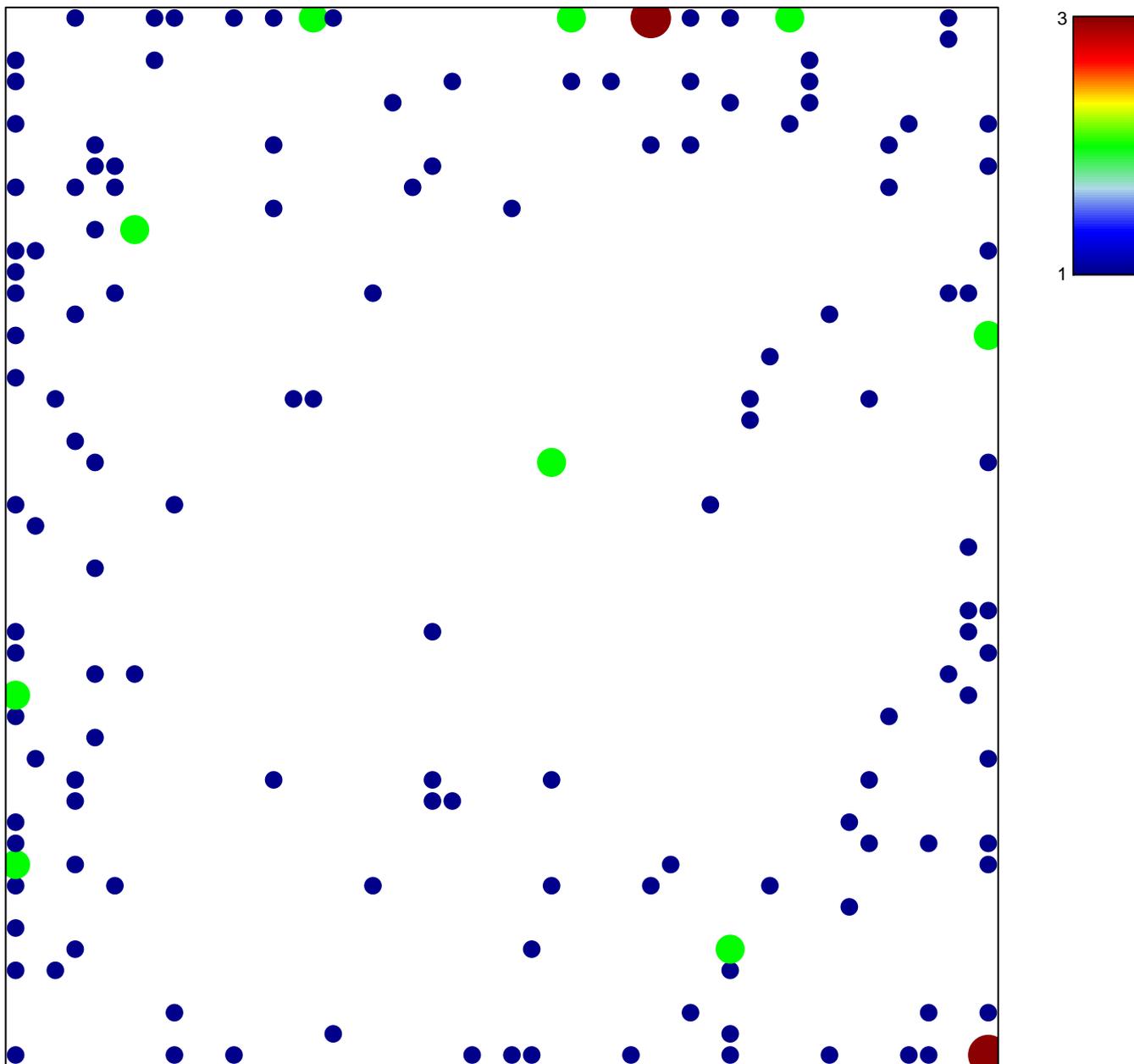
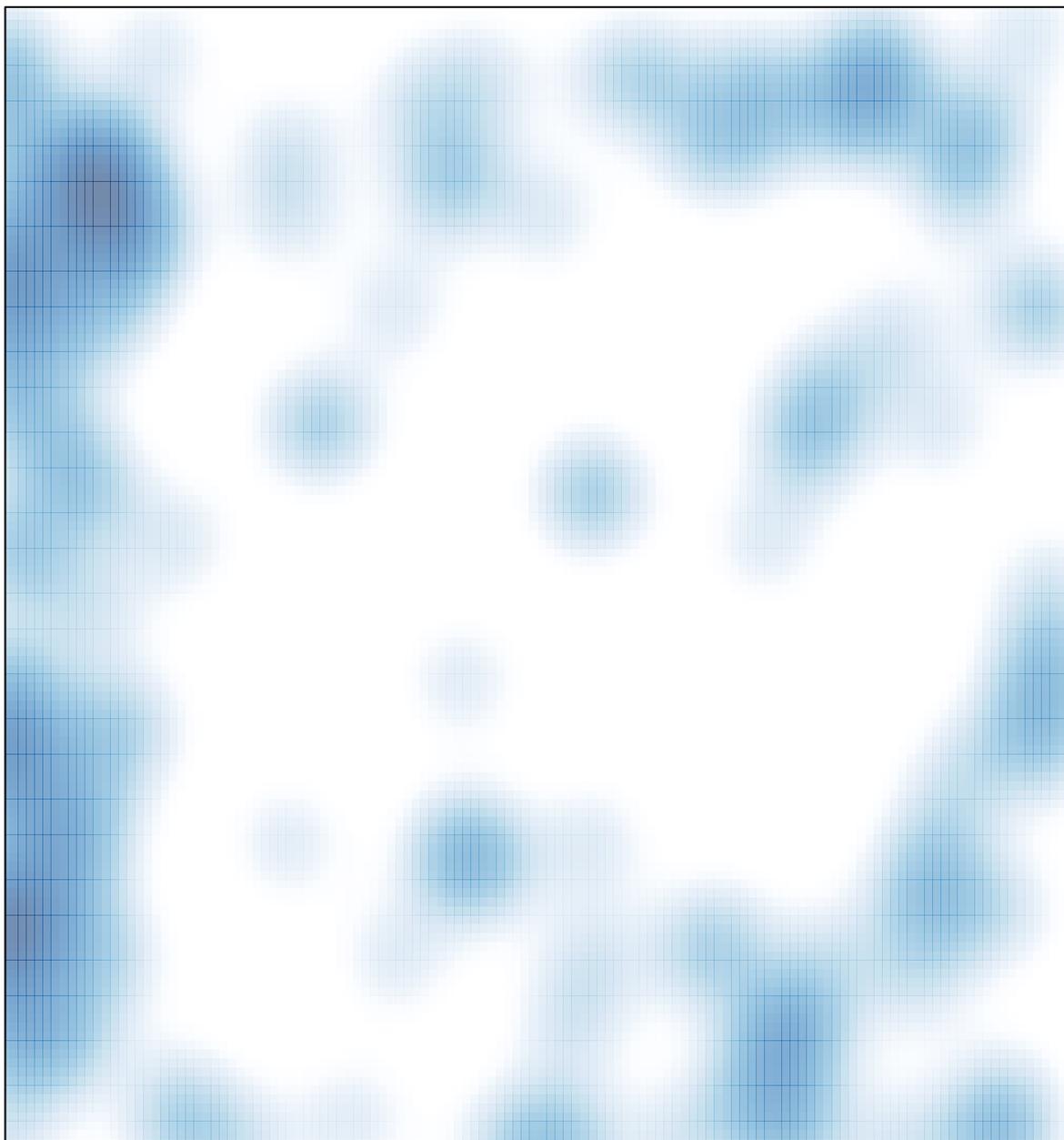


# WILLSCHER\_GBM\_proteomics\_wtOnly\_Differencelist



# features = 159  
chi-square p = 0.7

# WILLSCHER\_GBM\_proteomics\_wtOnly\_Differencelis



# features = 159 , max = 3