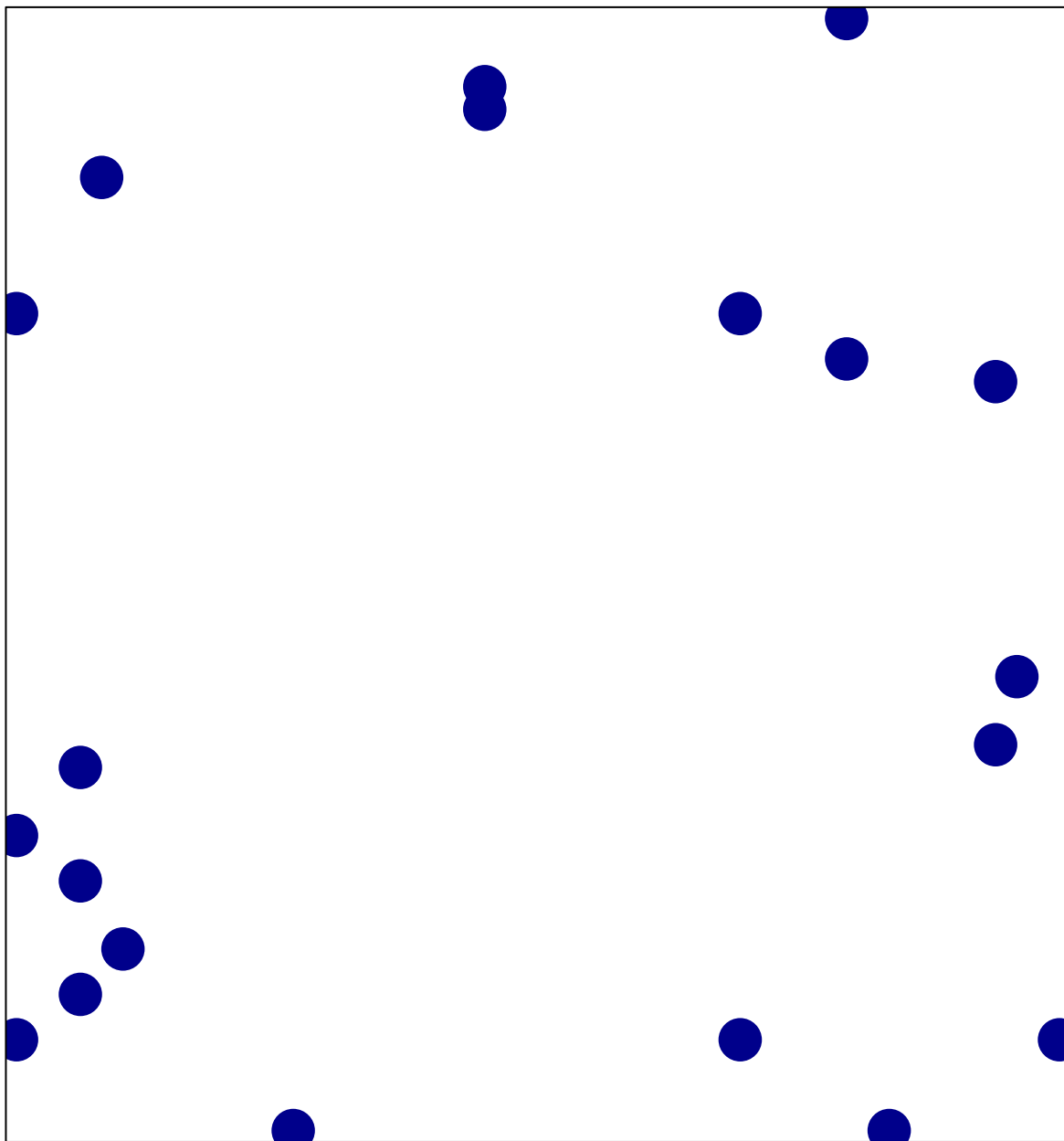
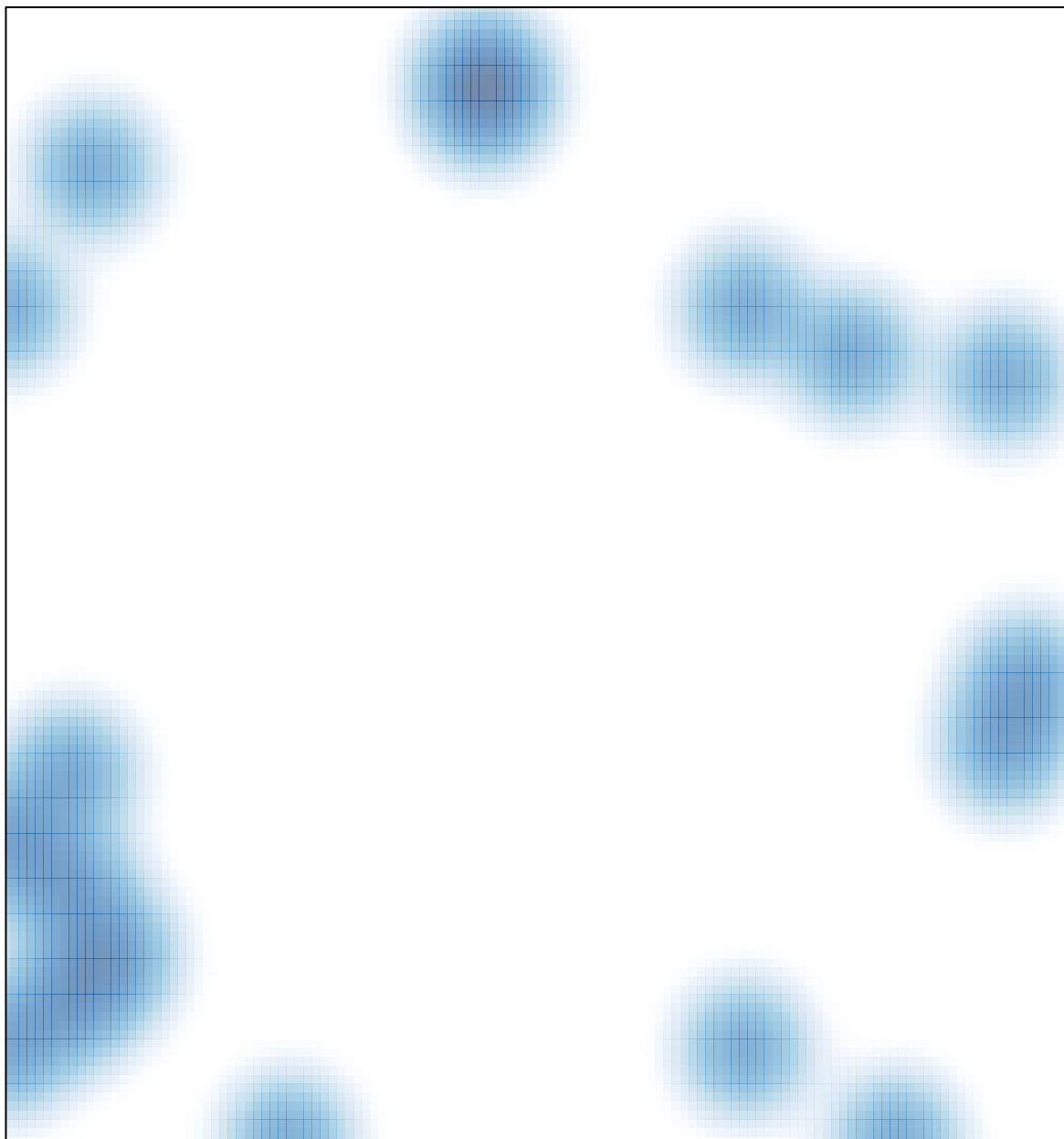


SA_B_CELL_RECEPTOR_COMPLEXES



features = 20
chi-square p = 0.82

SA_B_CELL_RECEPTOR_COMPLEXES



features = 20 , max = 1