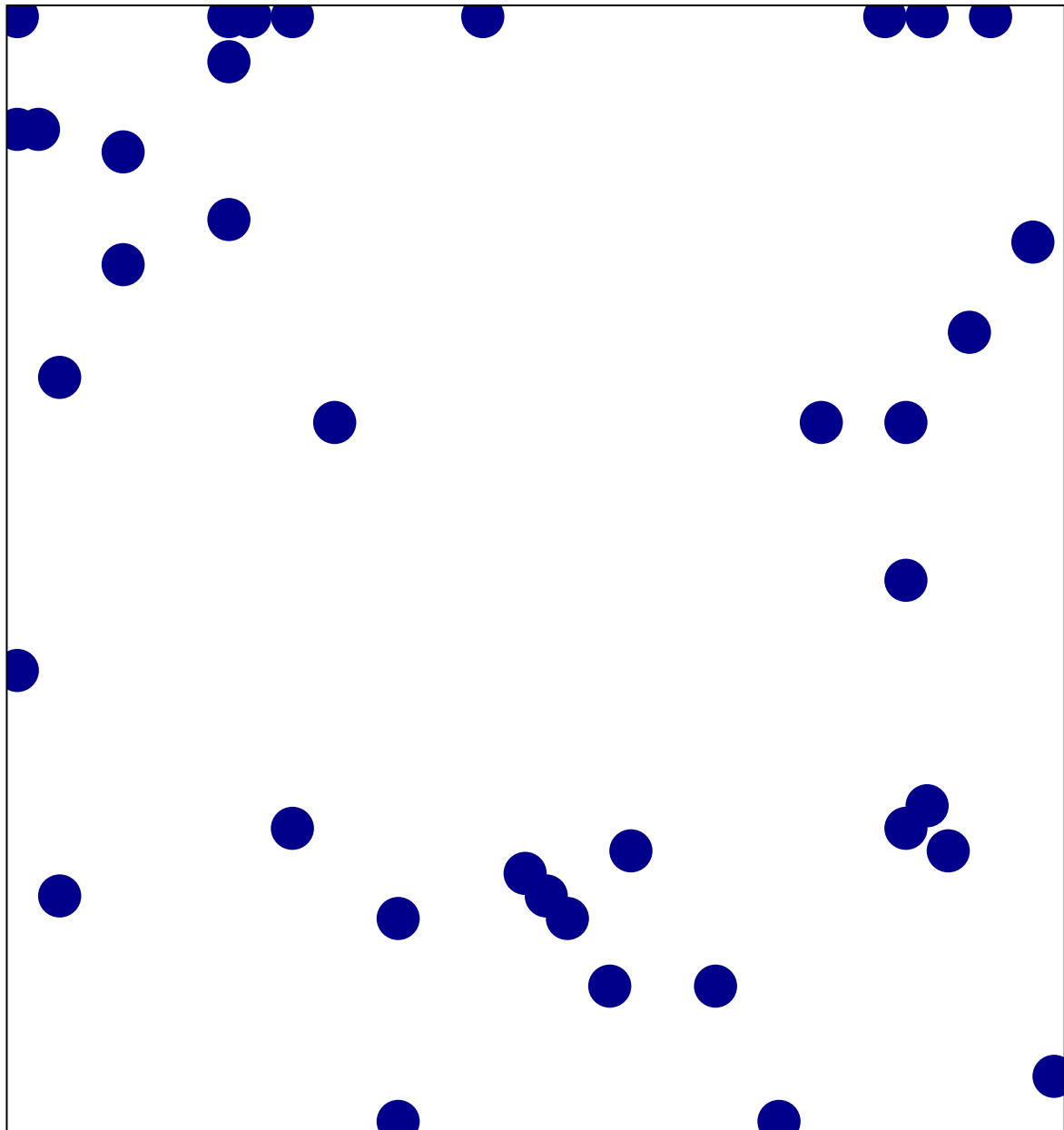
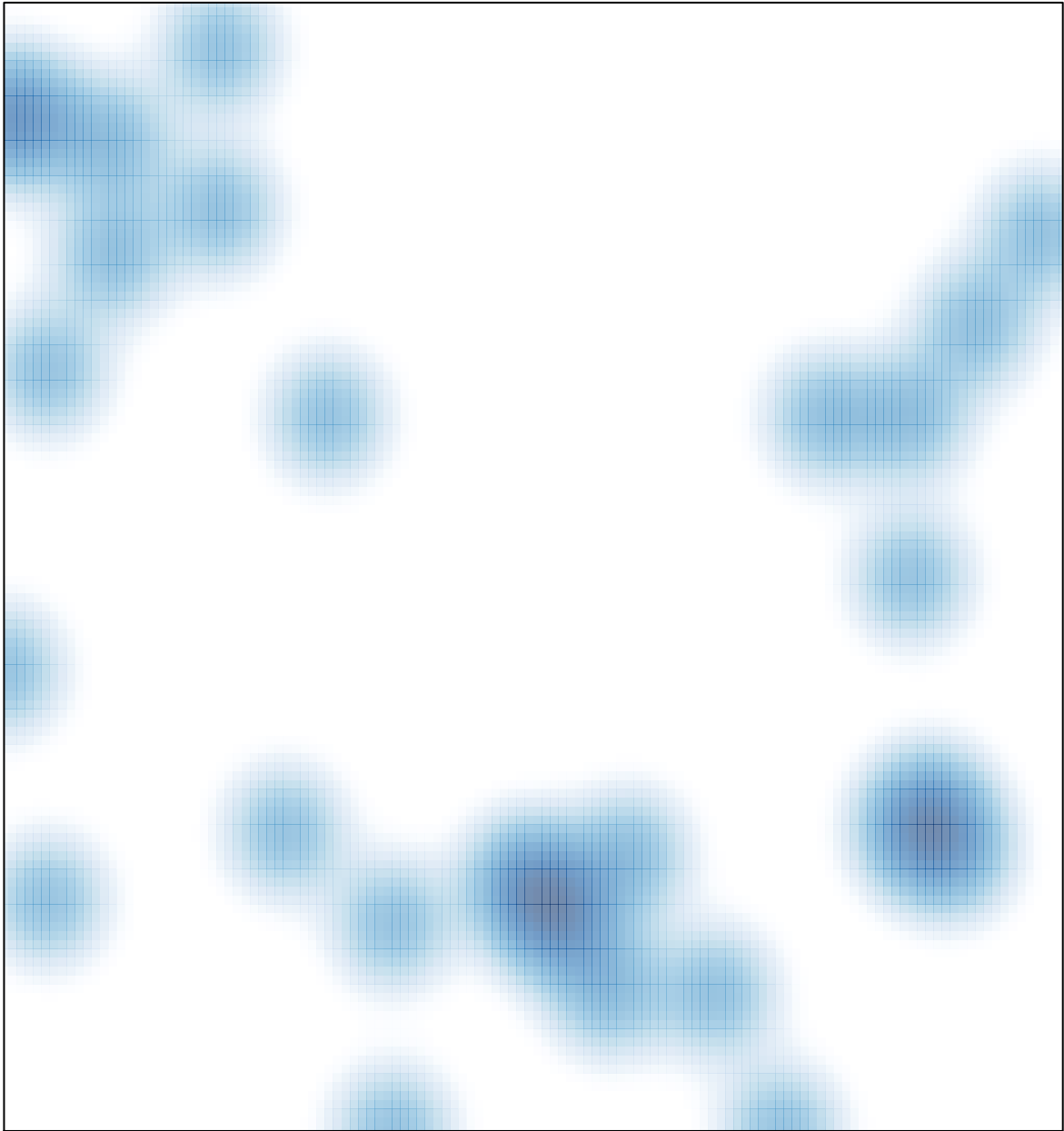


WILLSCHER_GBM_proteomics_wtOnly_SpotH



features = 37
chi-square p = 0.8

WILLSCHER_GBM_proteomics_wtOnly_SpotH



features = 37 , max = 1