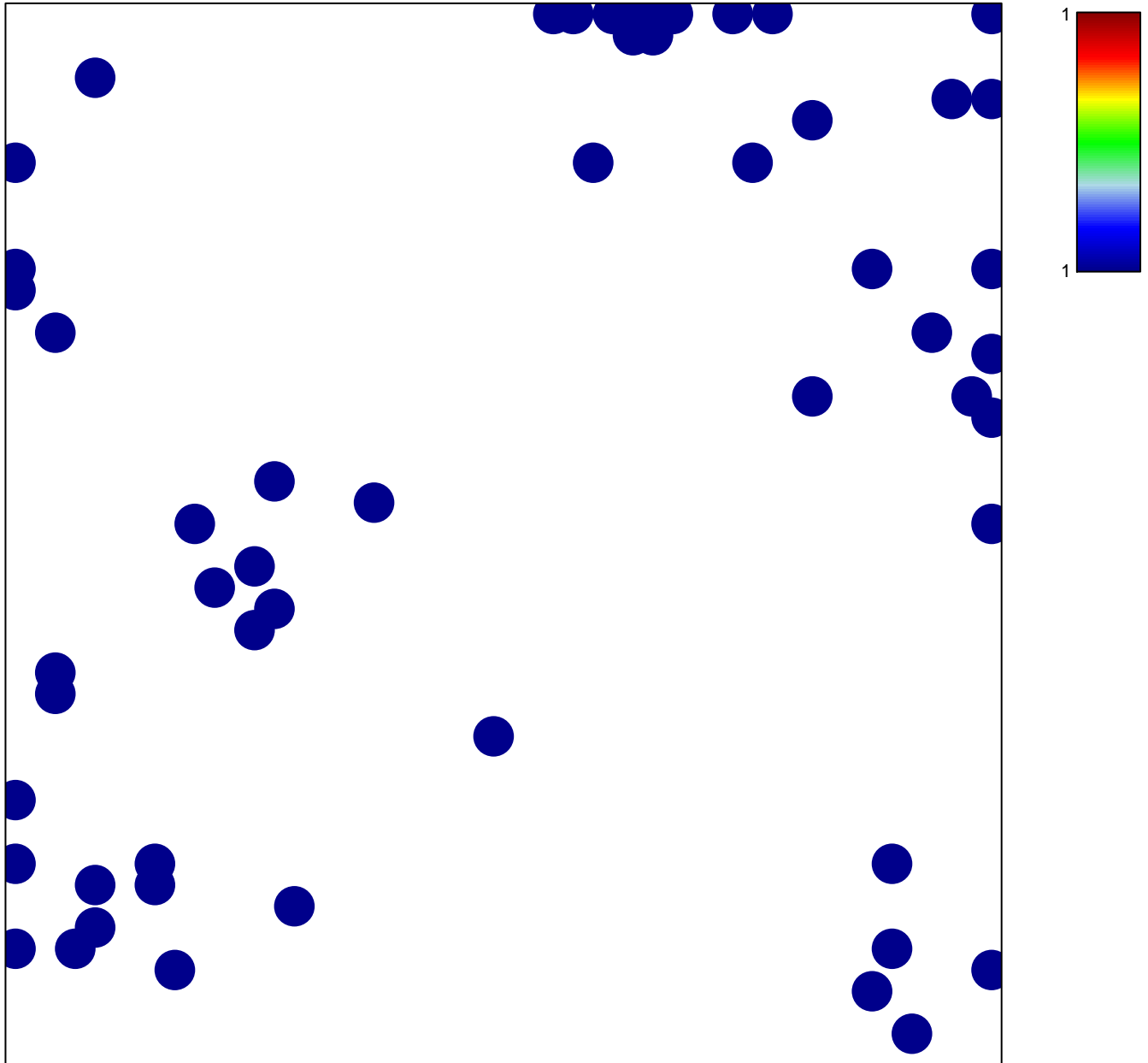
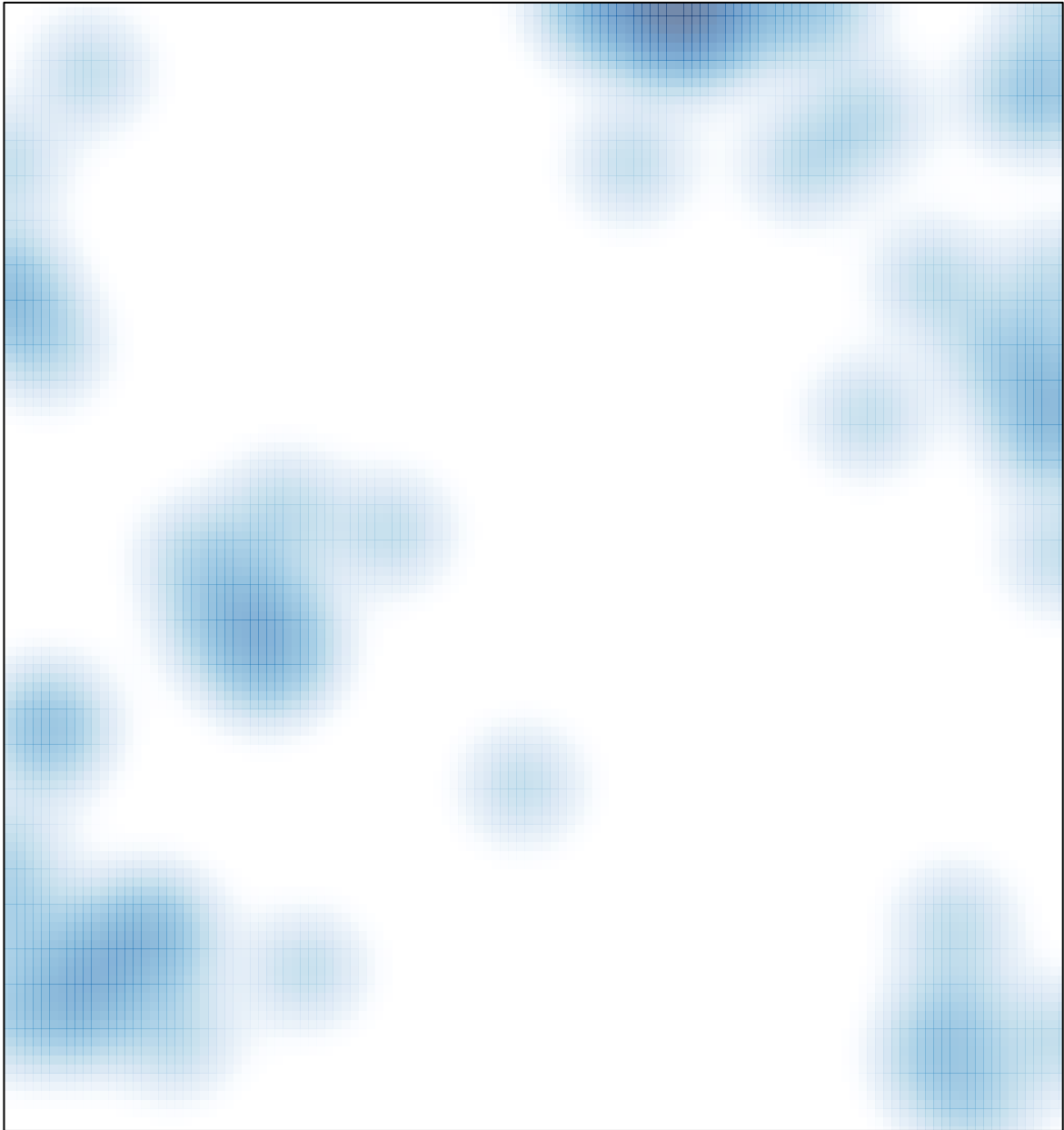


willscher_GBM_proteomics_wtOnly_SpotG



features = 54
chi-square p = 0.78

willscher_GBM_proteomics_wtOnly_SpotG



features = 54 , max = 1