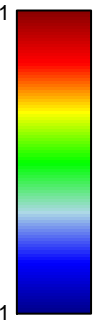
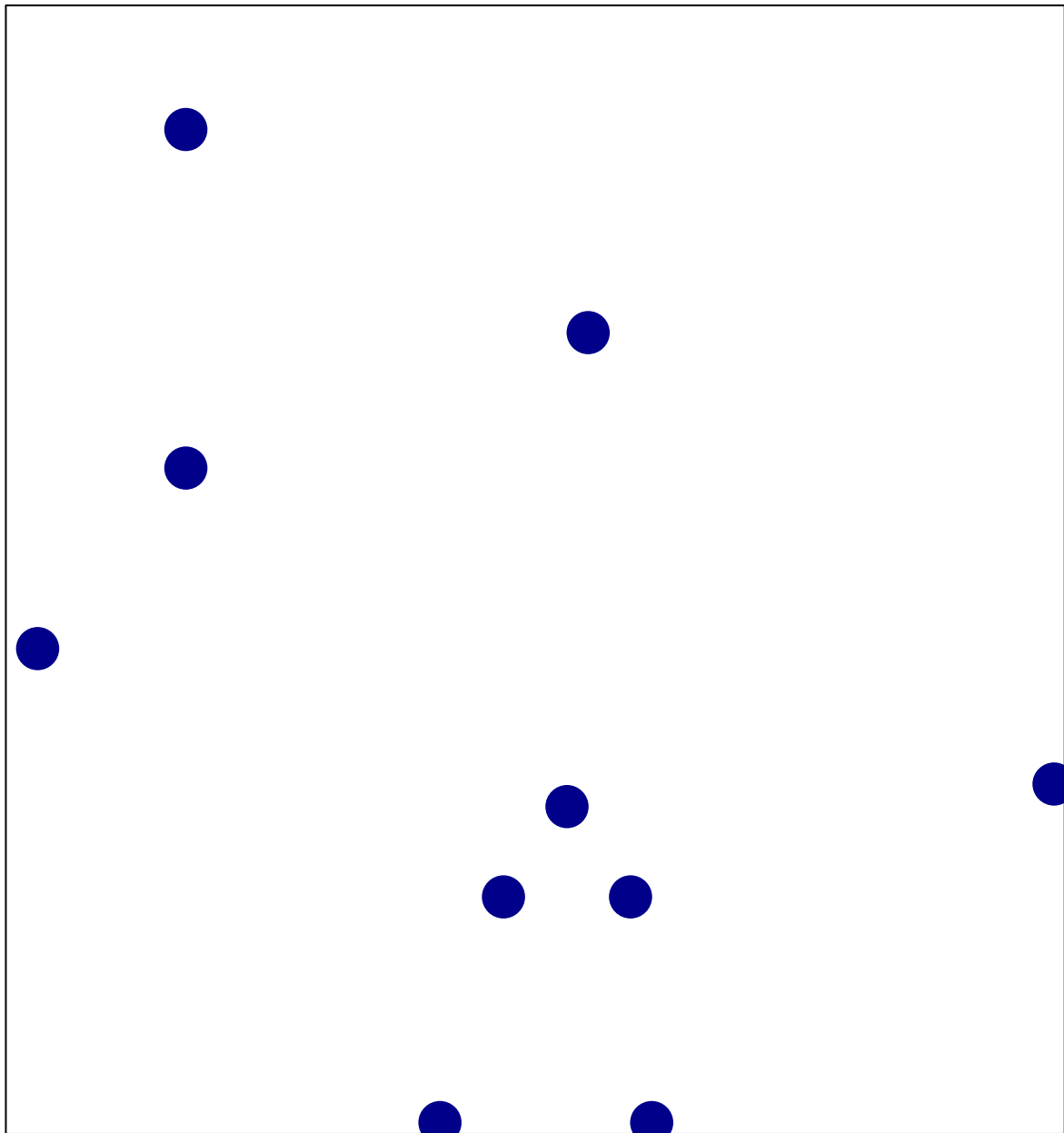
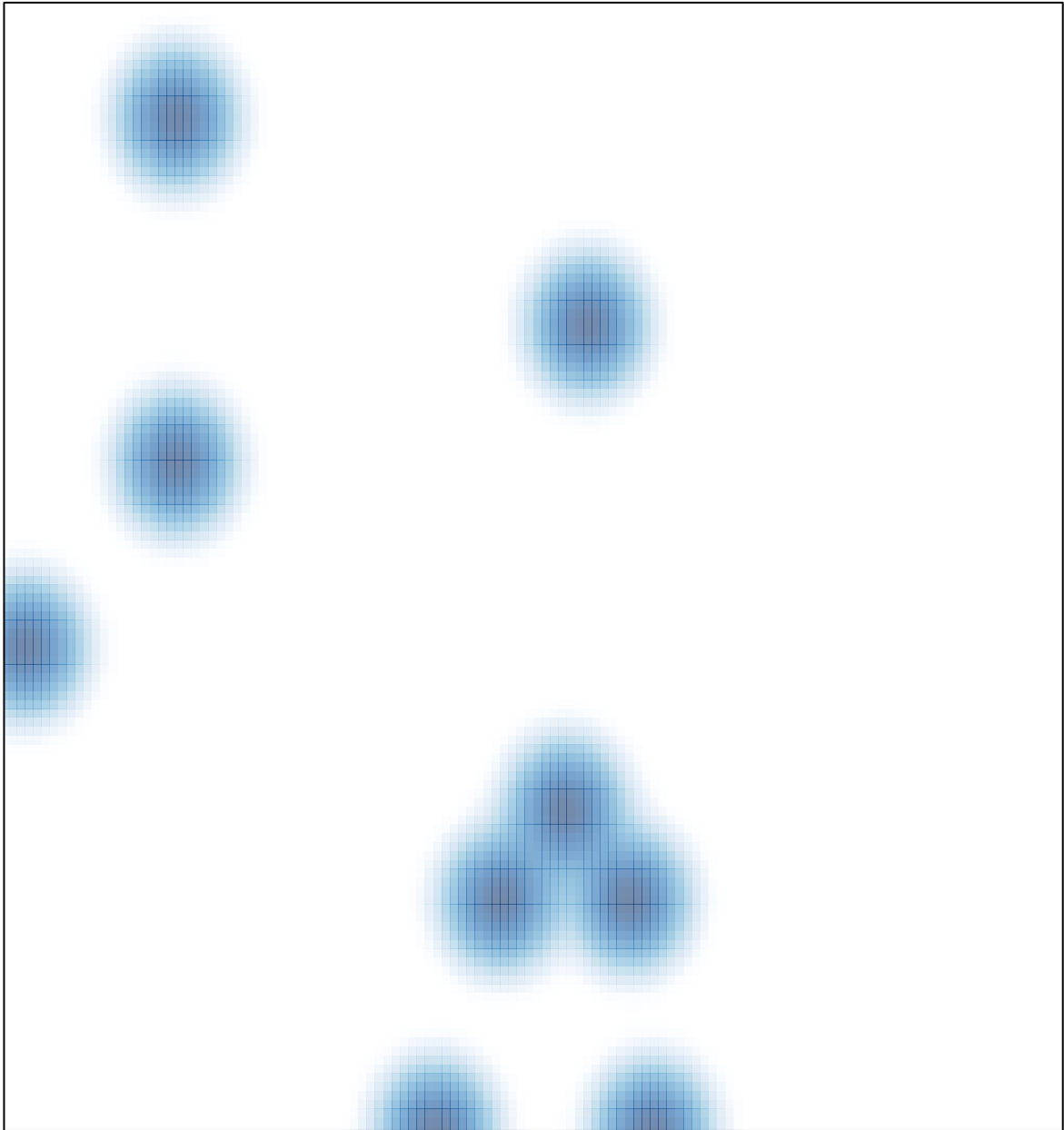


# alternative mRNA splicing, via spliceosome



# features = 10  
chi-square p = 0.83

# alternative mRNA splicing, via spliceosome



# features = 10 , max = 1