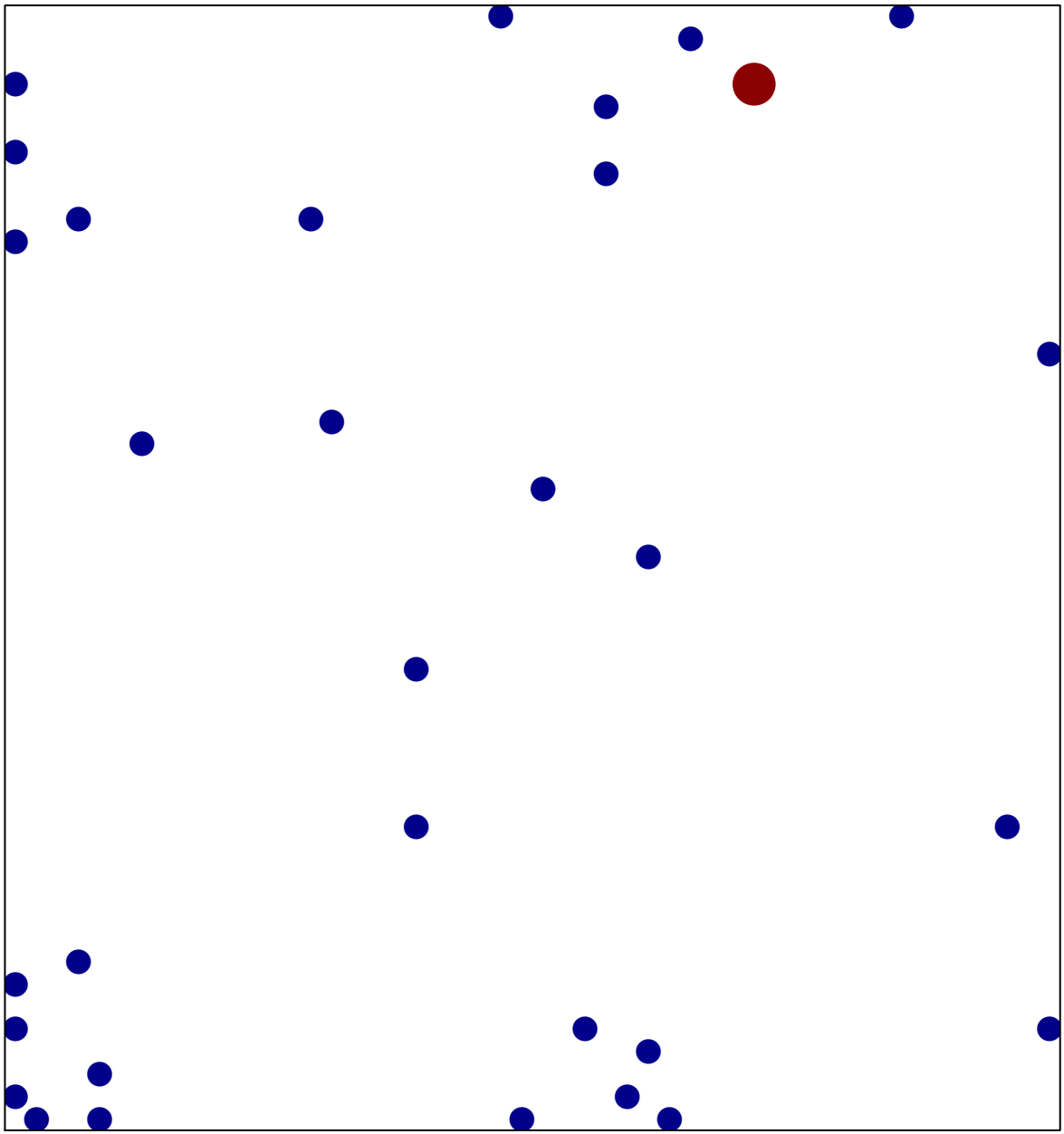
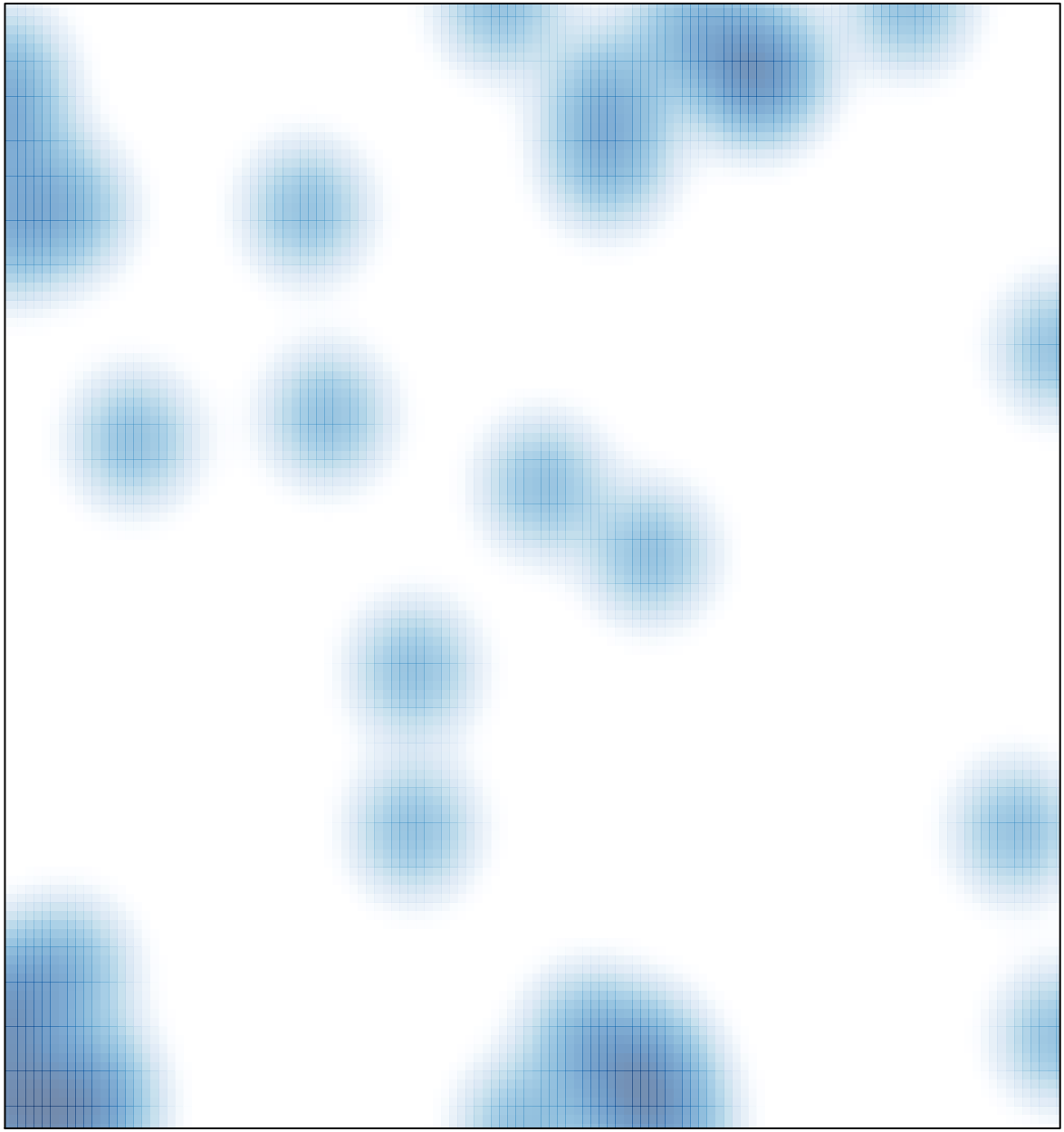
willscher_GBM_proteomics_wtOnly_SpotC



features = 33
chi−square p = 0.81

willscher_GBM_proteomics_wtOnly_SpotC



features = 33 , max = 2