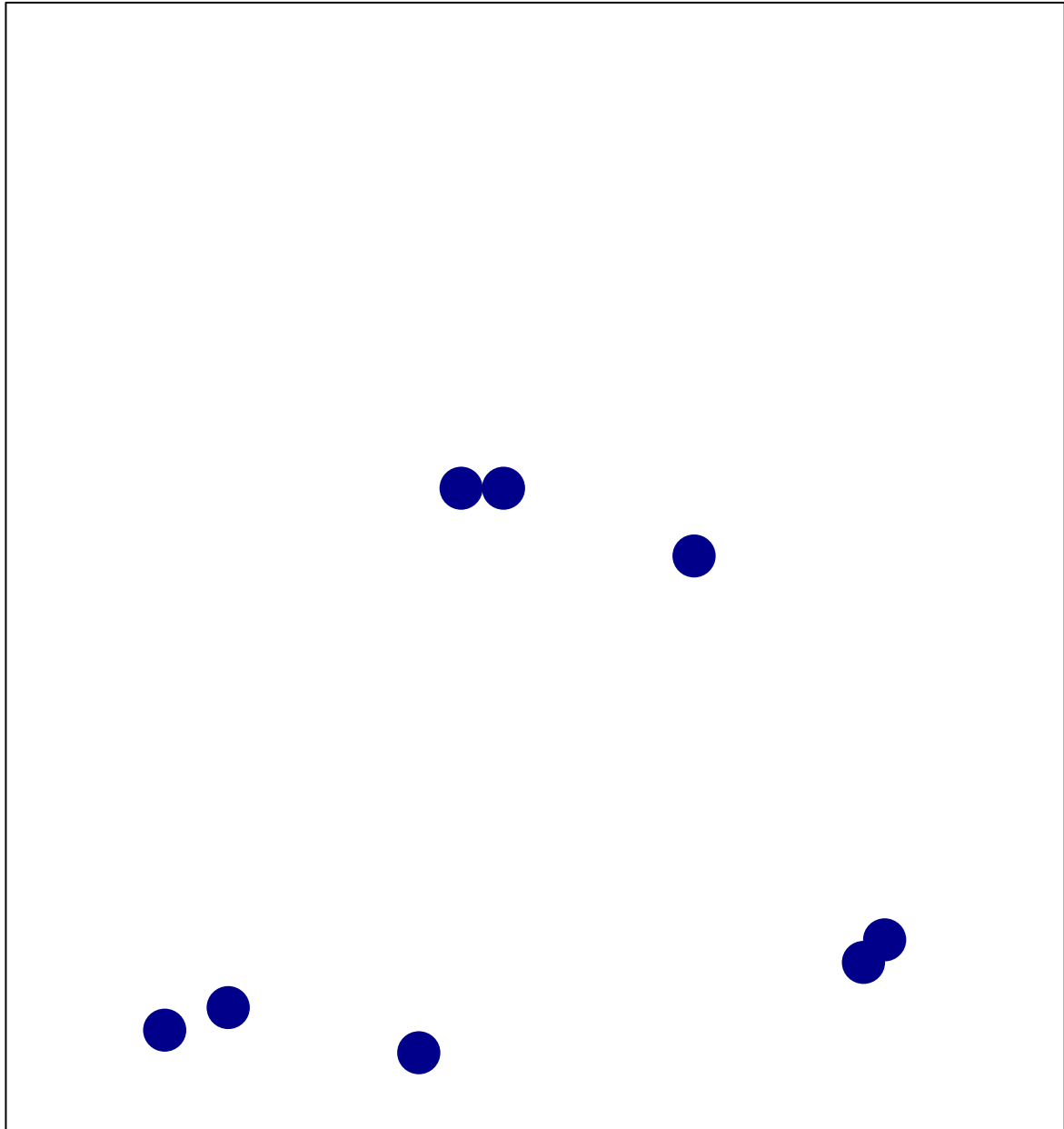
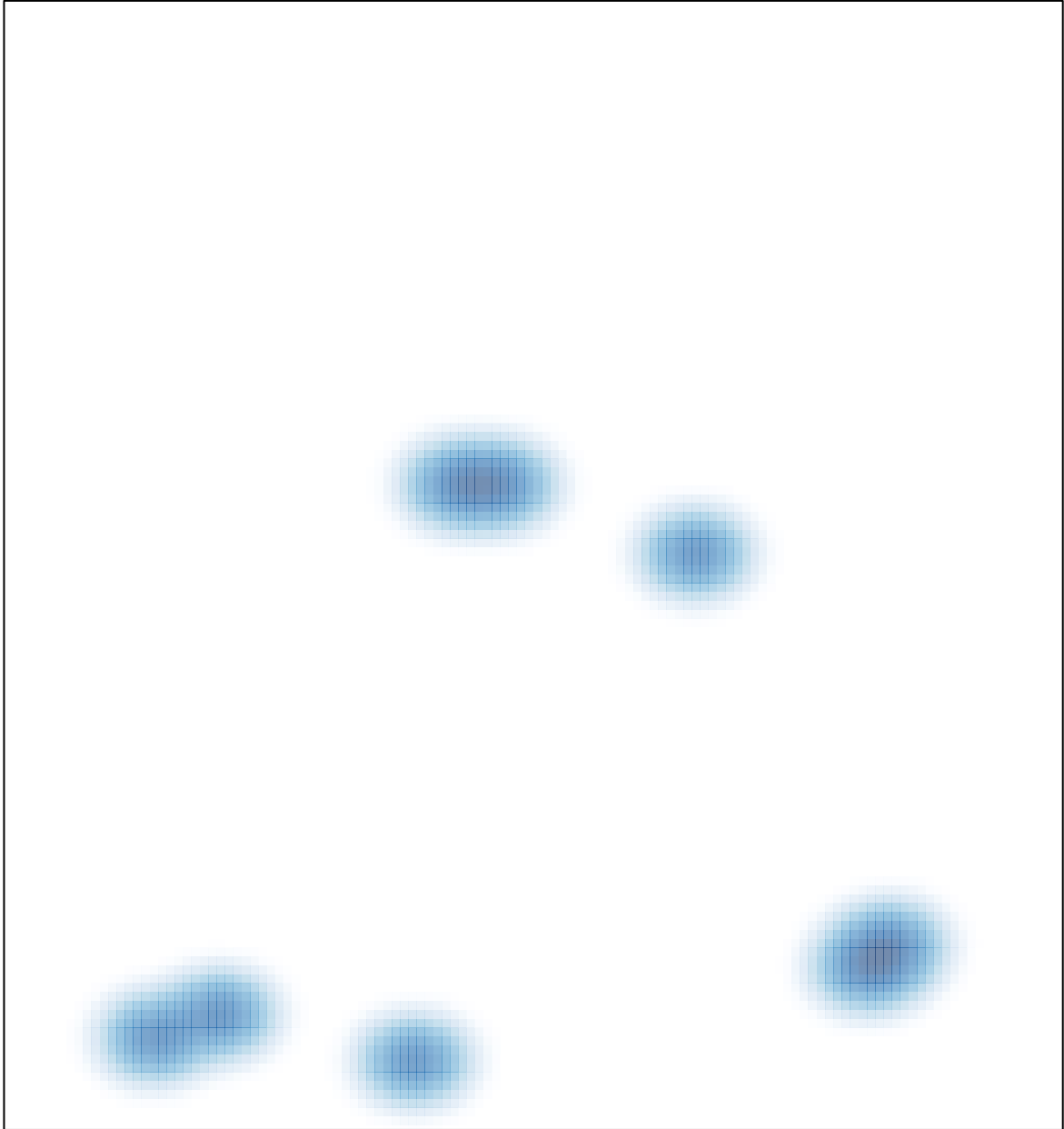


willscher_GBM_STSwt_proteomics-N_UP



features = 8
chi-square p = 0.84

willscher_GBM_STSwt_proteomics-N_UP



features = 8 , max = 1