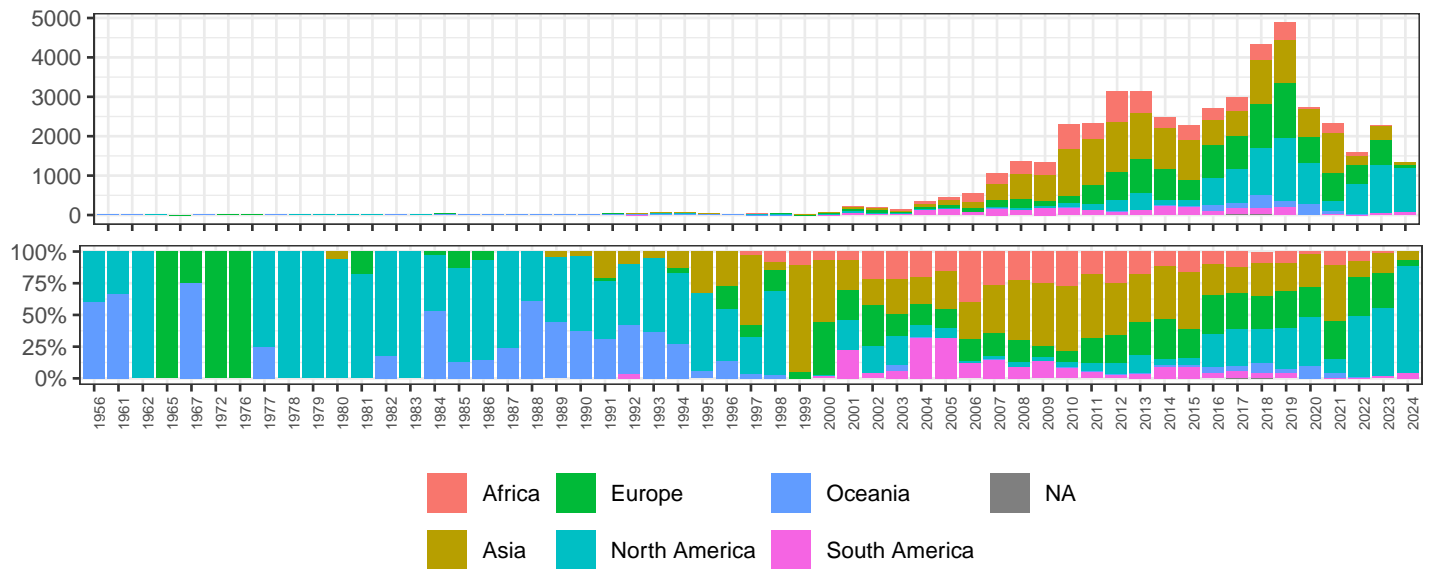
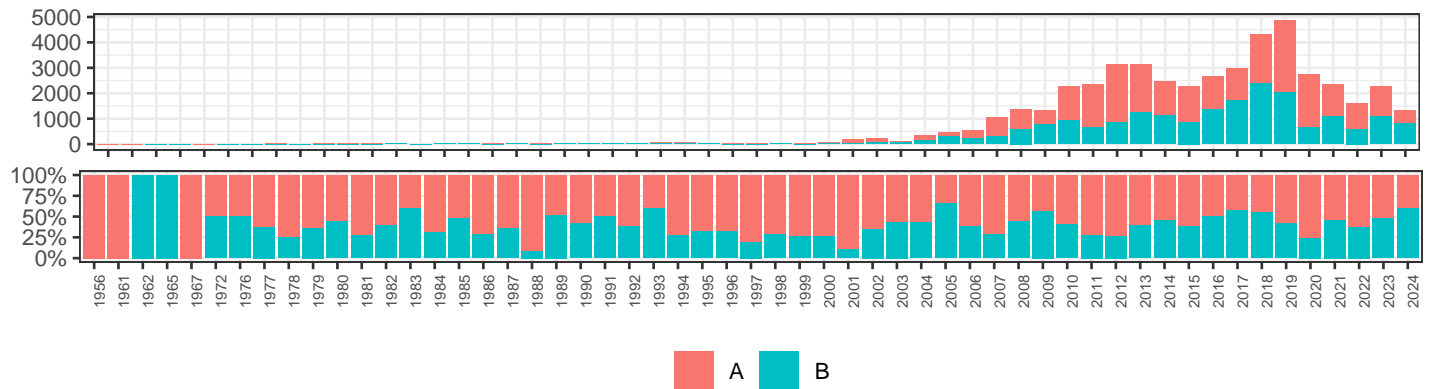


Exploratory Graphics

Across time and geography (n=47190)



RSV Type (n=47190)



Gene coverage (n=47190)

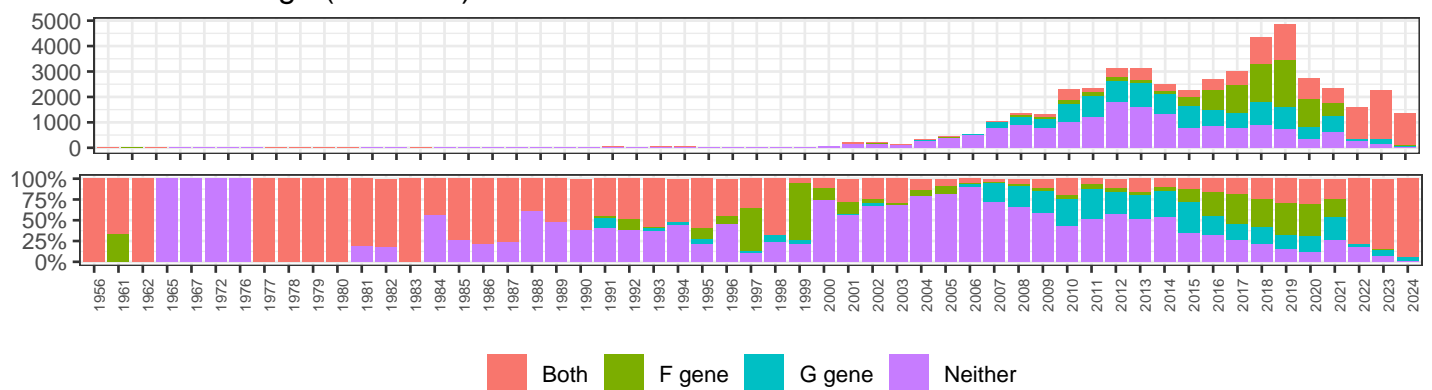


Table 1: Top 25 most frequent sequence submitters with their region and countries

authors	n	regions	countries
wilkins et al.	5265	Oceania, South America, North America, Europe, Africa, Asia	Australia, Brazil, Canada, Finland, Fra
langedijk et al.	1329	Europe, North America, Africa, South America, Asia, Oceania	Netherlands, Canada, Spain, South Afr
jia et al.	1307	Asia, NA	China, NA
song et al.	1237	Asia	China
agoti et al.	1180	Africa	Kenya
kunerth et al.	1171	North America	USA
frazar et al.	1078	North America	USA
yui et al.	1043	NA	NA
otieno et al.	1040	Africa	Kenya
lin et al.	850	North America, Asia, Europe	USA, China, United Kingdom, Netherl
chen et al.	842	Asia	China
ramaekers et al.	837	Europe	Belgium
kamau et al.	759	Africa	Kenya
thongpan et al.	710	Asia	Thailand
kondgen et al.	707	Europe	Germany
prifert et al.	684	Europe	Germany
fourati et al.	676	Europe, NA	France, NA
lauring et al.	654	North America	USA
gimferrer et al.	631	Europe	Spain
oketch et al.	591	Africa	Mali, South Africa, Zambia, Gambia, K
vidoni et al.	529	Europe	France
lee et al.	527	Asia	Taiwan, South Korea
das et al.	523	North America	USA
carr et al.	515	Europe	Ireland
hibino et al.	484	Asia	Japan, Lebanon, Myanmar, Malaysia

Table 2: Top 25 most frequent host

lower_field	n
homo sapiens	47361
NA	4637
pholidota	18

Lineages

Clade (n=47190)

