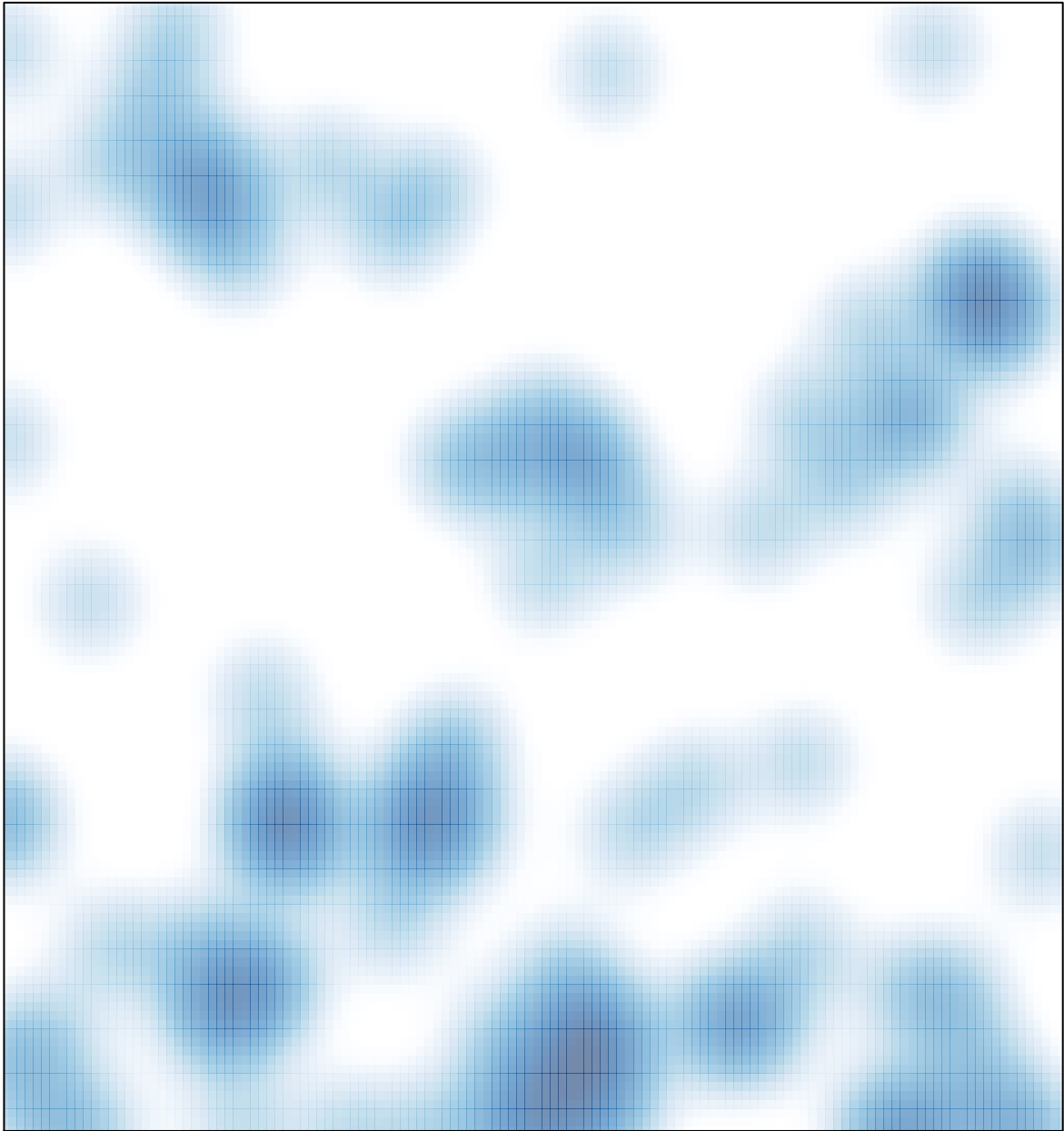


# features = 96  
chi-square p = 0.78

# RECEPTOR\_BINDING\_AND\_DOWNSTREAM\_TRANSM



# features = 96 , max = 3