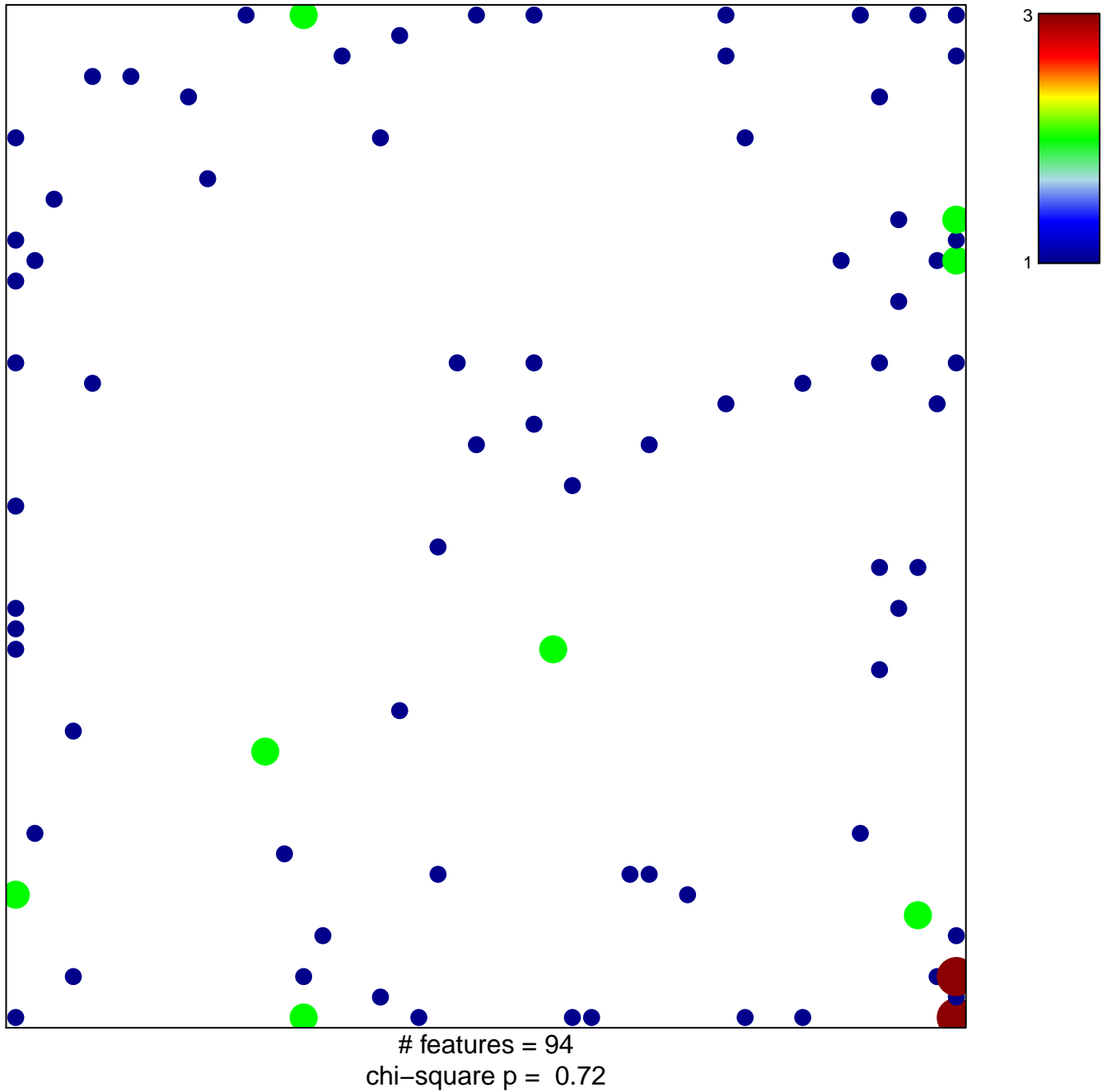
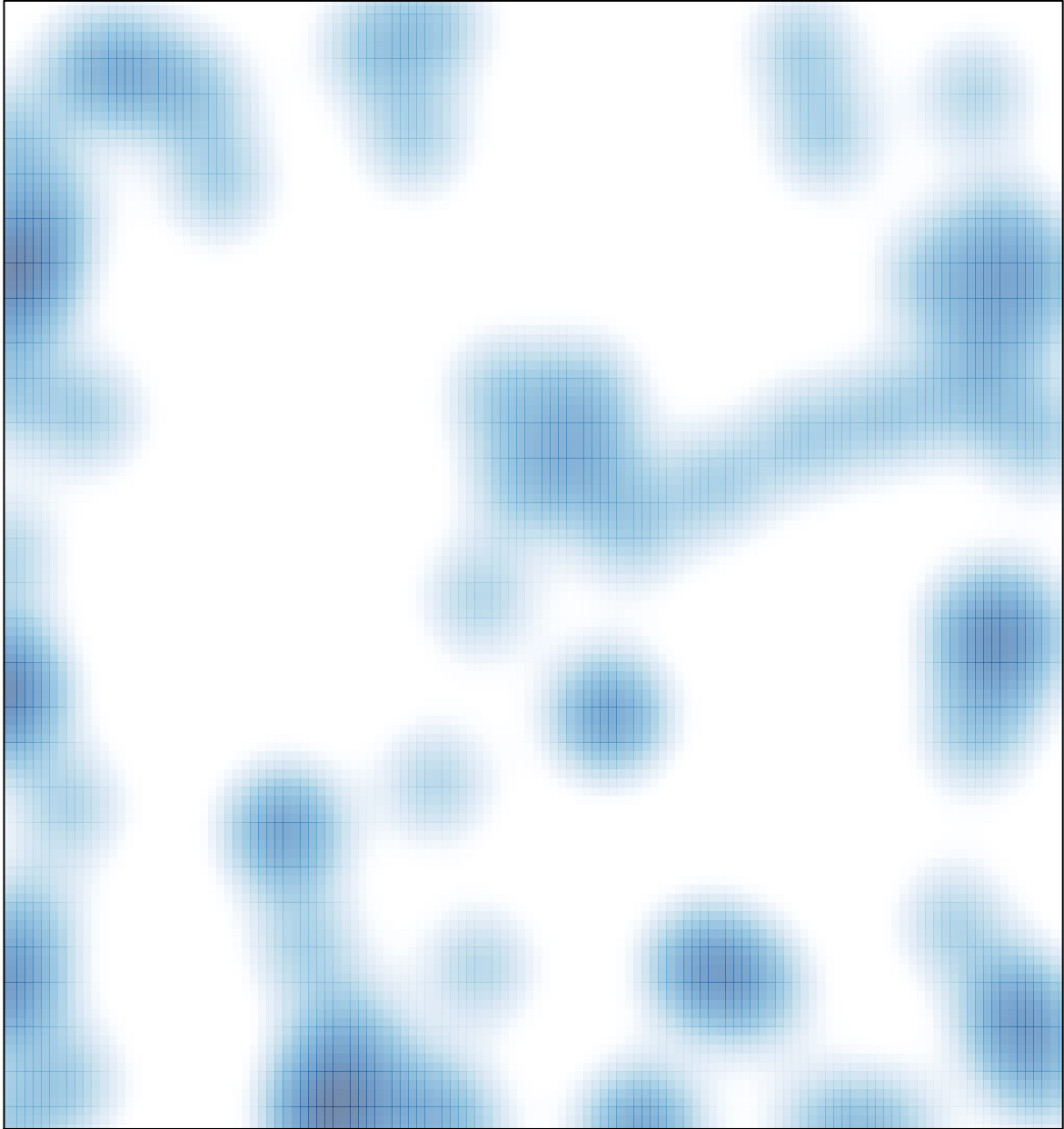


# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotB



# WILLSCHER\_GBM\_proteomics\_wtOnly\_SpotB



# features = 94 , max = 3