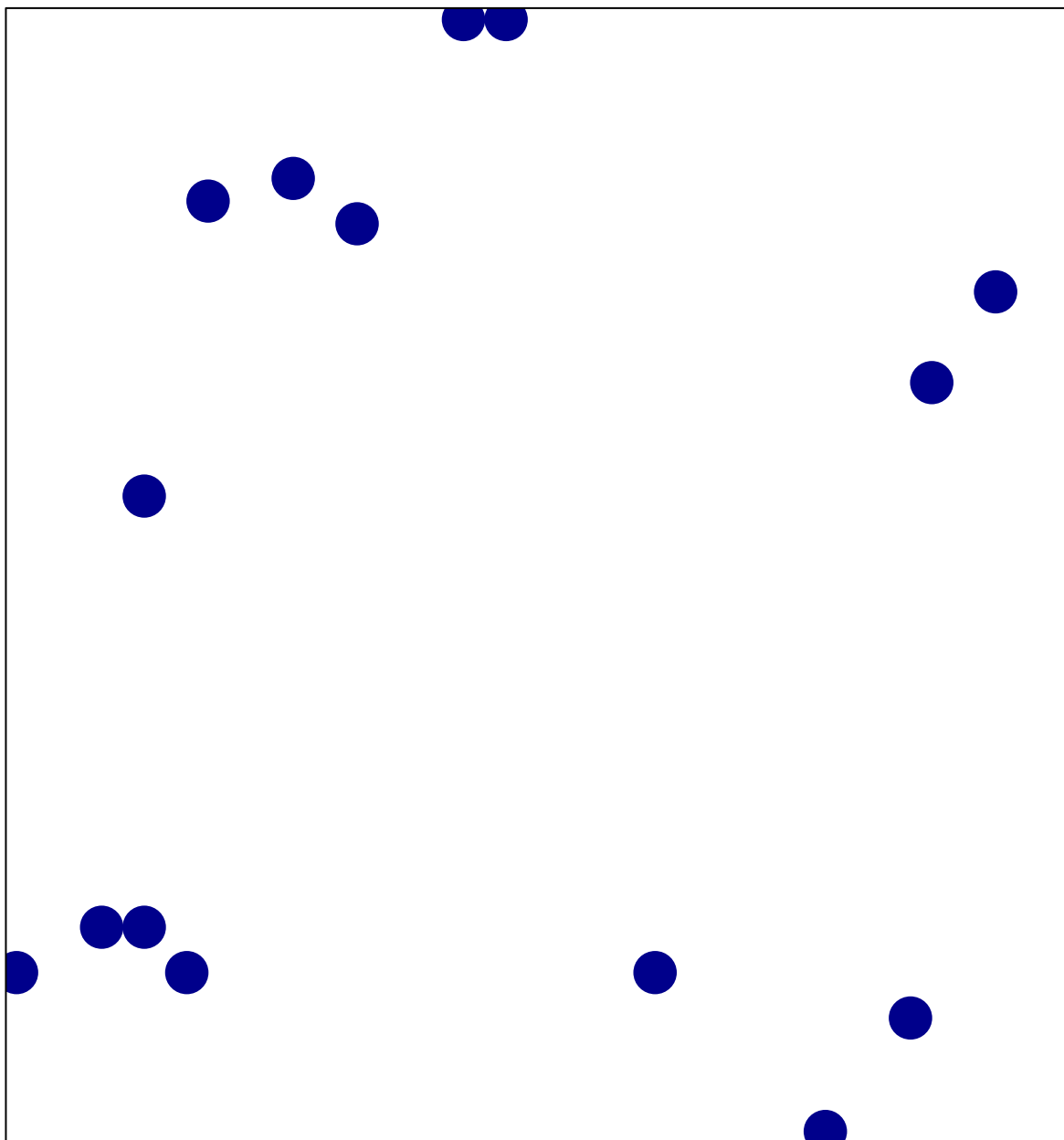
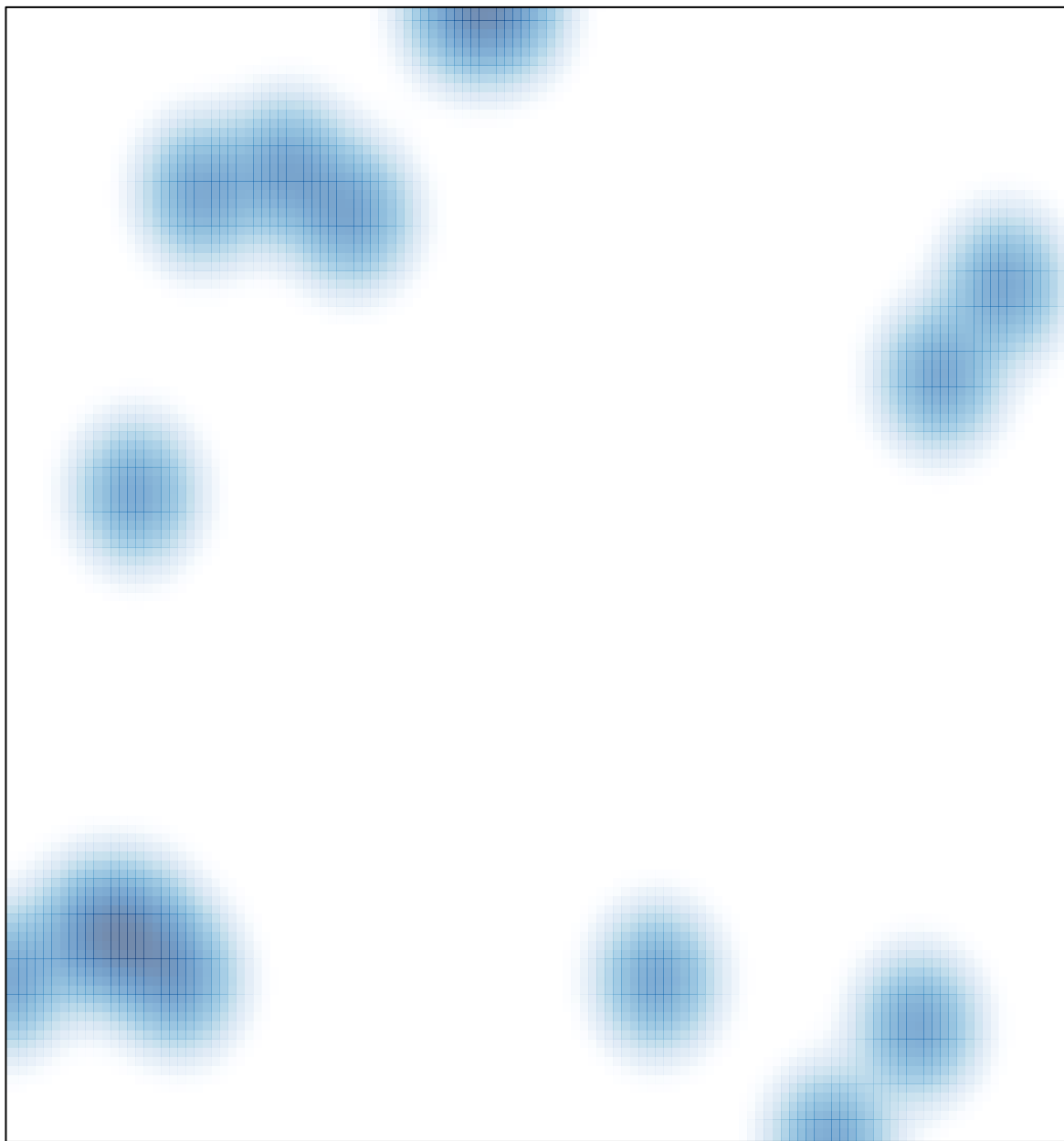


# SA\_B\_CELL\_RECEPTOR\_COMPLEXES



# features = 15  
chi-square p = 0.83

# SA\_B\_CELL\_RECEPTOR\_COMPLEXES



# features = 15 , max = 1