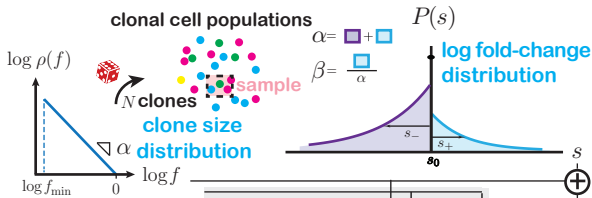


# A repertoire (frequencies):



# B sequenced blood sample (counts):

