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

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



Situation Report February 12th, 2025

The National Institute of Health Doutor Ricardo Jorge, I.P. (INSA) has analysed 50590 SARS-CoV-2 genome sequences so far.

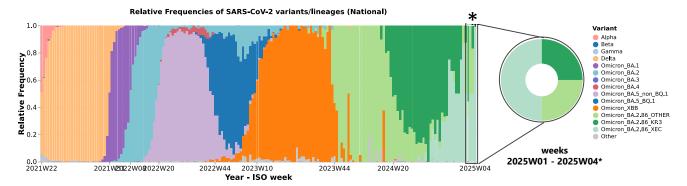
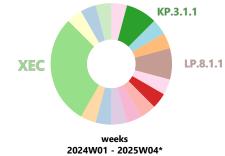


Figure 1: Evolution of the weekly relative frequency of the SARS-CoV-2 variants circulating in Portugal between ISO weeks 22/2021 (31/05/21 - 06/06/21) and 04/2025 (20/01/25 - 26/01/25), with emphasis on the latest weeks. *The presented relative frequencies refer to the period of ISO weeks 01/2025 to 04/2025. 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 **5.5%** of the sequences analyzed between weeks **01/2025** and **04/2025** (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 01/2025 to 04/2025), it accounted for 66.6% of the sequences analyzed, remaining the dominant lineage in Portugal. This lineage has also been detected in several countries, with an increasing global trend.



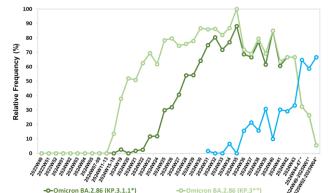


Figure 2: Evolution of the relative frequency of some sub-lineages of interest circulating in Portugal. The circular graph shows the distribution of the relative frequencies of SARS-CoV-2 sub-lineages in the period of ISO weeks 01/2025 and 04/2025 (30/12/24 – 26/01/25) highlighting the most frequent 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

Omicron BA.2.86 (XEC recombinant**)

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





