willscher_GBM_proteomics_wtOnly_SpotJ
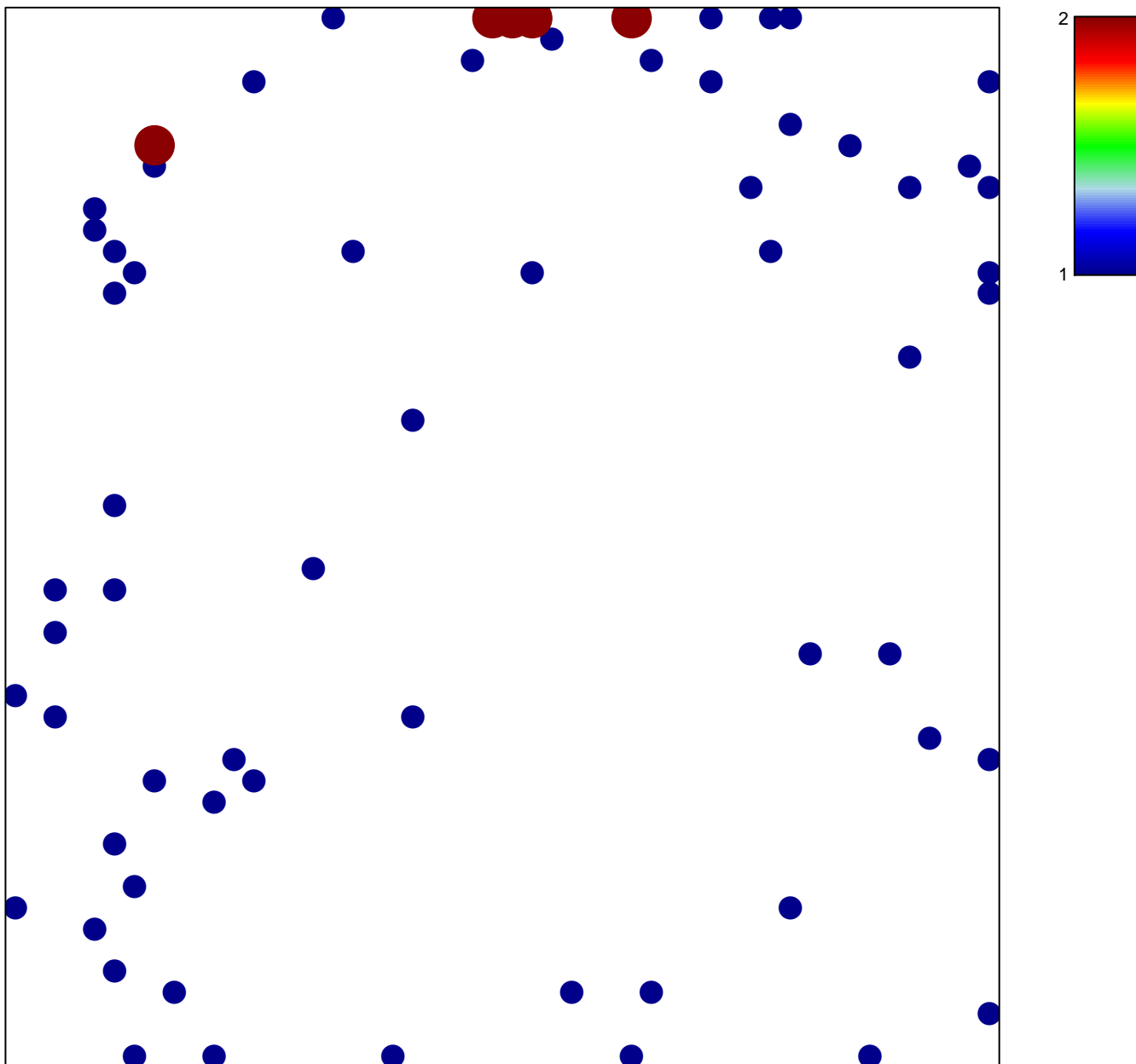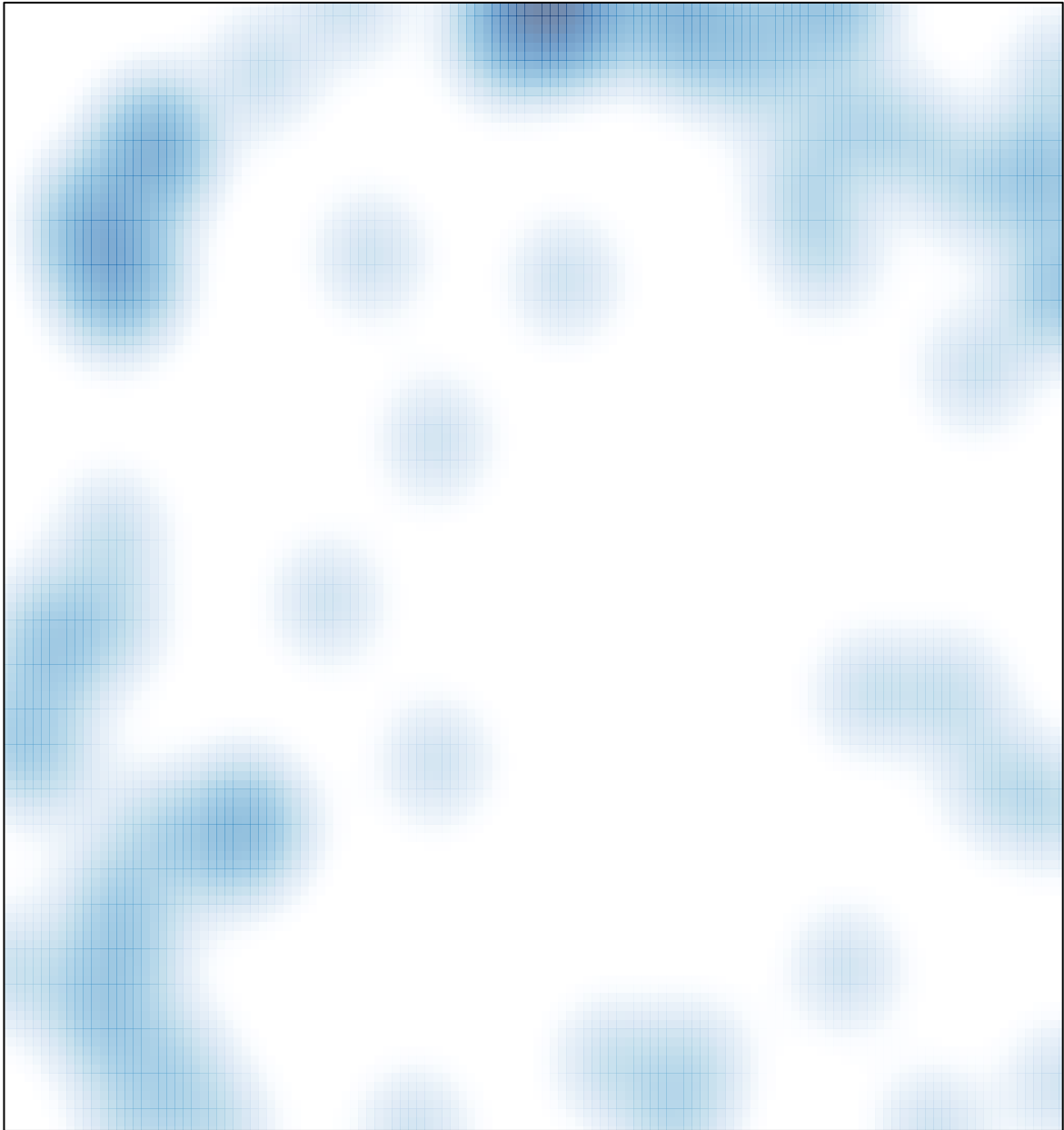
features = 70
chi−square p = 0.78

willscher_GBM_proteomics_wtOnly_SpotJ

features = 70 , max = 2