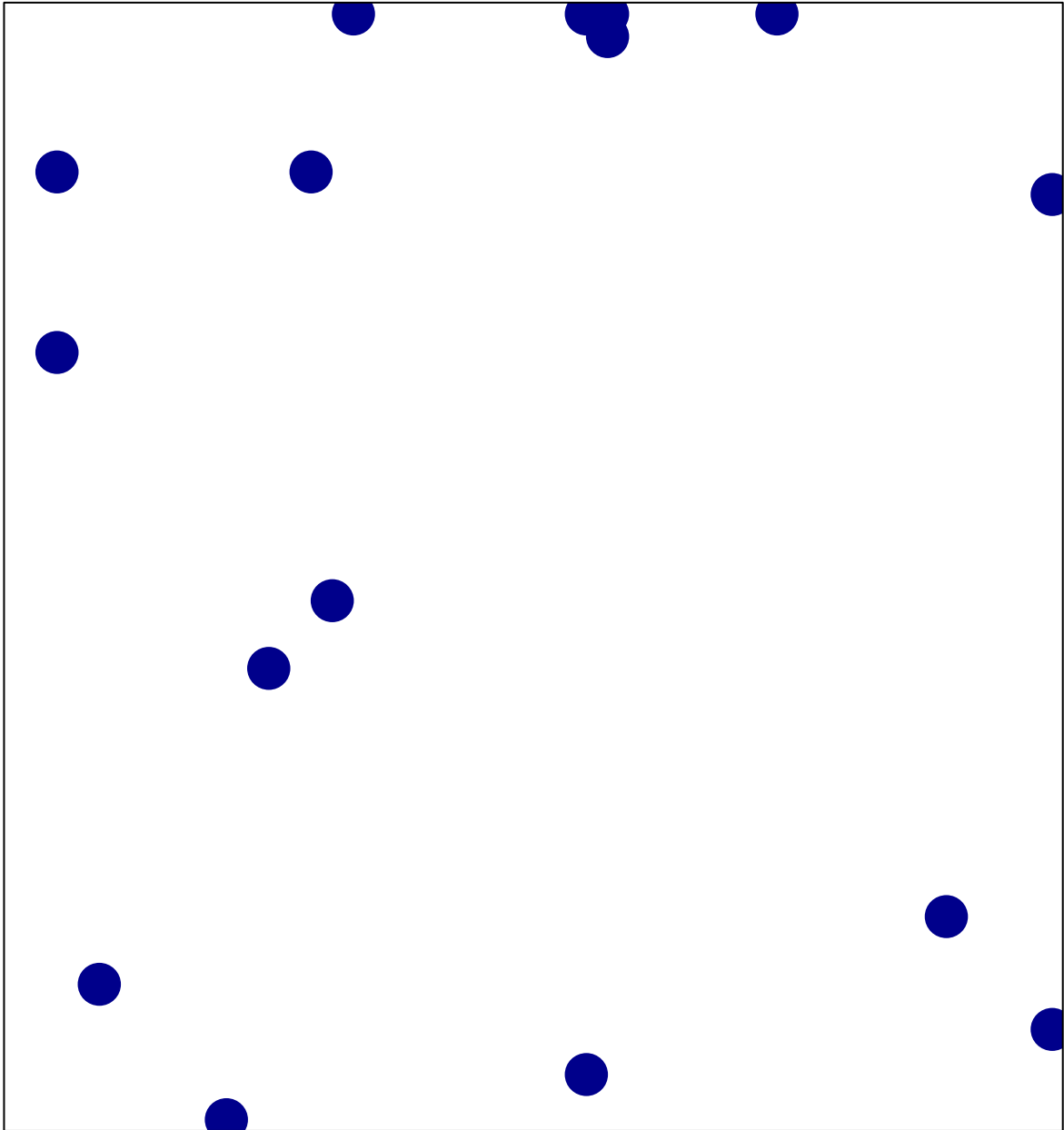
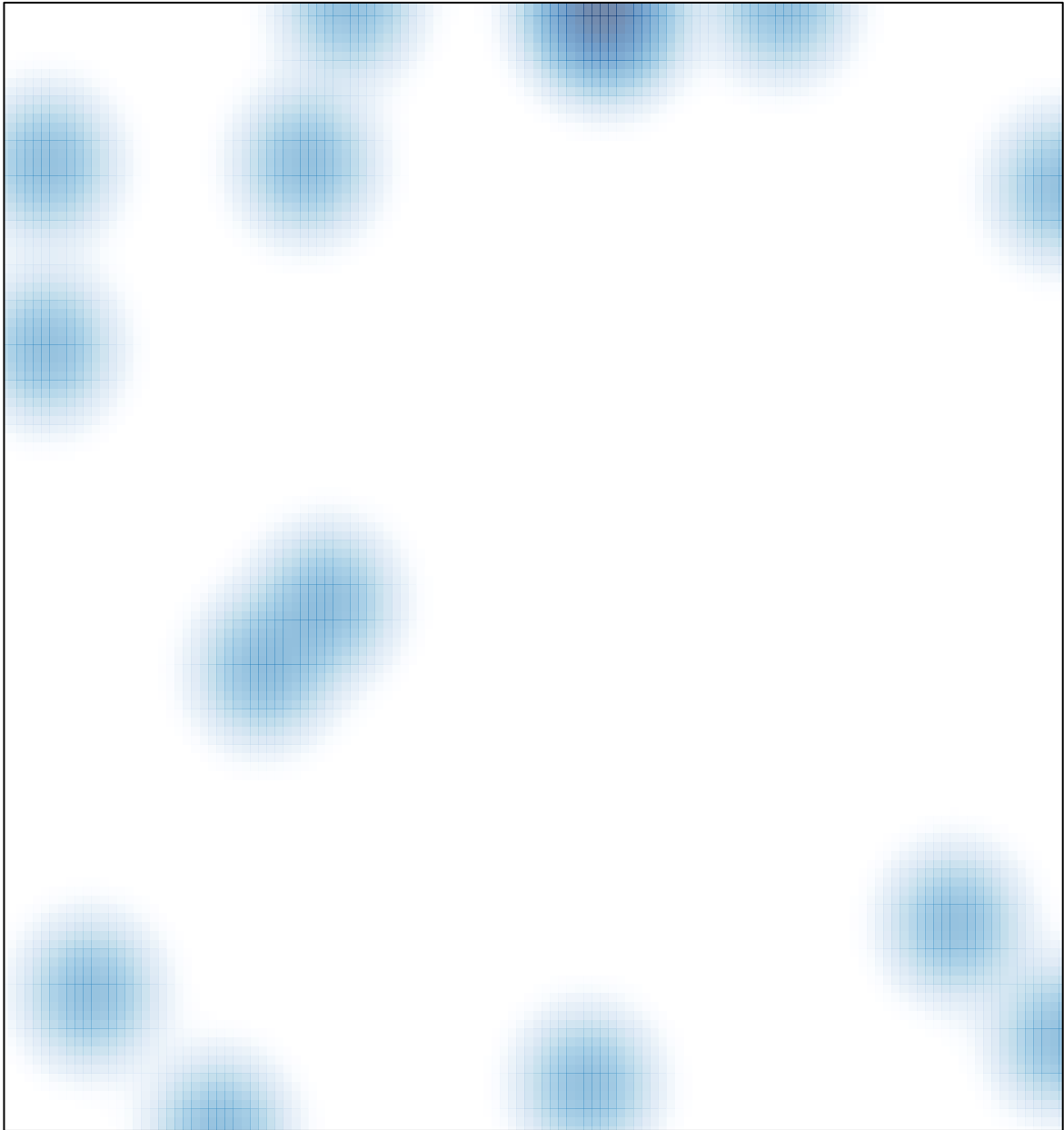


willscher_GBM_STSwt_proteomics-O_UP



features = 16
chi-square p = 0.82

willscher_GBM_STSwt_proteomics-O_UP



features = 16 , max = 1