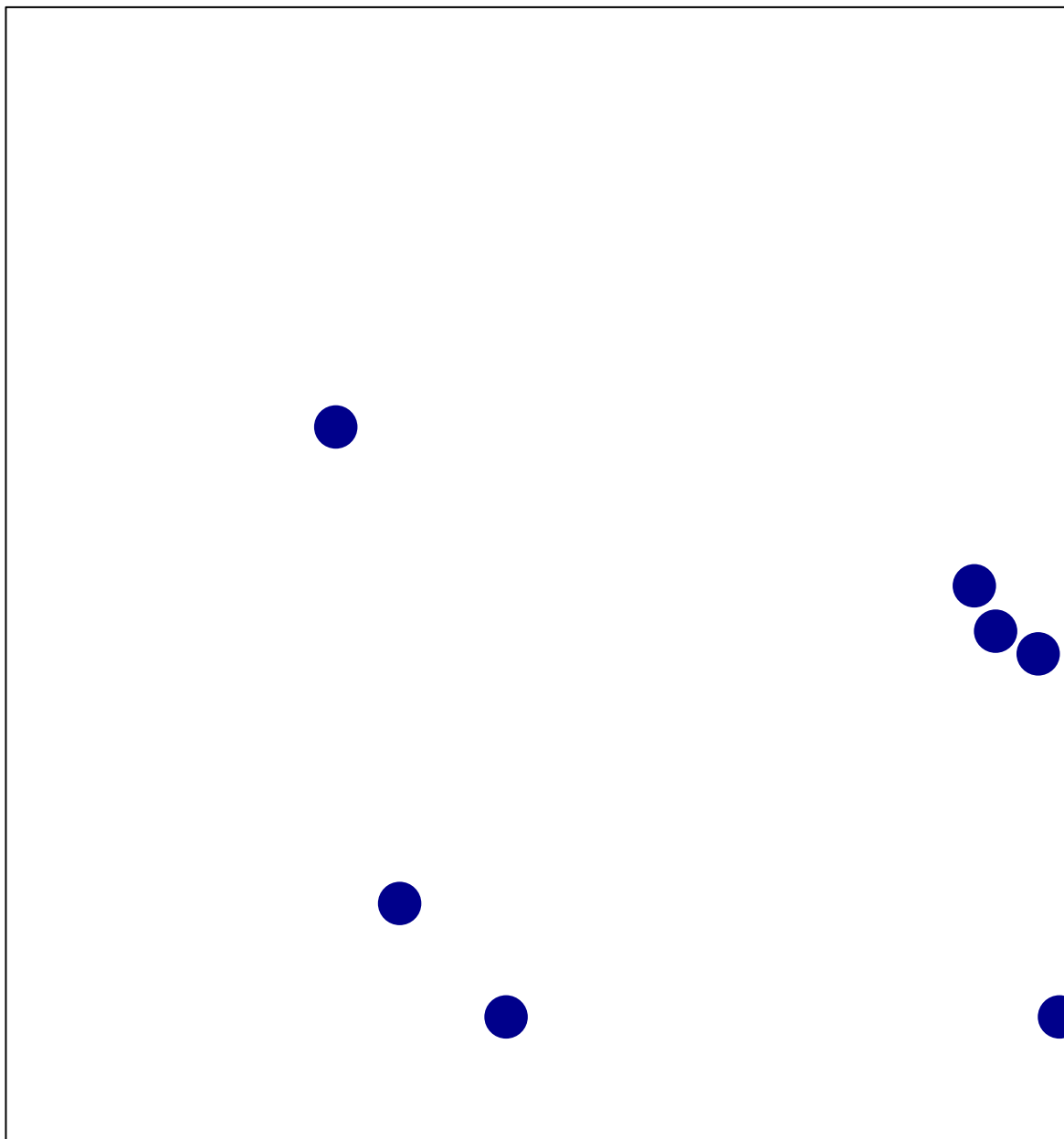
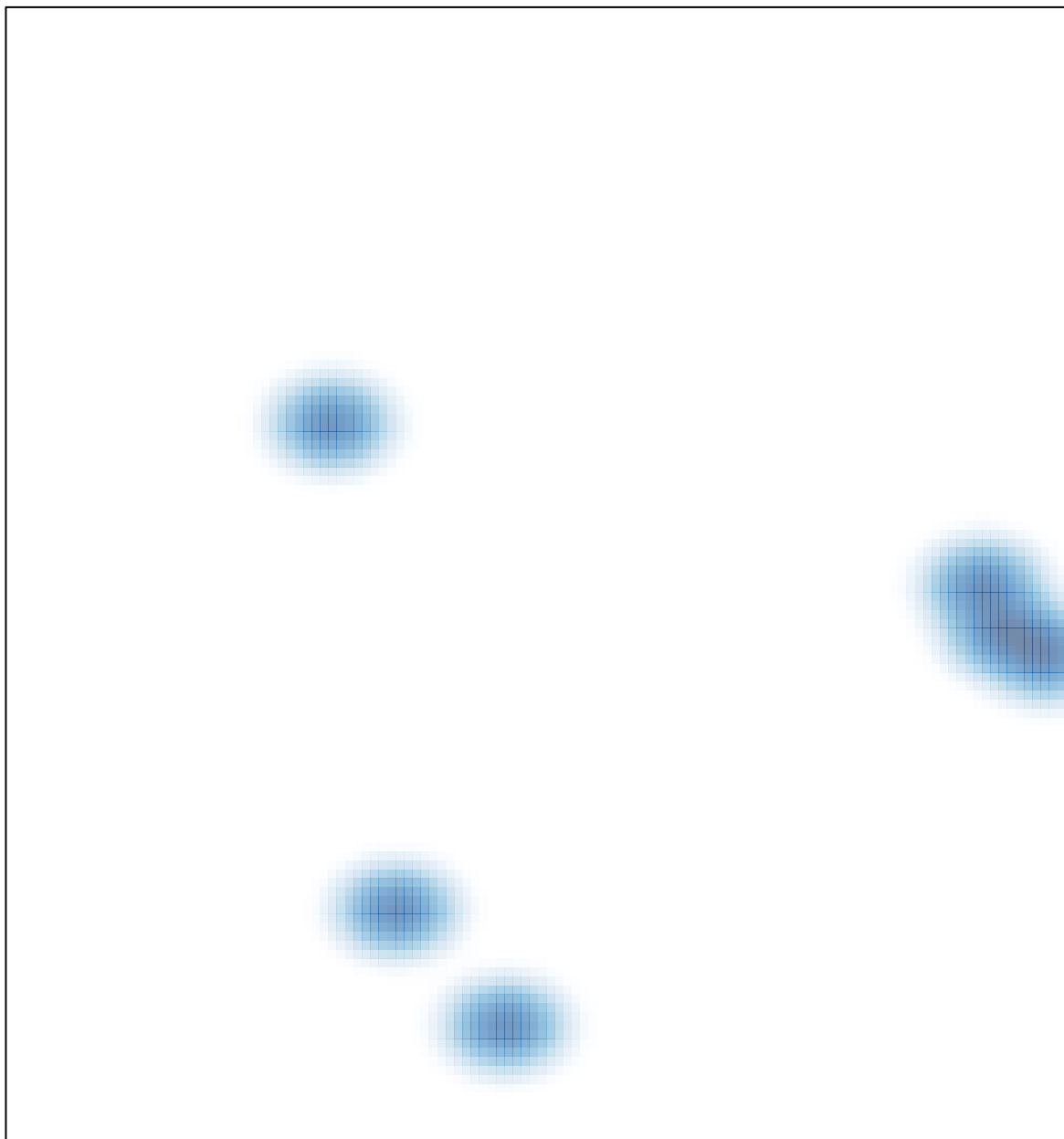


WILLSCHER_GBM_LTSwt_proteomics-G_UP



features = 7
chi-square p = 0.84

WILLSCHER_GBM_LTSwt_proteomics-G_UP



features = 7 , max = 1