



Genomic surveillance report

Update for Belgium, 20/09/2022

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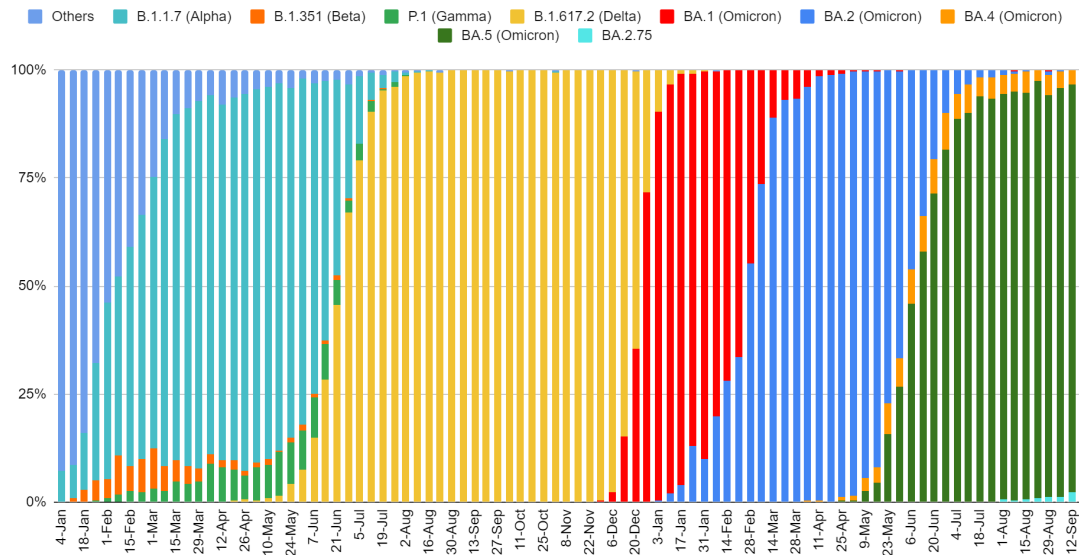
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Previous reports are available online using this [link](#).

Executive summary

Omicron BA.5 remains the dominant lineage in Belgium and accounts for around 94% of the most recent infections. The genomic situation is currently stable.



To date, BA.2.75 samples account for approximately 1% of the circulating strains.

1 Epidemiological context and indicators related to diagnostic activities

Omicron BA.2 and BA.2.75 can be distinguished from BA.4 and BA.5 as the latter variants present a deletion 69/70 in the Spike gene (S gene target failure, SGTF).

At this stage, SGTF samples (BA.5 and BA.4) still represent the vast majority of the most recent infections, but a slow decline in favor of non-SGTF samples (including BA.2.75) may have started since a couple of weeks (Figure 1). These recent changes must be interpreted with caution as there is currently a low number of samples analyzed by the federal PCR platform (figure 2). This signal will nevertheless be followed-up carefully in the coming weeks. Note that the numbers have been adapted compared to the previous weeks, as it has been noticed that some QC results had been included.

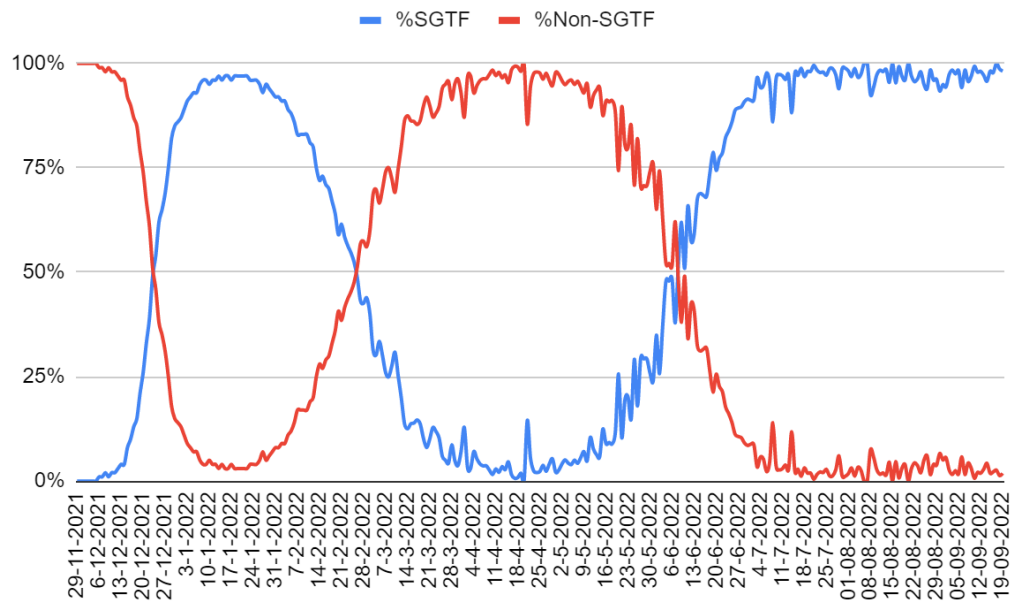


Figure 1: S gene target failure (SGTF, in blue: BA.4 and BA.5) and others (red: currently considered predominantly BA.2 and BA.2.75) among positive samples reported by the federal platform laboratories.

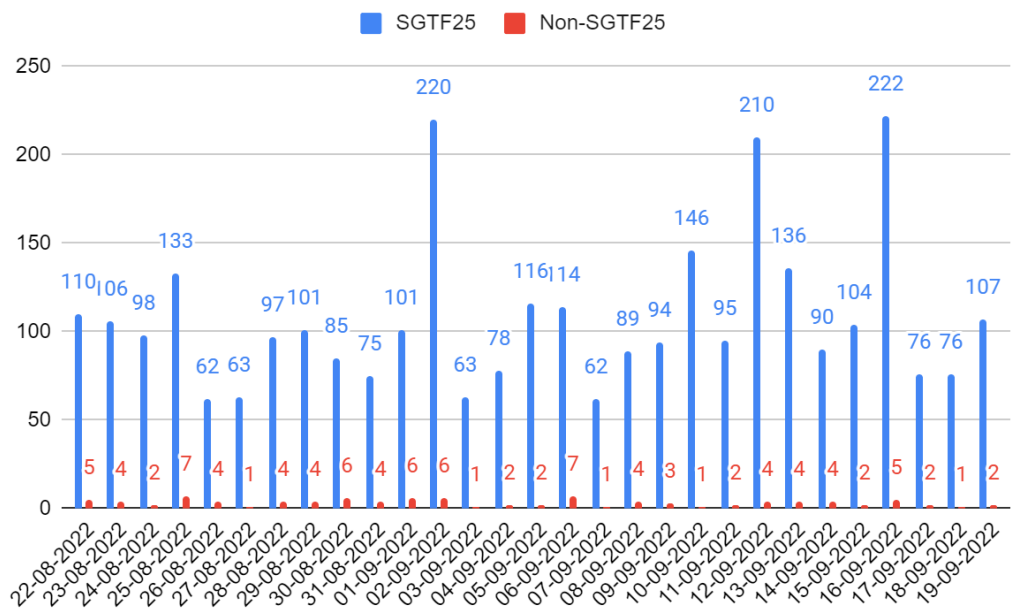


Figure 2: S gene target failure (SGTF, in blue: BA.4 and BA.5) and others (red: currently considered predominantly BA.2 and BA.2.75) among positive samples reported by the federal platform laboratories.

We have been informed that the government will stop the activity of the federal platform PCR laboratories in the coming weeks. This implies that in the future, figure 1 and figure 2 will not be updated and genomic surveillance in Belgium will solely rely on sequencing efforts.

2 Monitoring of Variants of Concern in Belgium

During the last two weeks of baseline surveillance - 05/09/2022 to 18/09/2022 - (533 sequences collected at this stage), BA.5 represented 94%, BA.4 represented 4% and BA.2.75 represented 1% of the samples sequenced. The BA.2.75 subvariant is currently classified as a variant of interest (VOI) by the ECDC, and seems to have a (modest) transmission advantage over BA.5.

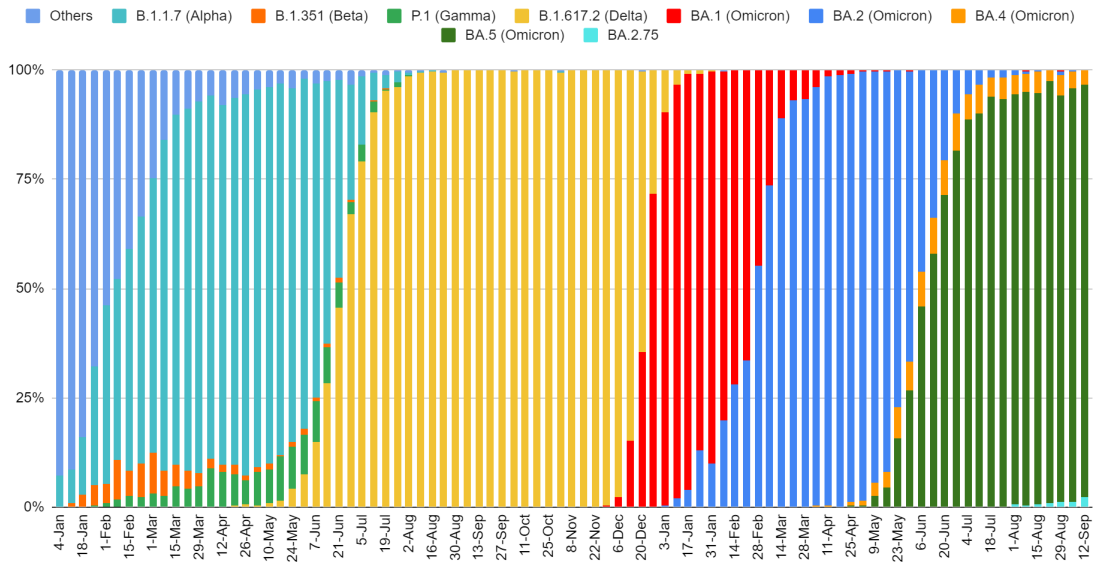


Figure 3: Share of variants of concern per week in Belgium since January 2021.