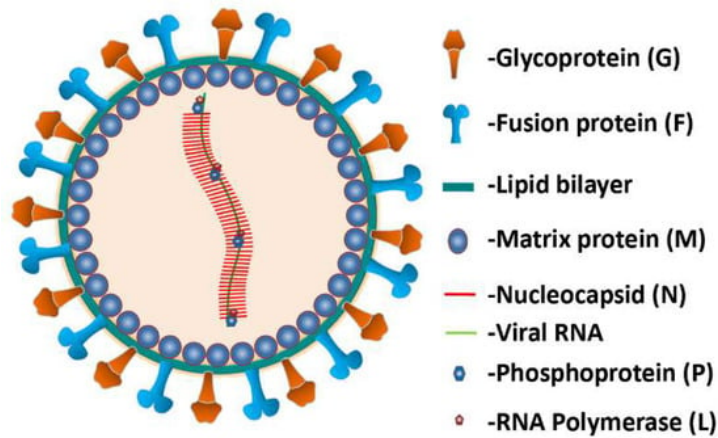


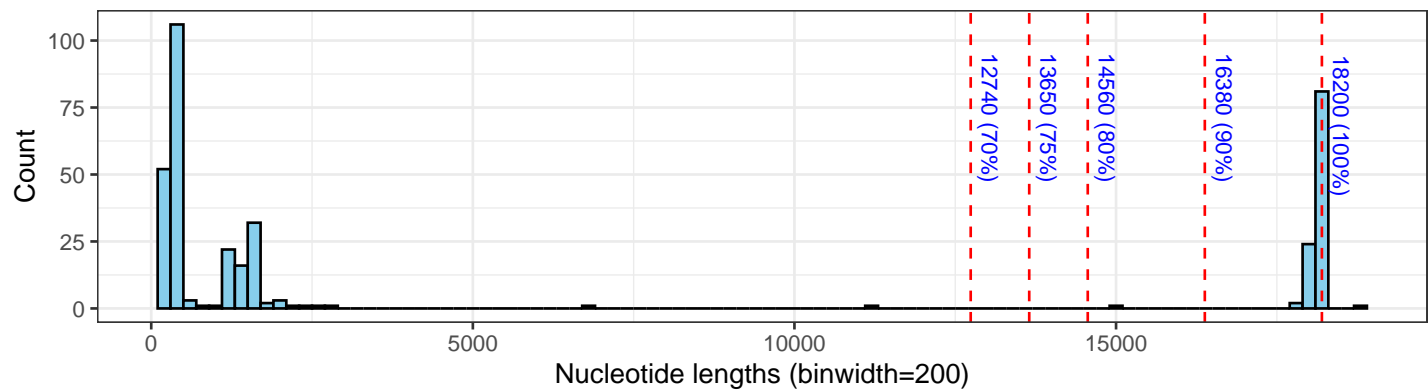
Background

Nipah genome based on "Nipah Virus: Past Outbreaks and Future Containment" by Vinood Soman Pillai, Gayathri Krishna, and Mohanan Valiya Veettil Figure 1:



NCBI GenBank records

Diagnostic plot to estimate min-length filter for phylogenetic analysis (all data = 353)



Frequency and proportion (n=266)

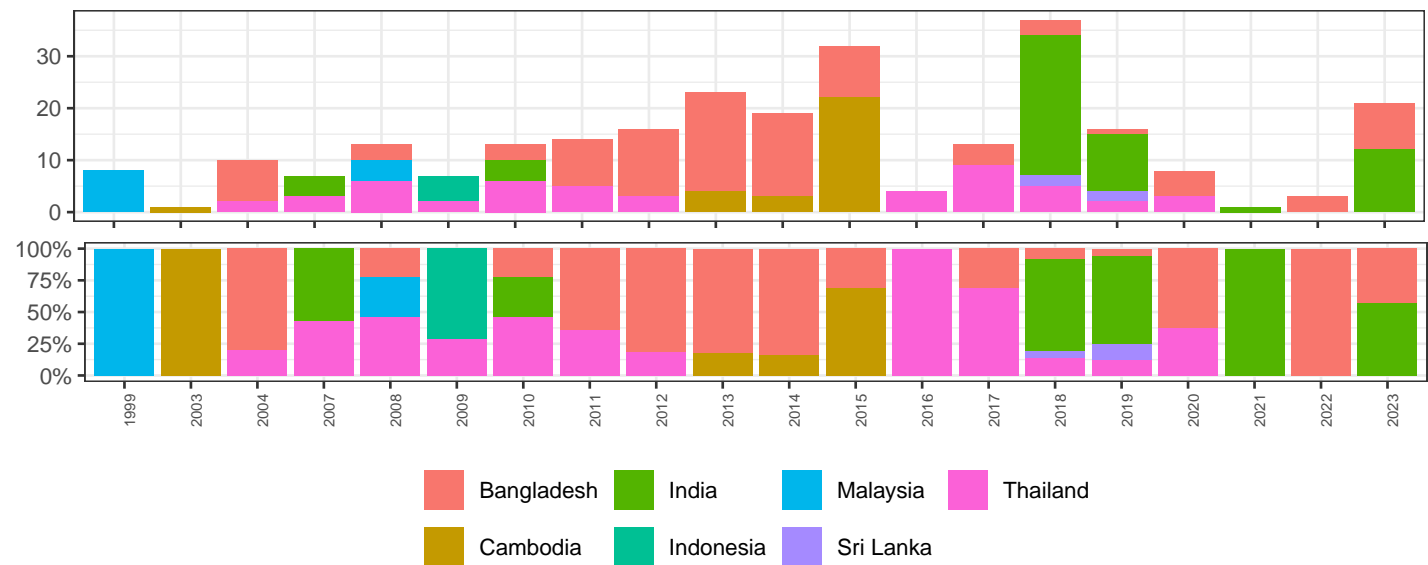


Table 1: Top 25 most frequent sequence submitters with their region and countries

authors	n	regions	countries
wacharapluesadee et al.	97	Asia	Thailand
rahman et al.	52	Asia	Malaysia, Bangladesh
yadav et al.	49	Asia	India
whitmer et al.	34	Asia	Malaysia, Bangladesh
azuero et al.	26	Asia	Cambodia
olival et al.	19	Asia	Bangladesh
anderson et al.	11	Asia	Bangladesh
lo et al.	9	Asia, NA	Bangladesh, NA
hassan et al.	7	Asia	Bangladesh
?	6	Asia	Thailand
arunkumar et al.	6	Asia	India
sendow et al.	5	Asia	Indonesia
abubakar et al.	4	NA, Asia	NA, Malaysia
arankalle et al.	4	Asia	India
kohl et al.	4	Asia	Sri Lanka
duong et al.	3	Asia	Cambodia
harcourt et al.	3	NA, Asia	NA, Bangladesh
sharifah et al.	3	Asia	Malaysia
chan et al.	2	Asia	Malaysia
chua et al.	2	NA, Asia	NA, Malaysia
reynes et al.	2	Asia	Cambodia
bennett et al.	1	NA	Missing
gaudino et al.	1	Asia	Cambodia
honko et al.	1	Asia	Malaysia
johnson et al.	1	Asia	Malaysia