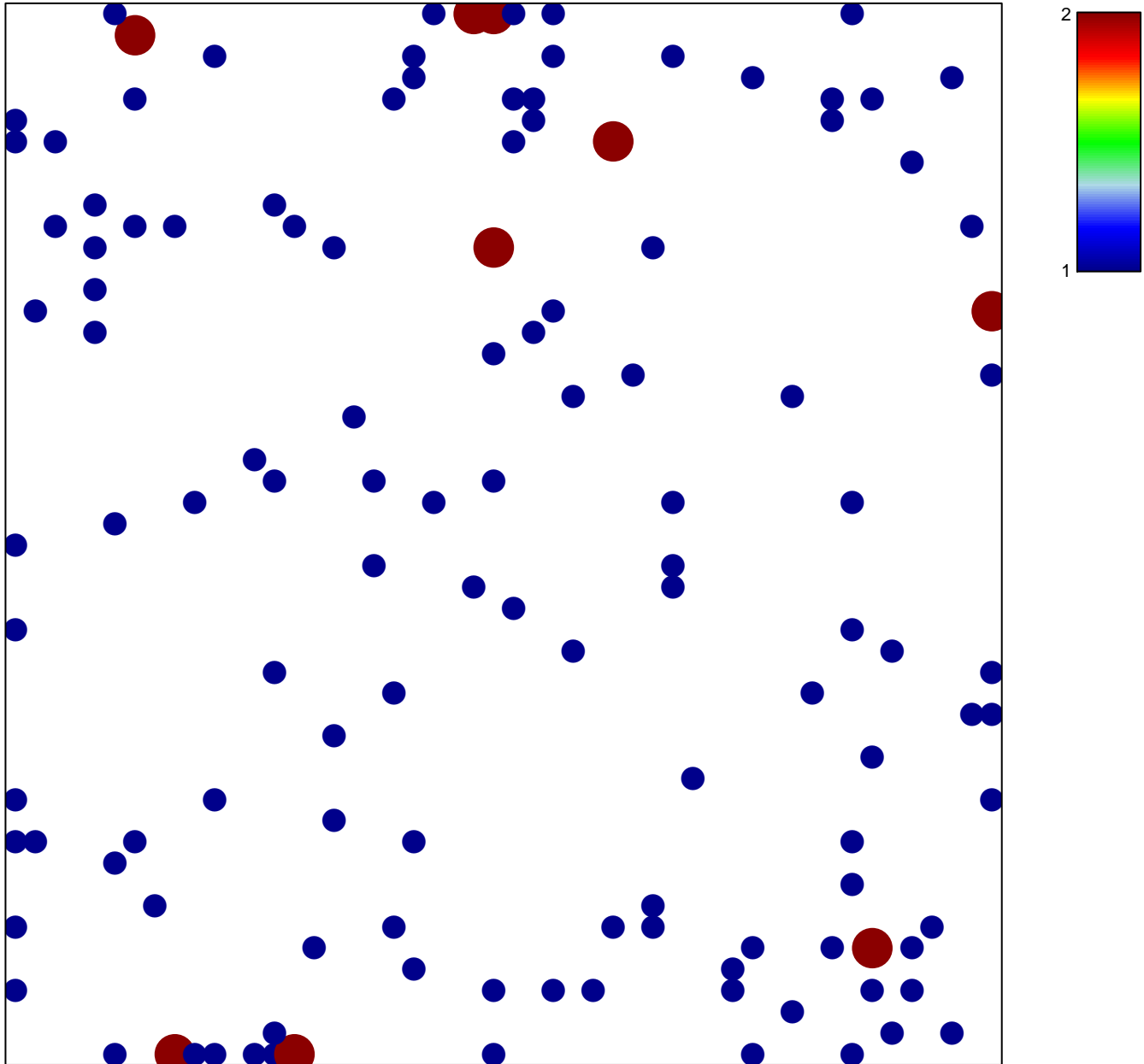
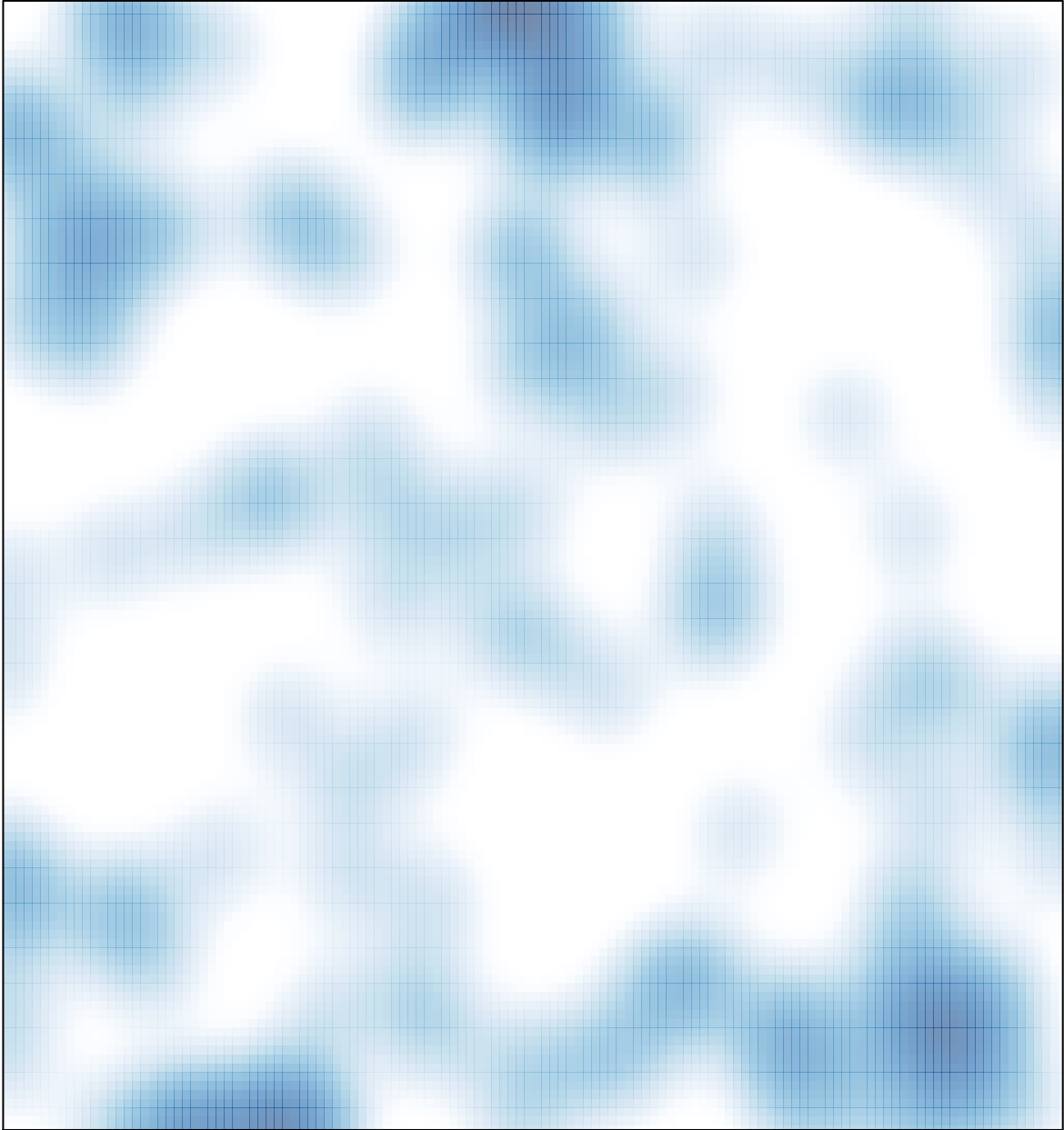


protein dephosphorylation



features = 135
chi-square p = 0.8

protein dephosphorylation



features = 135 , max = 2