willscher_GBM_proteomics_wtOnly_SpotB
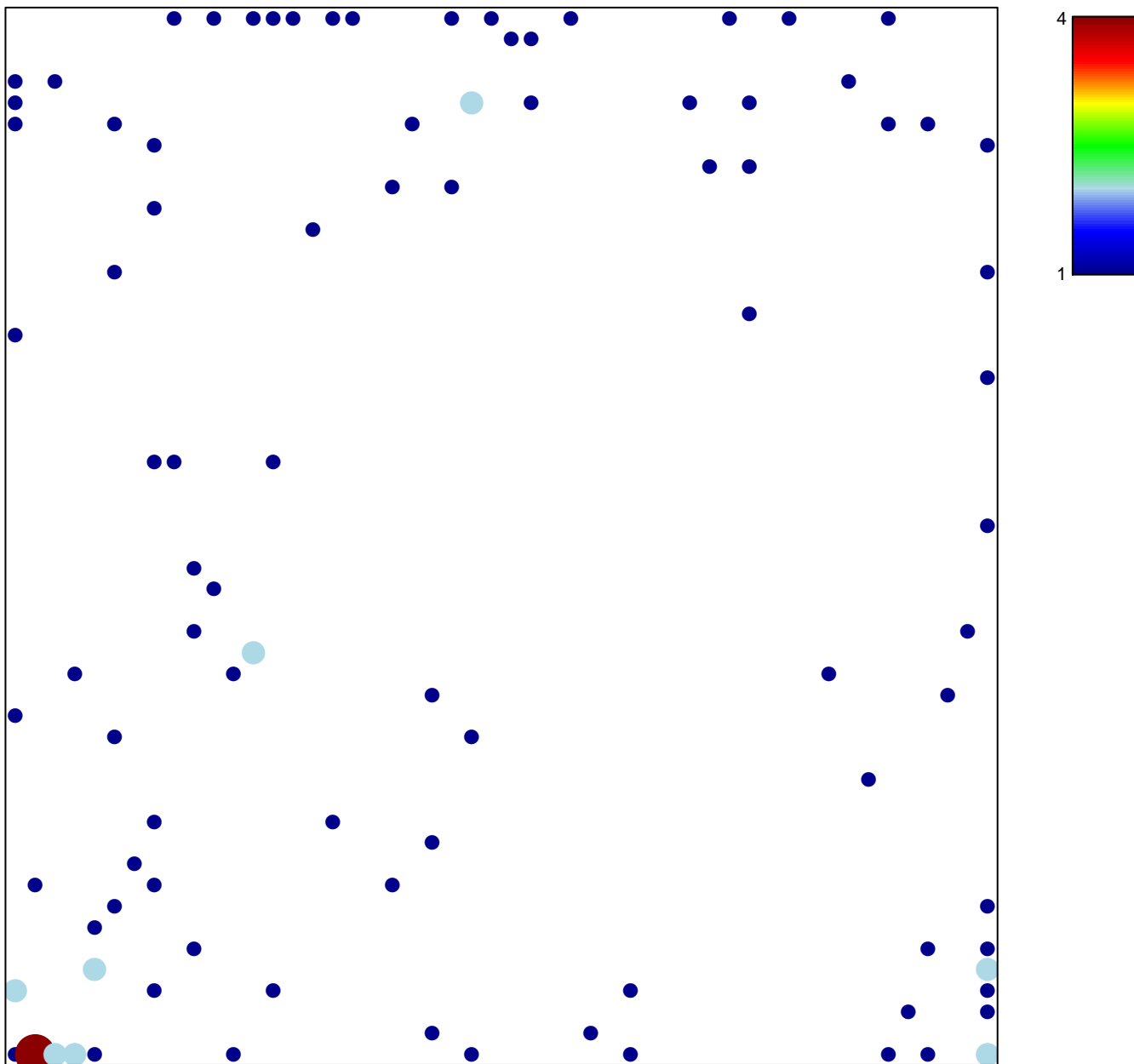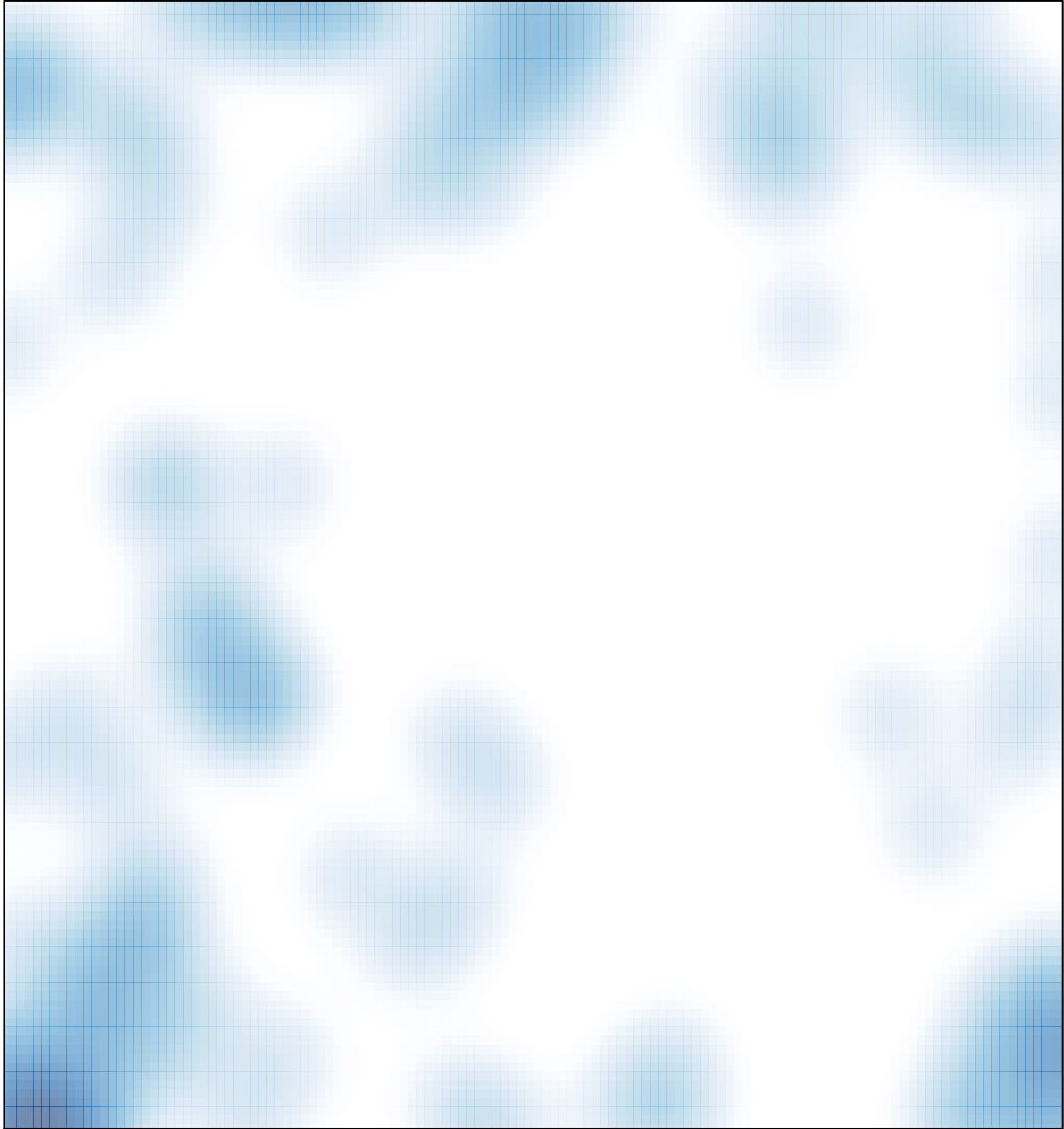
features = 105
chi−square p = 0.72

willscher_GBM_proteomics_wtOnly_SpotB

features = 105 , max = 4