regulation of alternative mRNA splicing, via spliceosome
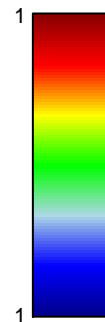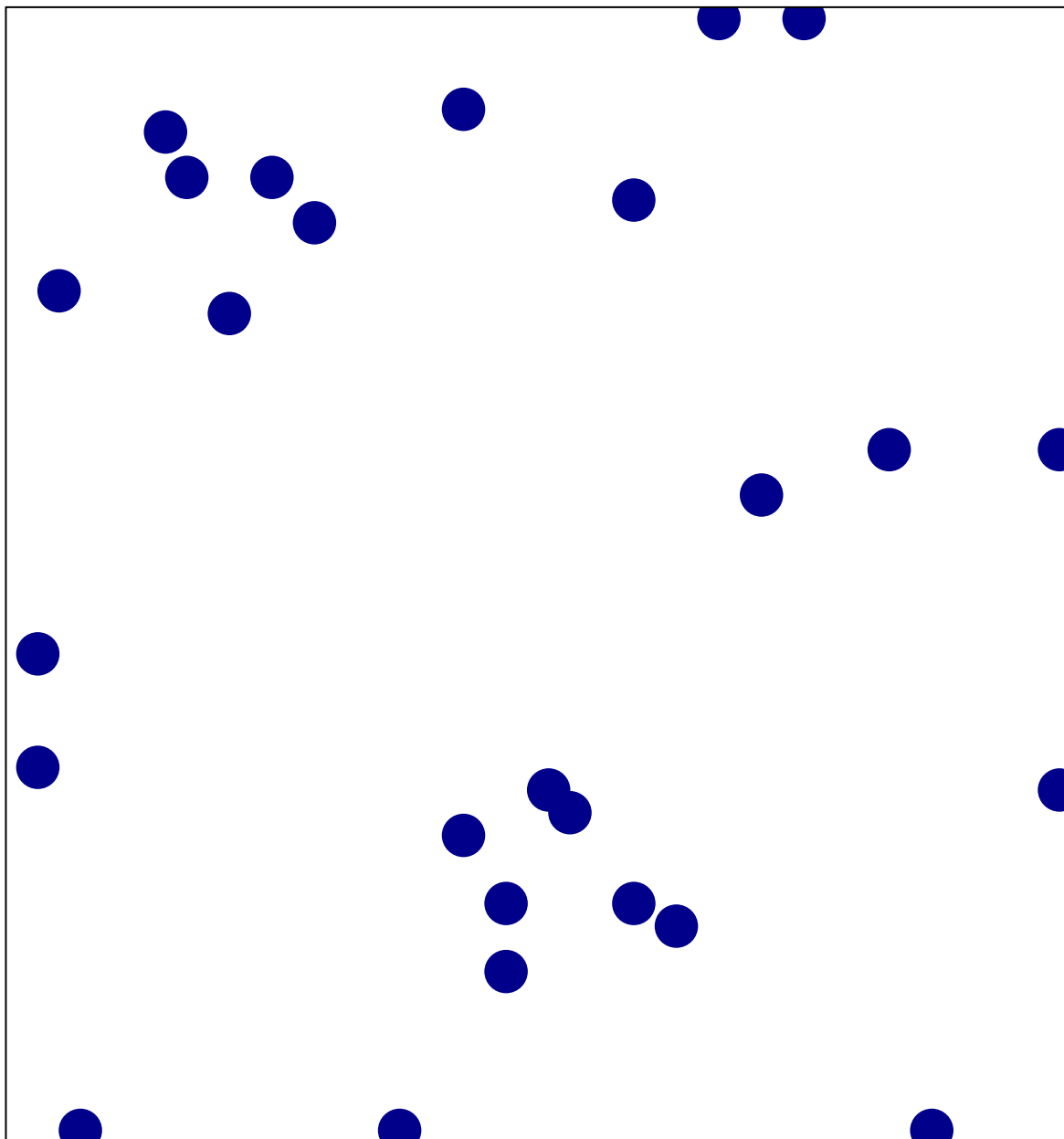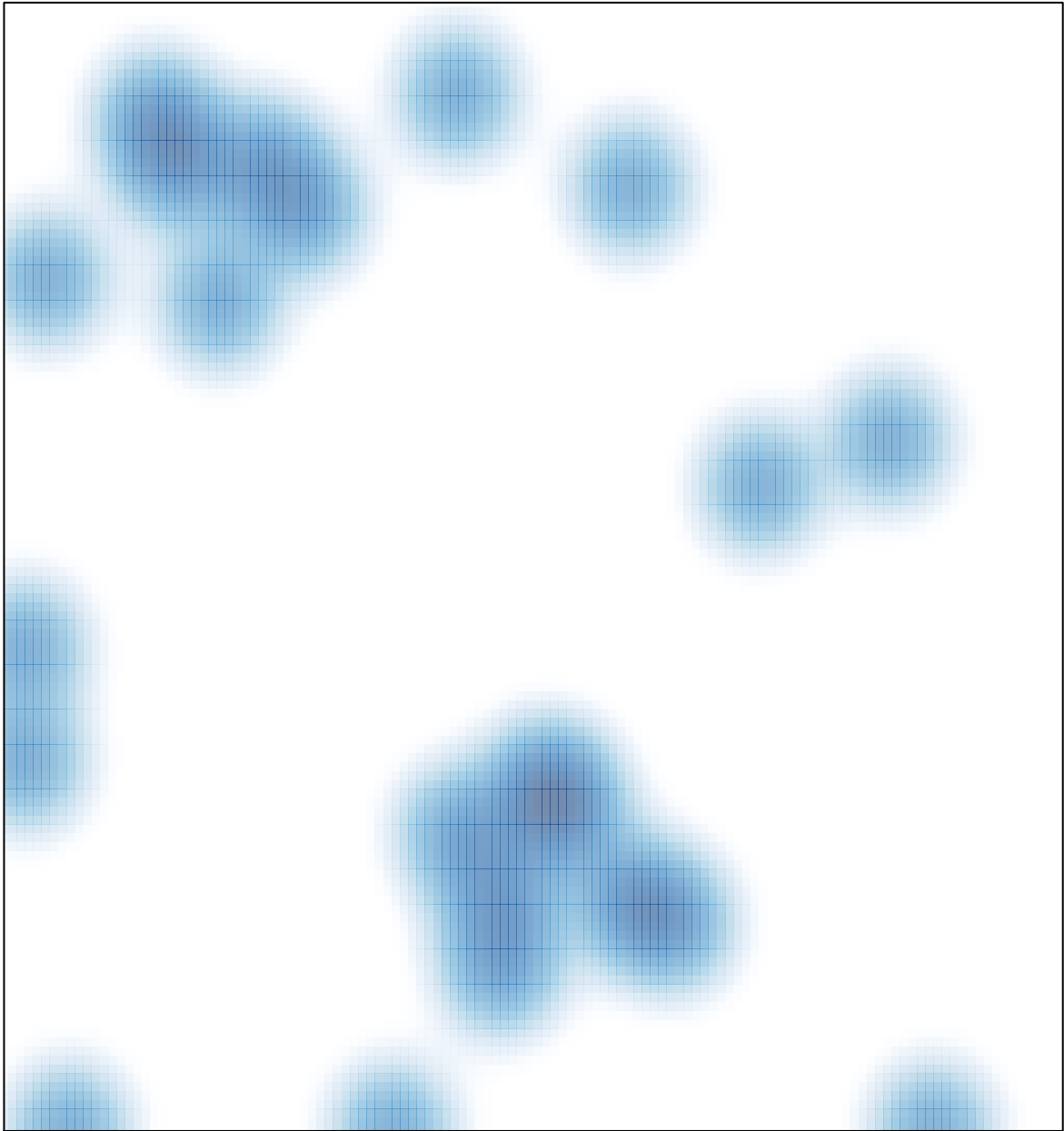

features = 26
chi−square p = 0.84

egulation of alternative mRNA splicing, via spliceoso

features = 26 , max = 1