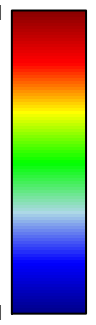
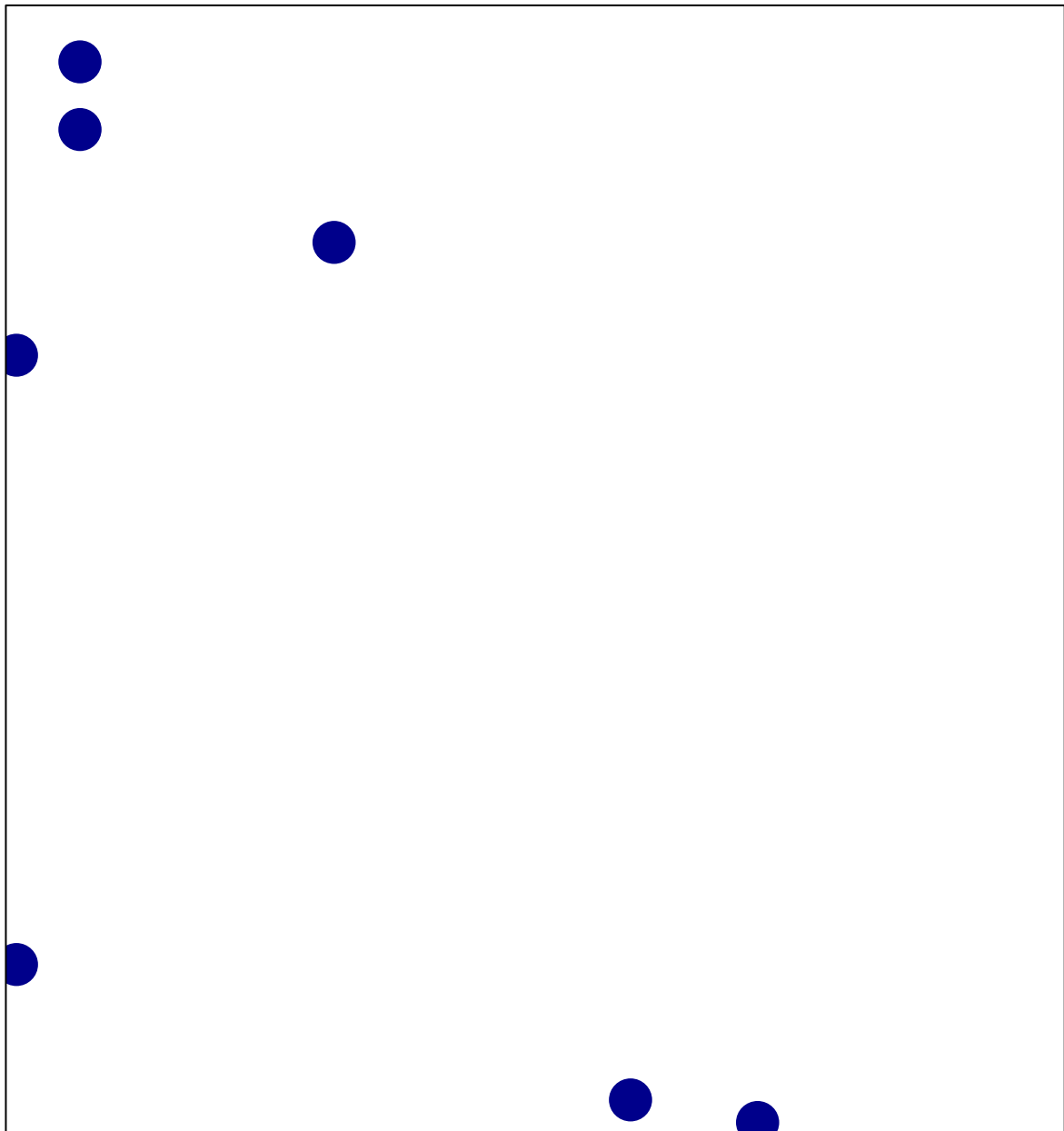
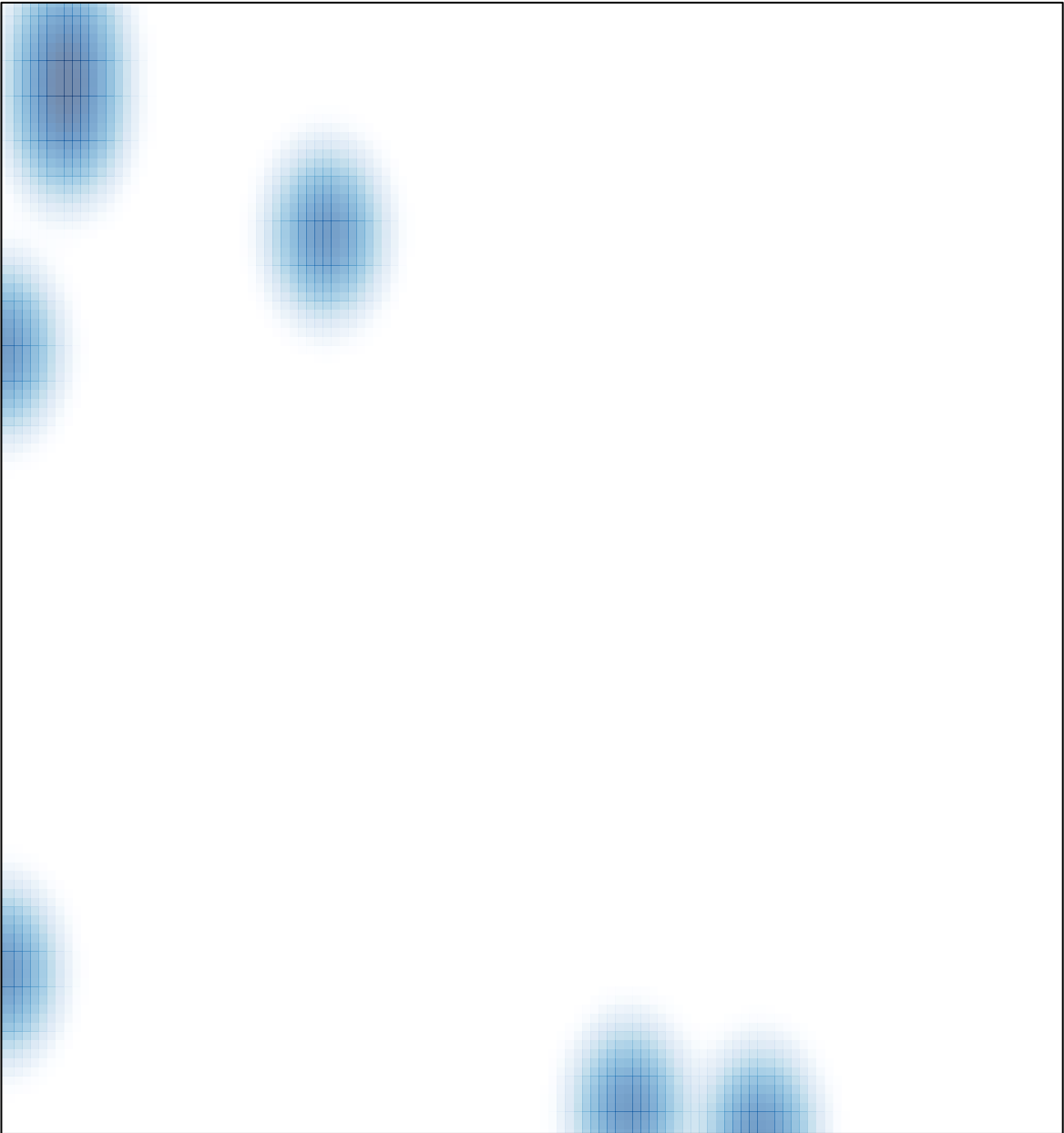


willscher_GBM_LTSmut_proteomics-P_DOWN



features = 7
chi-square p = 0.83

willscher_GBM_LTSmut_proteomics-P_DOWN



features = 7 , max = 1