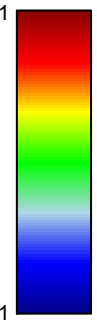
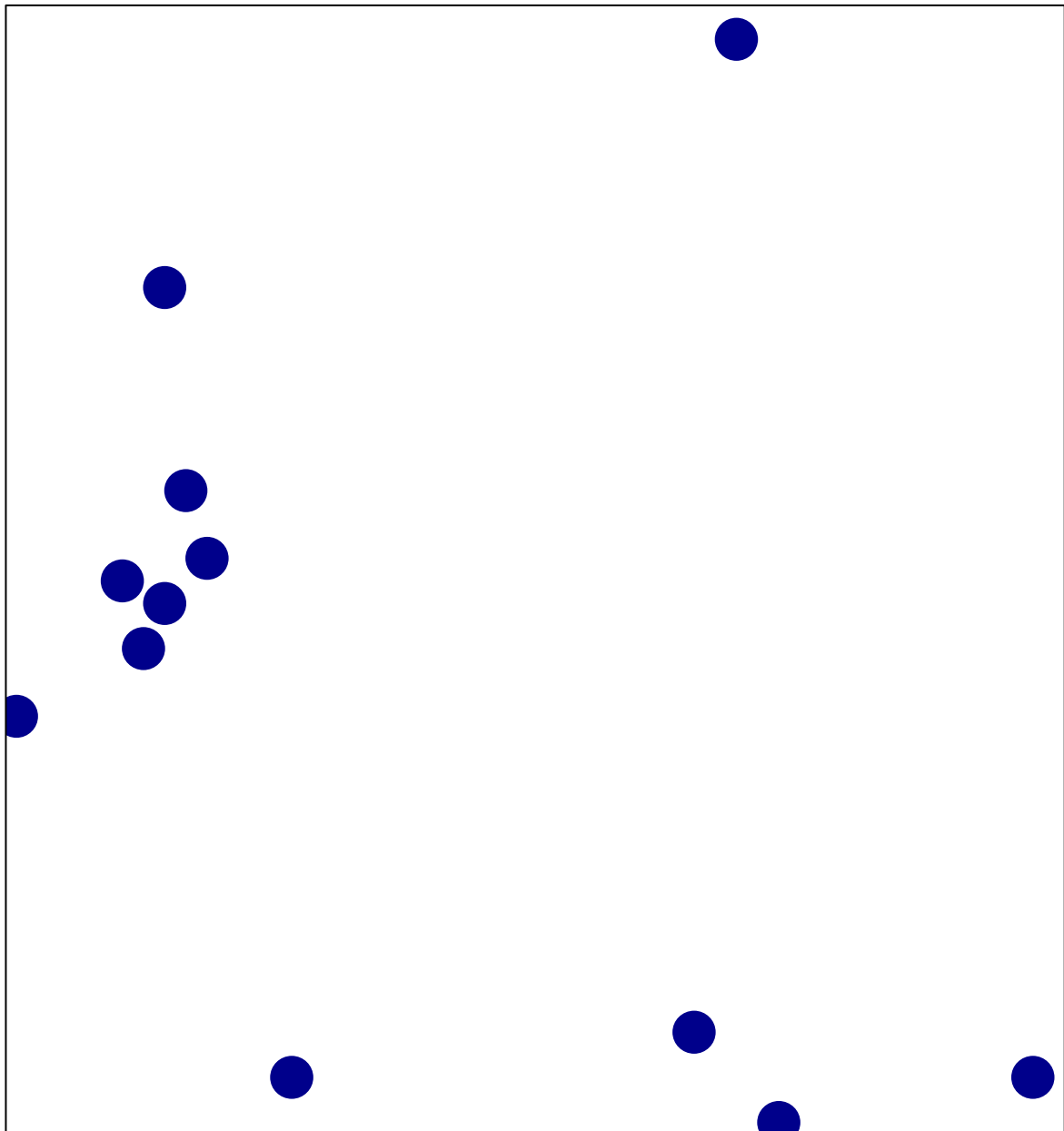
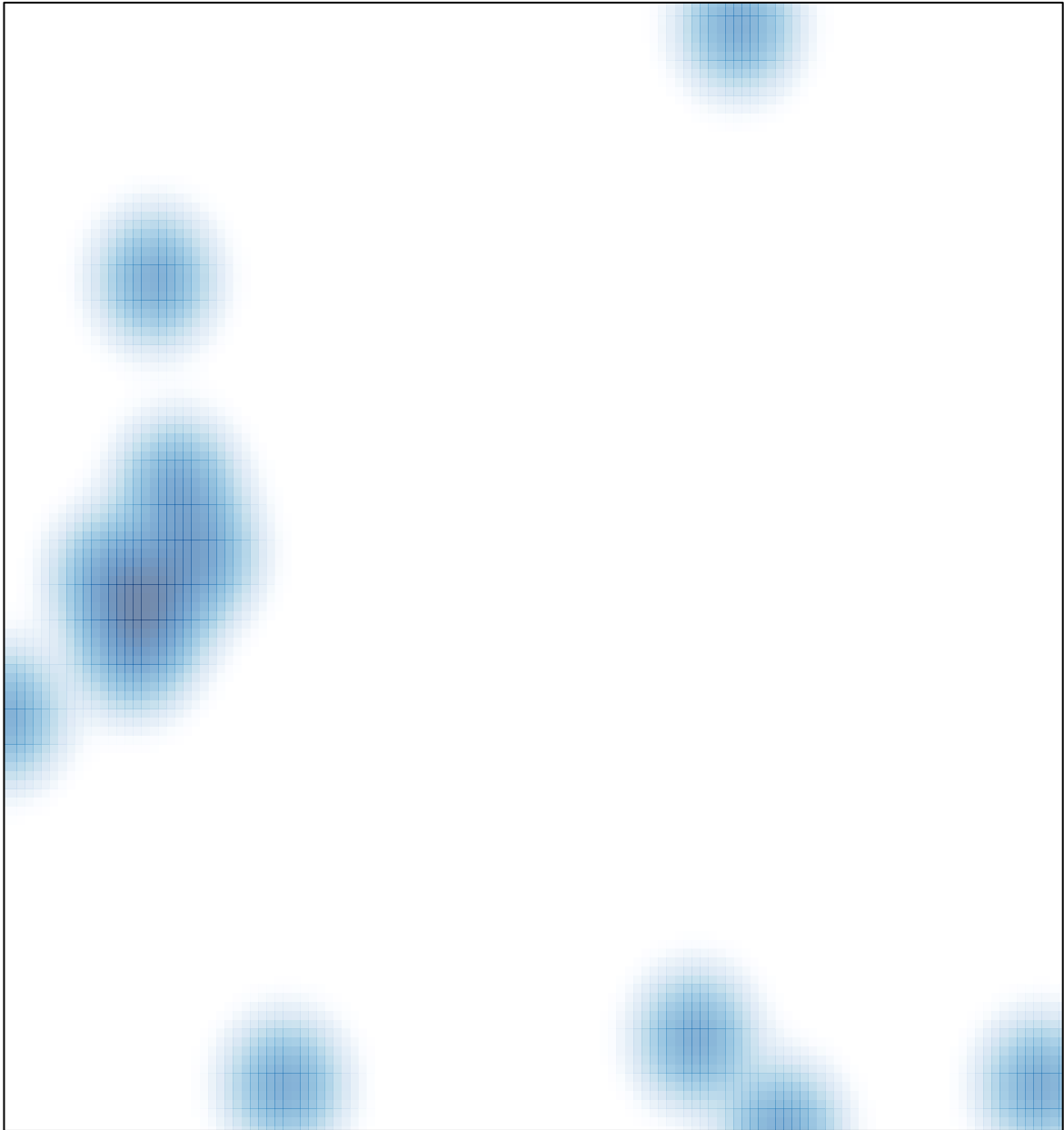


peptidyl-proline modification



features = 12
chi-square p = 0.85

peptidyl-proline modification



features = 12 , max = 1