Whole genomic sequencing of the SARS-CoV-2 variants in Sri Lanka

31.12.2021

176 samples were sequenced from 14th to 23nd of December from the community and arrival and departure samples from the airport.

Omicron: 41/176. While most of these samples were from BIA, some were from MOH Minuwangoda, MOH Mahara, CMC and Colombo

Detection of different delta sub lineages and locations:

AY.28 (Sri Lanka delta sub-lineage): 19/176. Divulapitiya, Giriulla, Colombo, Gampaha, Matara, Mirigama,

AY.104 (Sri Lanka delta sub-lineage): Batticaloa, Colombo, Puwakpitiya, Duwulapitiyta, Galle, Gampaha, Ja Ela, Ahungalla, Kuthulgala, Matara, Madampe, Baddegama, Thammitta

AY.106: Negombo

AY.120: Nilpathagama

AY.122: Kandy, Udugampola

AY.127: Galle

AY.1.617.2: Awissawella, Minuwagoda, Diwulapitiya, Dompe, Kandy, Kuruwita, Colombo, Negombo

AY.43: Galle, Chilaw

Timeline of the emergence and displacement of different variants in Sri Lanka

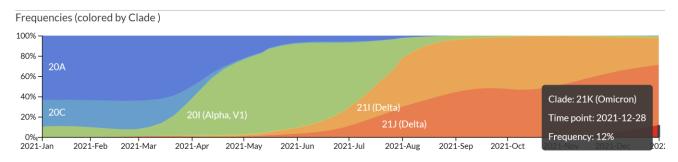


Figure 1: Change in the SARS-CoV-2 variants in Sri Lanka over time. The graph shows changes in the different variants

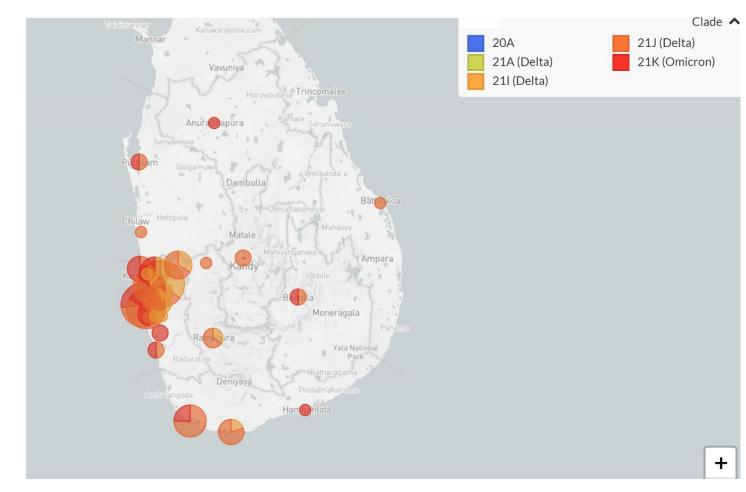
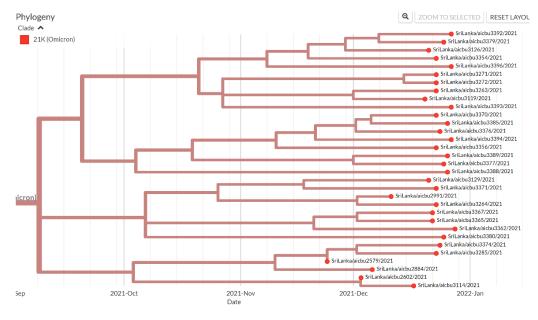


Figure 2: Locations of samples sequenced during the past week (3nd week of December)



Phylogenetic tree of Omicron samples in Sri Lanka

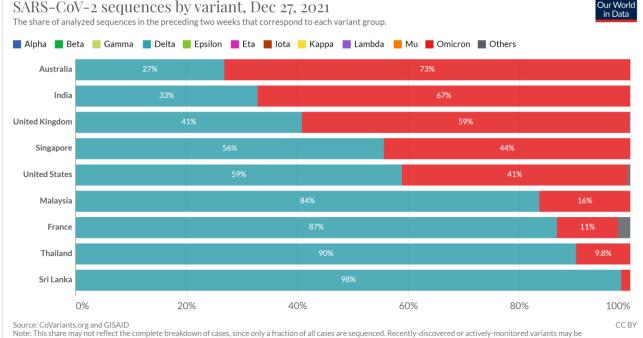


Figure 3: Countries reporting Omicron as of 24th December 2021.

As shown in the map, local transmission of Omicron is reported in many countries. Omicron is currently detected in over 100 countries in the world.

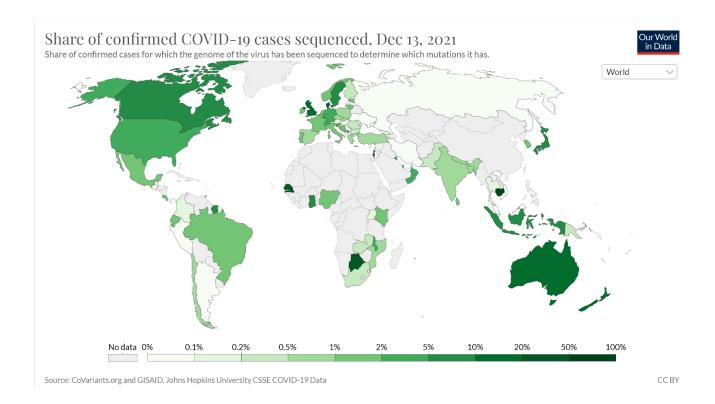
Country / territory	Sequences shared	Sequences shared with complete collection date	Reported COVID-19 cases	Reported deaths from COVID-19	% of cases sequenced and shared
Bangladesh	3,785	3,763	1,583,626	28,061	0.238
India	100,159	91,244	34,799,691	480,290	0.262
Indonesia	10,884	10,839	4,261,879	144,063	0.254
Maldives	818	818	95,032	261	0.861
Myanmar	106	62	529,813	19,257	0.012
Nepal	391	367	827,522	11,586	0.044
Sri Lanka	2,205	2,166	584,107	14,901	0.371
Thailand	10,566	9,573	2,214,712	21,630	0.432
Timor-Leste	356	356	19,833	122	1.795
Bhutan			2,660	3	

Table 1: Number of sequences shared by each country in the SEARO region. (excluding
samples sequenced in this round)



overrepresented, as suspected cases of these variants are likely to be sequenced preferentially or faster than other cases.

Figure 4: Prevalence of Omicron in different countries in the world based on sequencing data (The current round of SL data does not appear yet in this website)



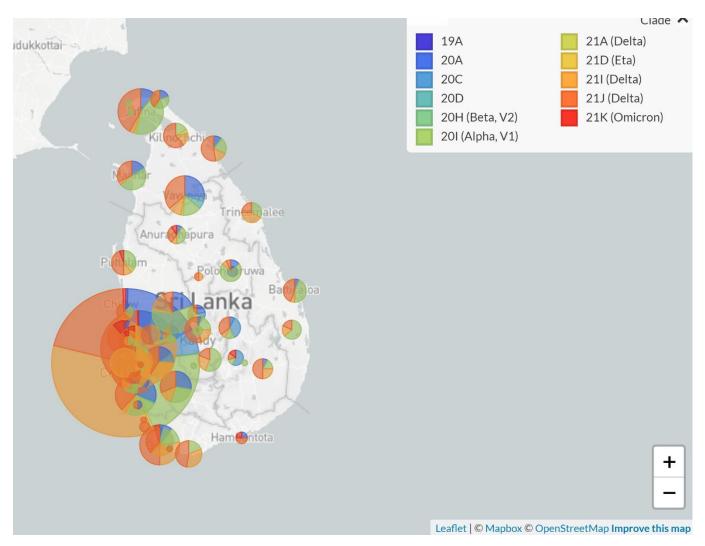


Figure 5: Sequencing of viruses carried out in Sri Lanka so far.

Other variants identified within Sri Lanka: B.1.411: Sri Lankan variant, B.1.1.25, B.1.258, B.1.428, B.4, B.4.7, B.1.1.365, B.1.525, B.1, B.1.1

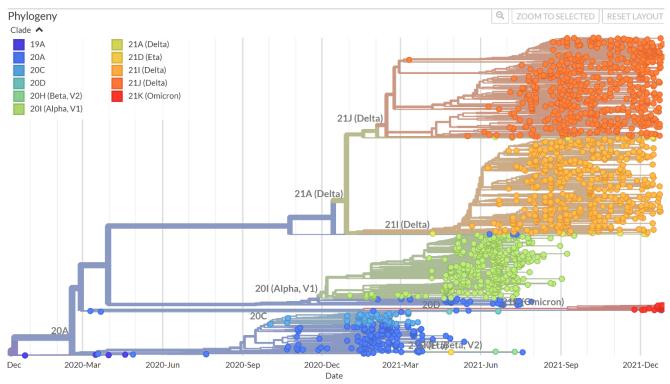


Figure 6: Phylogenetic tree of all SARS-CoV-2 sequences identified in Sri Lanka so far