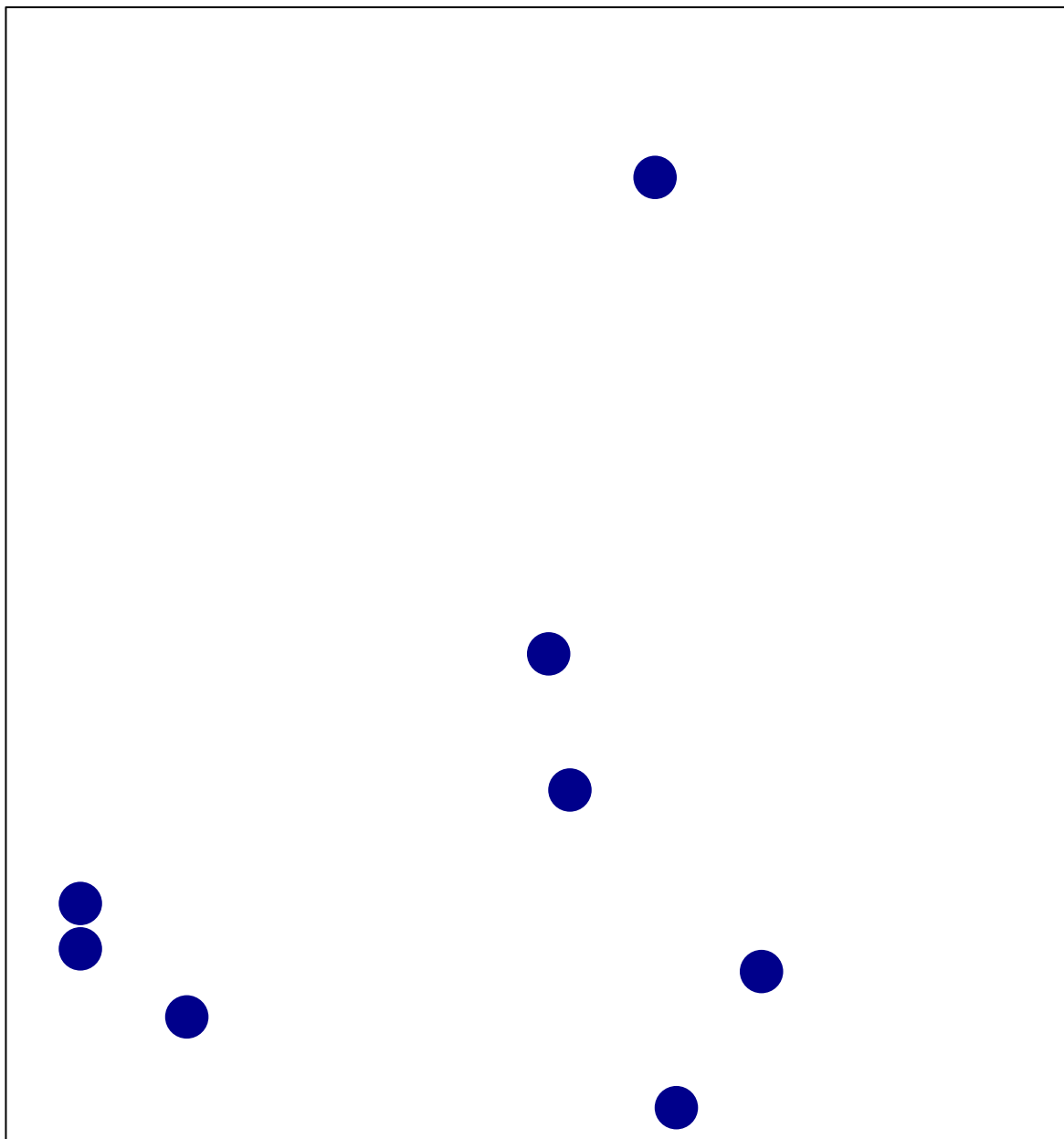


RECEPTOR_LIGAND_BINDING_INITIATES_THE_SECOND_PROTEOLYTIC_CLEAVAGE_OF_N



features = 8
chi-square p = 0.84

BINDING_INITIATES_THE_SECOND_PROTEOLYTIC_



features = 8 , max = 1