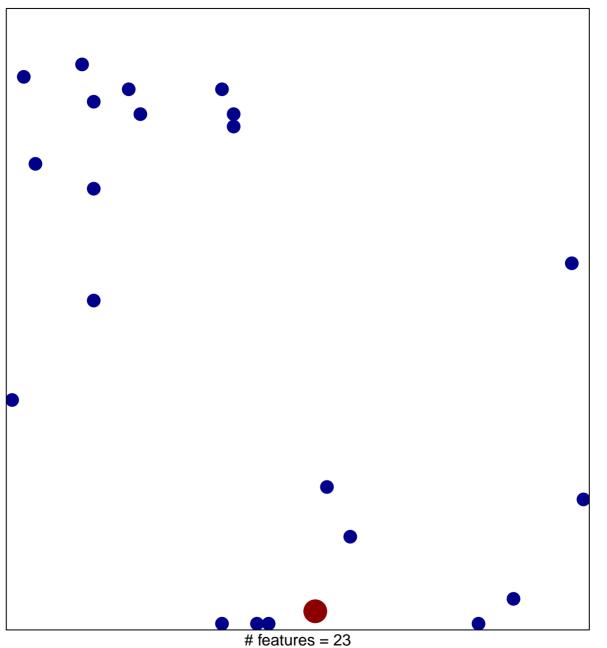
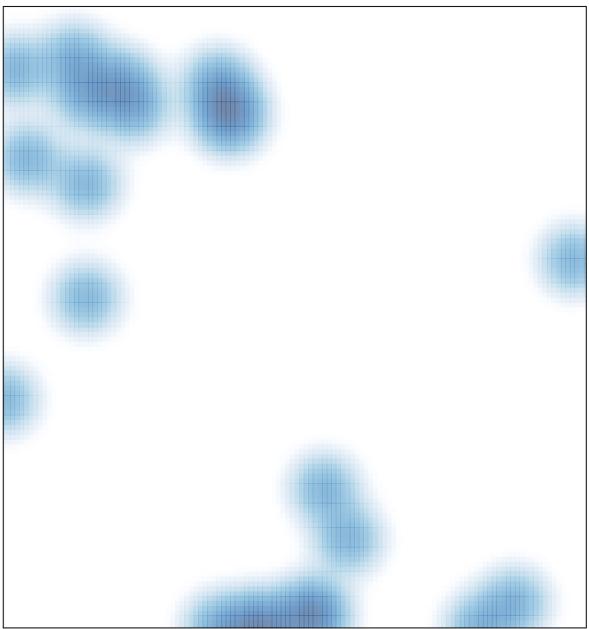
spliceosomal complex assembly



features = 23chi-square p = 0.83

spliceosomal complex assembly



features = 23 , max = 2