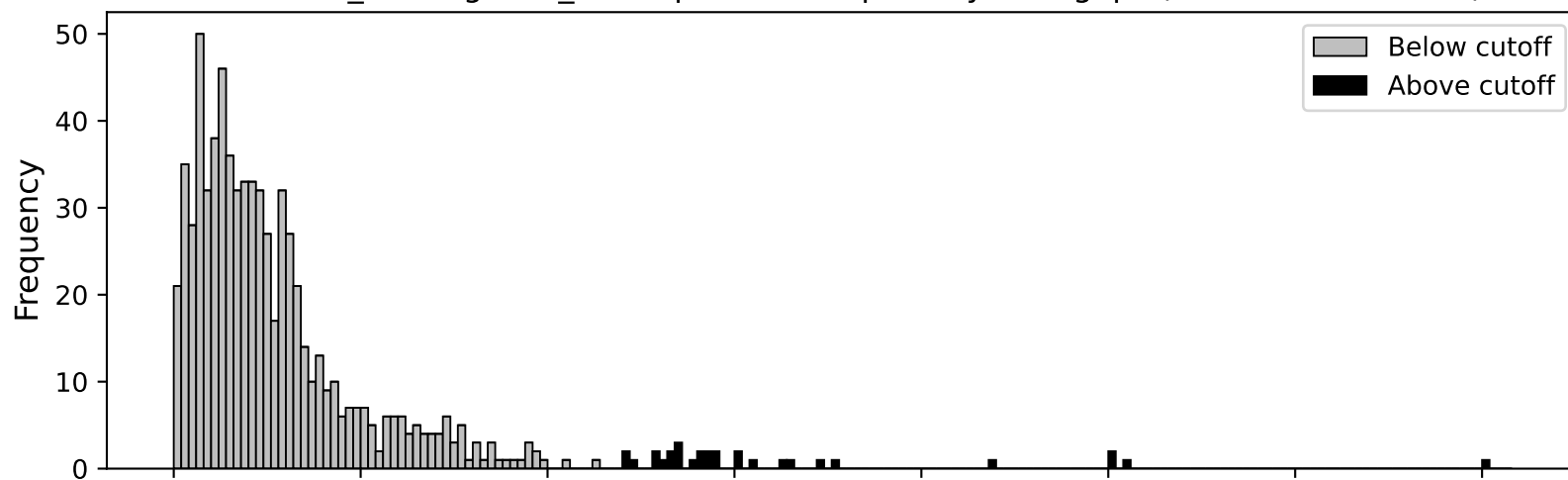


*Neisseria\_meningitidis\_MC58* plus strand primary DUS gaps (total # sites = 734)



*Neisseria\_meningitidis\_MC58* minus strand primary DUS gaps (total # sites = 743)

