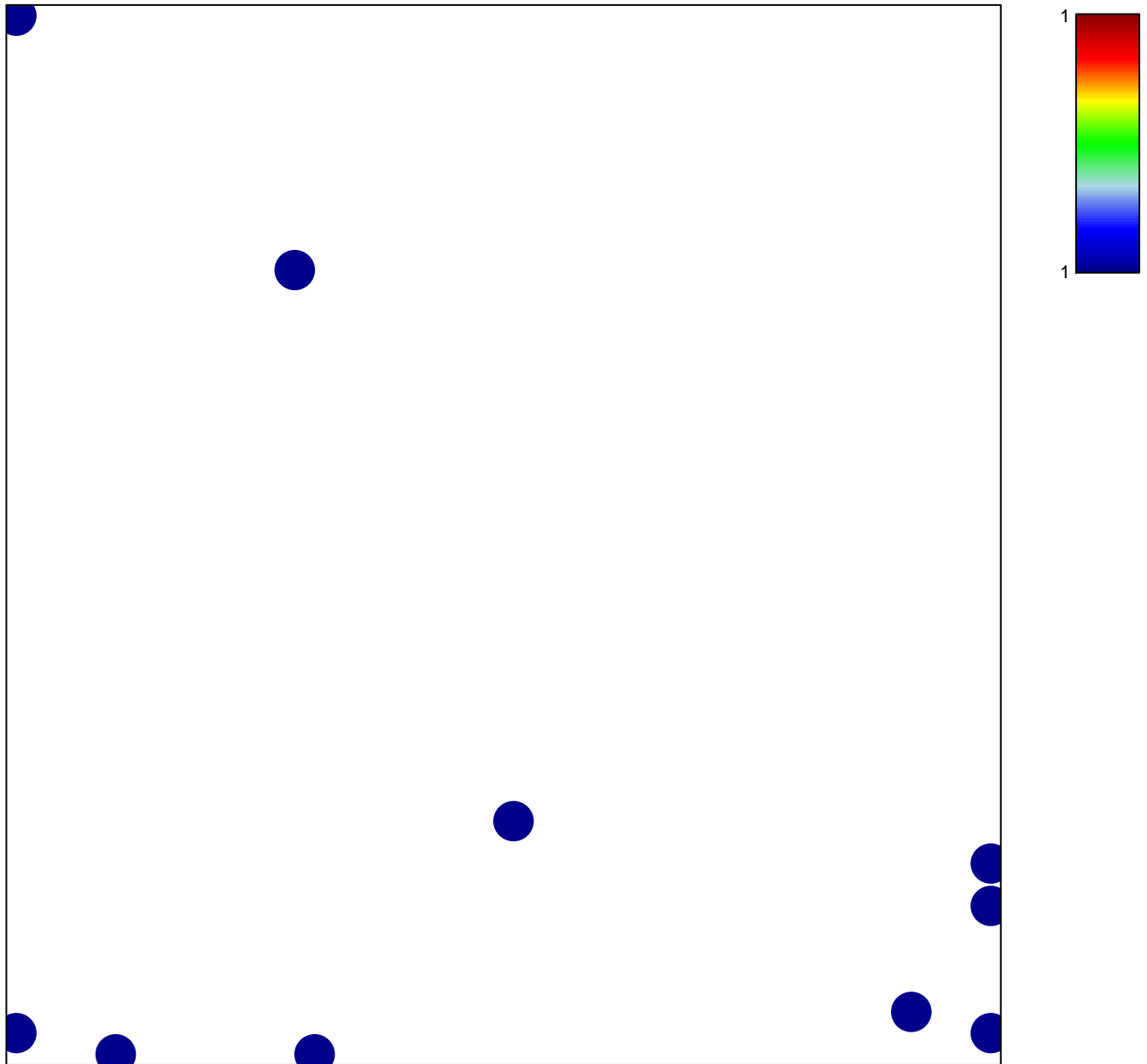
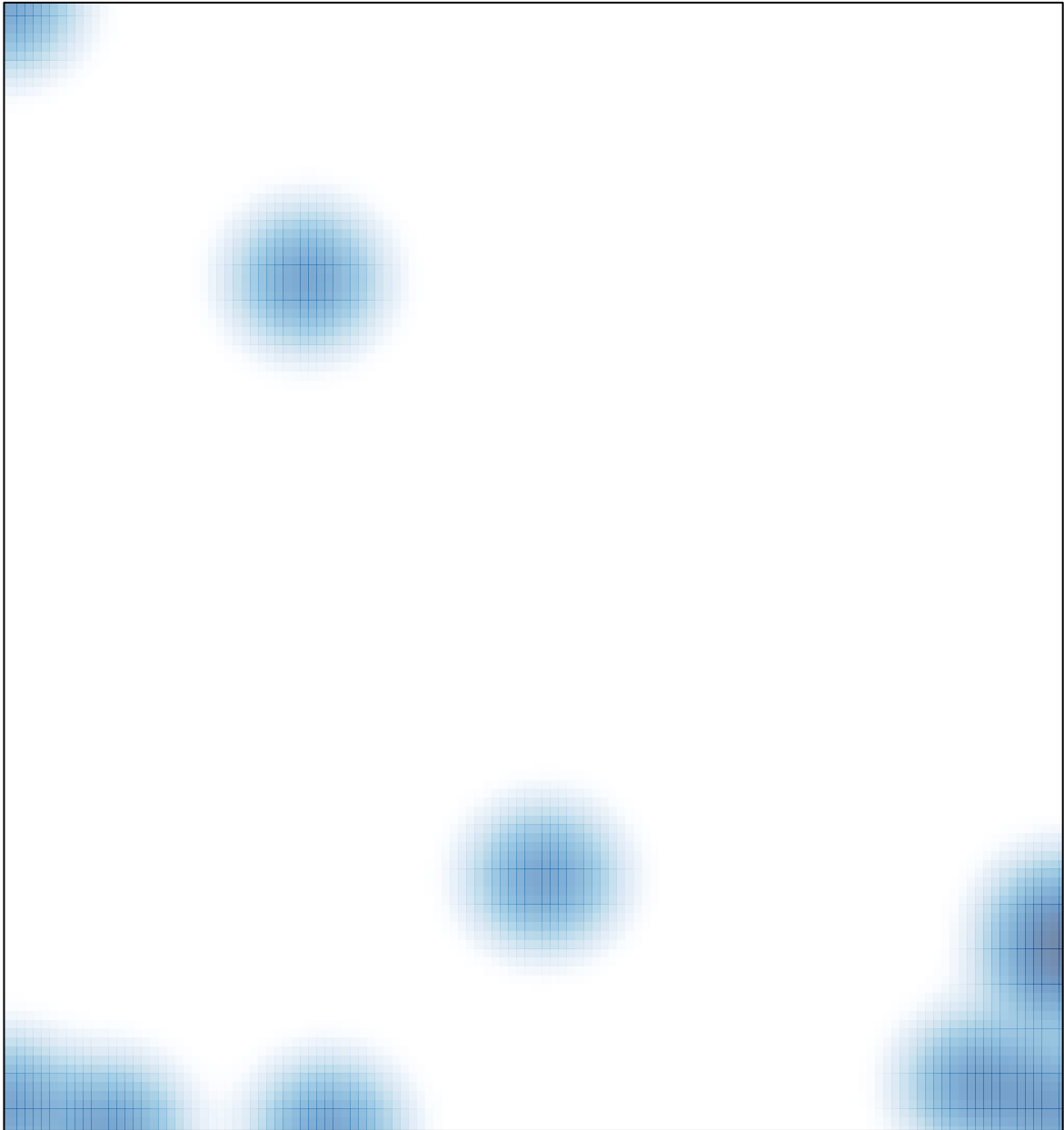


willscher_GBM_LTSwt_proteomics-G_UP



features = 10
chi-square p = 0.81

willscher_GBM_LTSwt_proteomics-G_UP



features = 10 , max = 1