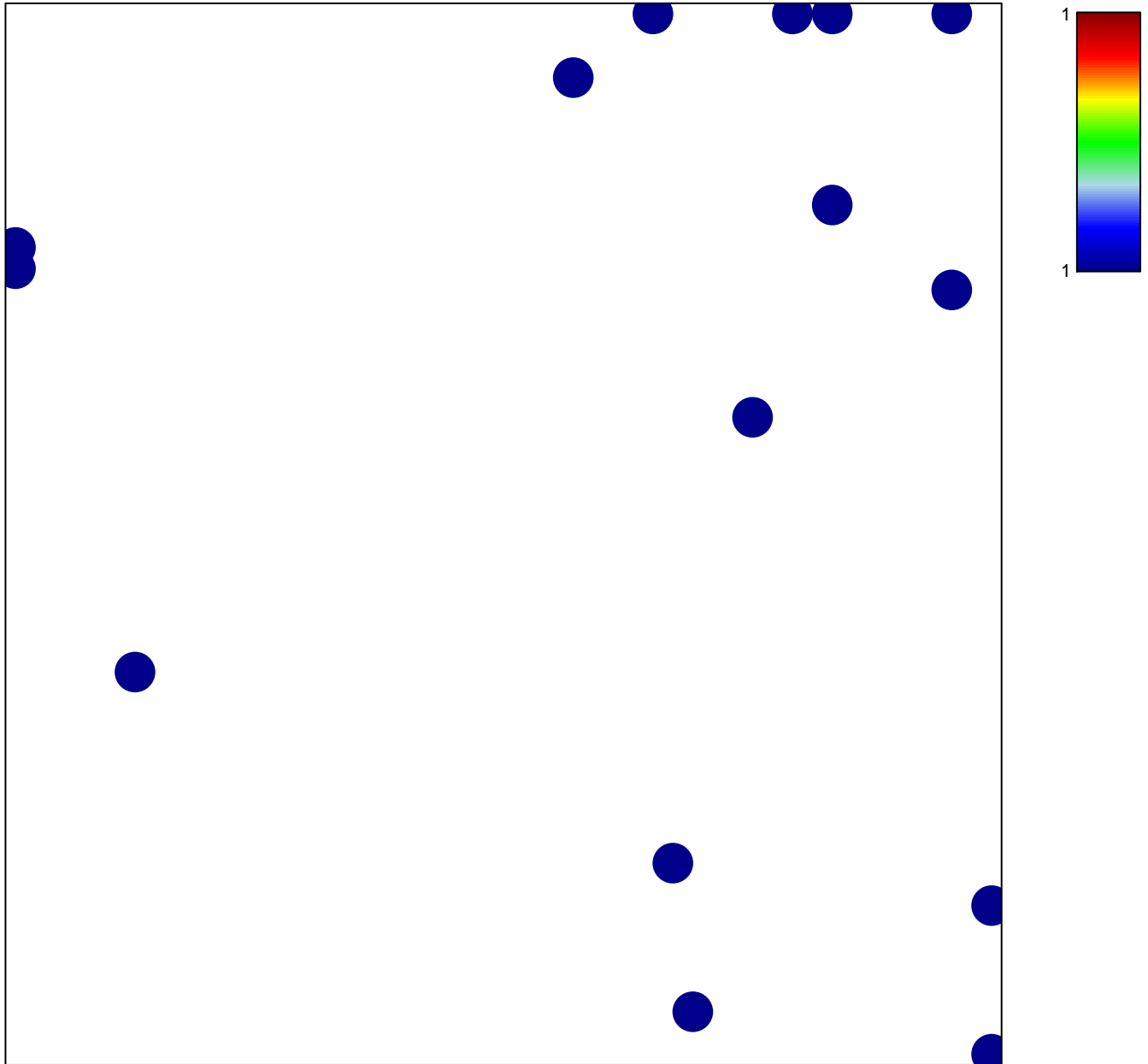


WILLSCHER_GBM_STSwt_proteomics-O_UP



features = 15
chi-square p = 0.82

WILLSCHER_GBM_STSwt_proteomics-O_UP



features = 15 , max = 1