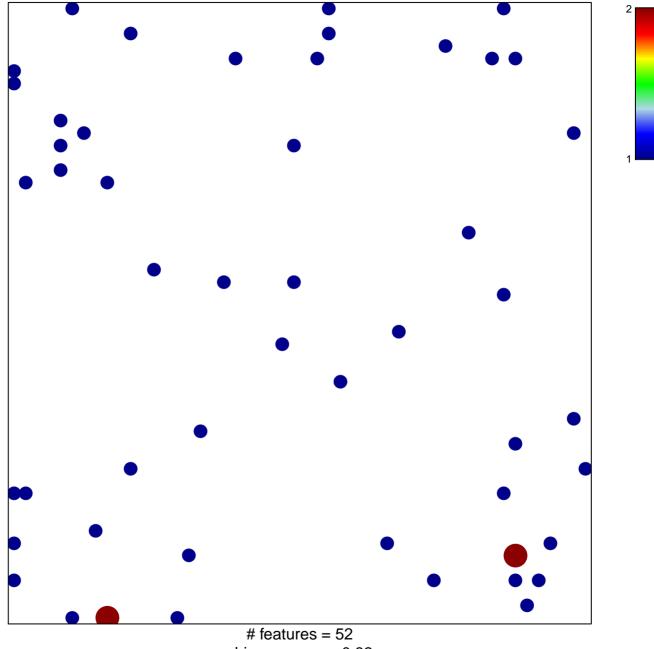
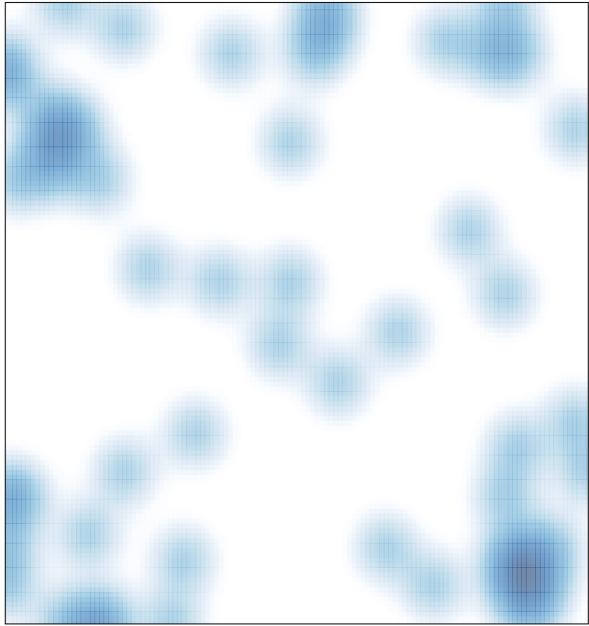
peptidyl-tyrosine dephosphorylation



chi-square p = 0.82

peptidyl-tyrosine dephosphorylation



features = 52, max = 2