Genetic diversity of the novel coronavirus SARS-CoV-2 (COVID-19) in Portugal

More information at https://insaflu.insa.pt/covid19



Situation Report January 21th, 2025

The National Institute of Health Doutor Ricardo Jorge, I.P. (INSA) has analysed 50568 SARS-CoV-2 genome sequences so far.

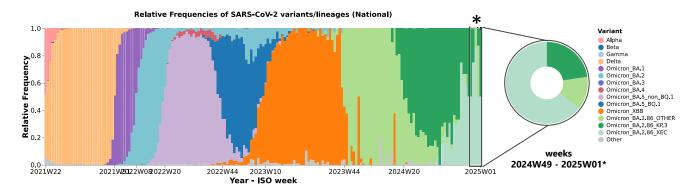
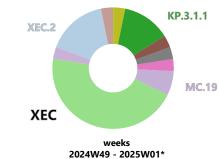


Figure 1: Evolution of the weekly relative frequency of the SARS-CoV-2 variants circulating in Portugal between ISO weeks 22/2021 (31/05/21 - 06/06/21) and 01/2025 (30/12/24 - 05/01/25), with emphasis on the latest weeks. *The presented relative frequencies refer to the period of ISO weeks 49/2024 to 01/2025. This and other graphs can be explored interactively on the website.

Main highlights

- The lineage BA.2.86 of the Omicron variant has been dominant in Portugal since week 44 of 2023, following its first detection in week 33/2023. Among its lineages, KP.3 stands out, as it included in the list of variants of interest by the ECDC (https://www.ecdc.europa.eu/en/covid-19/variants-concern). The relative frequency of KP.3 (and its sublineages) has been showing a declining trend in Portugal, accounting for 22.6% of the sequences analyzed between weeks 49/2024 and 01/2025 (Figure 1).
- The **recombinant XEC lineage** of the *Omicron* variant, which resulted from the recombination between two BA.2.86 sublineages (KS.1.1 and KP.3.3) was recently added to the **ECDC's variants under monitoring** list. Therefore, it is presented here independently from its ancestral lineage for better monitoring and interpretation of this report. **XEC** was first detected in Portugal in week 31/2024 (**Figure 2**), and in the latest sampling (weeks **49/2024** to **01/2025**), it accounted for **64.5%** of the sequences analyzed, remaining **the dominant lineage in Portugal**. This lineage has also been detected in several countries, with an **increasing global trend**.



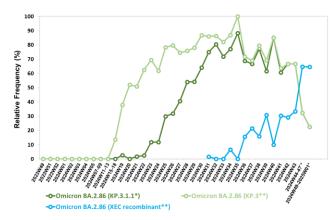


Figure 2: Evolution of the relative frequency of some sub-lineages of interest circulating in Portugal. The circular graph shows the distribution of the relative frequencies of SARS-CoV-2 sub-lineages in the period of ISO weeks 49/2024 and 01/2025 (02/12/24 – 05/01/25) **highlighting the most frequent sub-lineages in this period.** The evolution of relative frequencies of KP.3 and KP.3.1.1 lineages and the recombinant lineage XEC during the last weeks is shown in the line plot. *The presented relative frequencies correspond to the sub-lineages and their descendants. **Other graphs can be explored interactively on the website.**

Autorship

Genomics and Bioinformatics Unit Department of Infectious Diseases NATIONAL INSTITUTE OF HEALTH DOUTOR RICARDO JORGE Avenida Padre Cruz, 1649-016 Lisboa, PORTUGAL

Recommended citation

National Institute of Health (INSA) Doutor Ricardo Jorge. Genetic diversity of the novel coronavirus SARS-CoV-2 (COVID-19) in Portugal. Lisbon, Portugal INSA; 2022. Available at: https://insaflu.insa.pt/covid19

Useful links

https://www.ecdc.europa.eu/en/covid-19/situation-updates/variants-dashboa https://www.who.int/activities/tracking-SARS-CoV-2-variants https://cov-lineages.org/lineage_list.html https://cutreak.info/





