A Review on SARS-COV-2 – Emerging Variants

Akansha Massey^{1*}, Chaitanya Prakash Lodha²

¹Nursing Tutor, Government College of Nursing, GSVM Medical College, Kanpur, India ²Principal, Regency Institute of Nursing, Kanpur, India

Abstract: This paper presents review on emerging new SARS Co-2 Variants.

Keywords: Delta, Delta Plus, Gamma, Kappa, Lambda, Eta.

1. Introduction

SARS-CoV-2 is a highly transmissible and pathogenic coronavirus that first appeared in late 2019 and has since produced a pandemic of acute respiratory sickness known as "coronavirus disease 2019" (COVID-19), which poses a threat to human health and public safety. Severe respiratory distress syndrome (SARDS) is a severe form of acute respiratory distress. The virus that causes coronavirus disease 2019 (COVID-19), coronavirus 2 (SARS-CoV-2), has several variations, some of which are thought to be of particular concern due to their potential for greater transmissibility, higher virulence, or lower vaccine effectiveness.

Although recombination events between a bat SARS-like coronavirus and a pangolin coronavirus may have resulted in the formation of SARS-CoV-2, mutations have been proven to have an essential role in the ongoing evolution and generation of novel SARS-CoV-2 variants.

Researchers believe the variant first sampled and detected in Wuhan, China, differs from the progenitor genome "by three variants." A rise in transmissibility, an increase in fatality, and a considerable drop in the effectiveness of vaccines, therapy, and other health interventions, according to the World Health

Organization. As a result, many different SARS-CoV-2 lineages have emerged.

Broad Categories of SARS-CoV-2 lineages

- Variant of concern (VOC)
- Variant of interest (VOI)
- Variant under monitoring

A. Variants of Concern

There is substantial evidence that these variations have a considerable impact on transmissibility, severity, and/or immunity, which is likely to affect the epidemiological situation in the EU/EEA. The evidence for these traits, which includes genetic, epidemiological, and in-vitro research, inspires at least moderate confidence.

1) B.1.1.7 (Alpha) is a variation of B.1.1.7.

This variation was first discovered in Southern England in 2020, according to Johns Hopkins data. When compared to patients with non-Alpha strains of the virus, individuals with bouts of the Alpha variety had a higher risk of CCU admission and 28-day mortality, according to a Lancet study. The study also claims that "mortality appeared to be independent of viral strain" for those getting critical care.

2) B.1.351, also known as the Beta variation

South Africa was the first country to introduce the Beta variant. The viral strain may have the capacity to re-infect people who have recovered from previous strains of the virus, according to Johns Hopkins research. Some vaccines may be

Table 1 Variant of concern (VOC)

| WHO label | Country First detected | Year and Month First detected | Evidence on impact of transmissibility | Transmission in EU/EEA | | | |
|-----------|------------------------|-------------------------------|--|------------------------|--|--|--|
| Alpha | United Kingdom | September 2020 | Yes | Community | | | |
| Beta | South Africa | September 2020 | Yes | Community | | | |
| Gamma | Brazil | December 2020 | Yes | Community | | | |
| Delta | India | December 2020 | Yes | Dominant | | | |

Table 2 Variant of interest (VOI)

| WHO label | Country First detected | Year and Month First detected | Evidence on impact of transmissibility | Transmission in EU/EEA |
|-----------|------------------------|-------------------------------|--|------------------------|
| Eta | Nigeria | December 2020 | Yes | Community |
| Theta | The Philippines | January 2021 | Yes | Sporadic/Travel |
| Kappa | India | December 2020 | Yes | Outbreaks |
| lambda | Peru | December 2020 | Yes | Detected |

Table 3

Variant under monitoring

| WHO label | Country First detected | Year and Month First detected | Evidence on impact of transmissibility | Transmission in EU/EEA |
|-----------|------------------------|-------------------------------|--|------------------------|
| Lota | USA | December 2020 | Yes | Community |
| Zeta | Brazil | January 2021 | Yes | Community |

^{*}Corresponding author: akanshamassey17@gmail.com

resistant to this version. According to CIDRAP, the strain has a higher viral load than its predecessor.

B. Variants of Interest

A VOI is a SARS CoV-2 variant with a genetic capacity that influences virus features such as illness severity, immunological escape, transmissibility, and diagnostic escape, according to the WHO. The World Health Organization also found that a VOI results in a significant amount of community transmission. A global surge in instances poses a significant threat to global public health. There is evidence on genomic features, epidemiological evidence, or in-vitro evidence for these variants that could imply a significant impact on transmissibility, severity, and/or immunity, potentially affecting the epidemiological situation in the EU/EEA. However, the evidence is either preliminary or fraught with significant uncertainty.

C. Variants under Monitoring

These additional SARS-CoV-2 variants were discovered as signals thanks to epidemic intelligence, rules-based genomic variant screening, and early scientific data. There is some evidence that they may have qualities comparable to VOCs, however the evidence is either insufficient or has not yet been evaluated by ECDC.

2. About Third Wave

The second wave of coronavirus swept the country, wreaking havoc and killing thousands. While we aren't out of the woods yet, there are several rumors of a third wave of COVID on the way, which may be just as severe and put a strain on healthcare resources.

Many pediatric instances of COVID-19 emerged during the second wave of coronavirus, causing infections in children who were previously thought to be immune. Many people believed that after older folks and younger, healthy people, the following wave would be far more harmful to children due to the nature of the illnesses and the increase in instances.

The United Kingdom was getting set to open up when the first instances of the SARS-CoV-2 Delta strain were discovered in mid-April. COVID-19 cases, hospitalizations, and fatalities were all on the decline, due to months of quarantine and one of the world's most rapid immunization programmes. Two months later, the variation, which was initially discovered in India, has triggered a third wave in the United Kingdom, forcing the government to postpone the complete reopening of society, which had been scheduled for June 21.

Delta, also known as B.1.617.2, is a viral lineage that was initially discovered in India in April and May amid a virulent wave of infections. In other regions of the nation, the lineage spread quickly and exhibited symptoms of partial vaccine resistance. However, researchers found it impossible to separate the variant's inherent characteristics from other variables propelling India's confirmed cases above 400,000 per day, such as huge gatherings.

Although the Delta variety has been connected to a COVID-19 comeback in Nepal, Southeast Asia, and elsewhere, its expansion in the United Kingdom has provided scientists with a clear picture of the harm it presents. Delta appears to be 60 percent more transmissible than the already extremely contagious Alpha strain (also known as B.1.1.7) discovered in late 2020 in the United Kingdom.

Every 11 days, the number of cases of the Delta variety in the United Kingdom doubles. The slower increase in hospital admissions should comfort countries with sufficient vaccination inventories, according to Wenseleers. People who have had one vaccination dose are 75 percent less likely to be hospitalised than those who have not been vaccinated, according to a new Public Health England study1, while those who have been fully protected are 94 percent less likely to be hospitalised. Delta is relatively vaccination resistant, especially in those who have only had a single dose. The third wave of coronavirus infections are largely reliant on vaccination coverage and exposure rates. Having said that, it is impossible to anticipate the severity or risk of an infectious wave. Fresher waves, on the other hand, are weaker than preceding ones, according to scientific study. Even though a third wave of the virus is unavoidable, the magnitude and severity of the third wave, especially with genetic changes fuelling the spread, may be vastly different from the second wave, and could even be well-managed provided all preventative precautions are taken.

3. Conclusion

This paper presented an overview on SARS-COV-2 emerging variants.

References

- SARS Cov-2 Variants/ Lineages, Accessed 30 May 2021, available from https://en.wikipedia.org/wiki/Variants_of_SARS-CoV-2.
- SARS Cov-2 third wave information, Accessed 2 30 May 2021, available from https://timesofindia.indiatimes.com/life-style/health-fitness/healthnews/coronavirus-what-do-we-know-about-the-possible-third-wave-ofcovid-19.
- SARS Cov-2 third wave harmful effects, Accessed 2 10 June 2021, [3] available from https://www.google.co.in/search?q=harmful+effects+of+covid+variants+3rd+wave.
- SARS Cov-2 hits UK, China, India, Accessed 12 June 2021, available from https://www.google.co.in/search?q=sars-cov-2+meaning.
- Covid CDC Guidelines, Accessed 12 June 2021, available from https://www.google.co.in/search?q=cdc+covid+guidelines.
- [6] SARS Cov-2 Delta variant, Accessed 15 June 2021, available from https://www.google.co.in/search?q=delta+plus+variant