

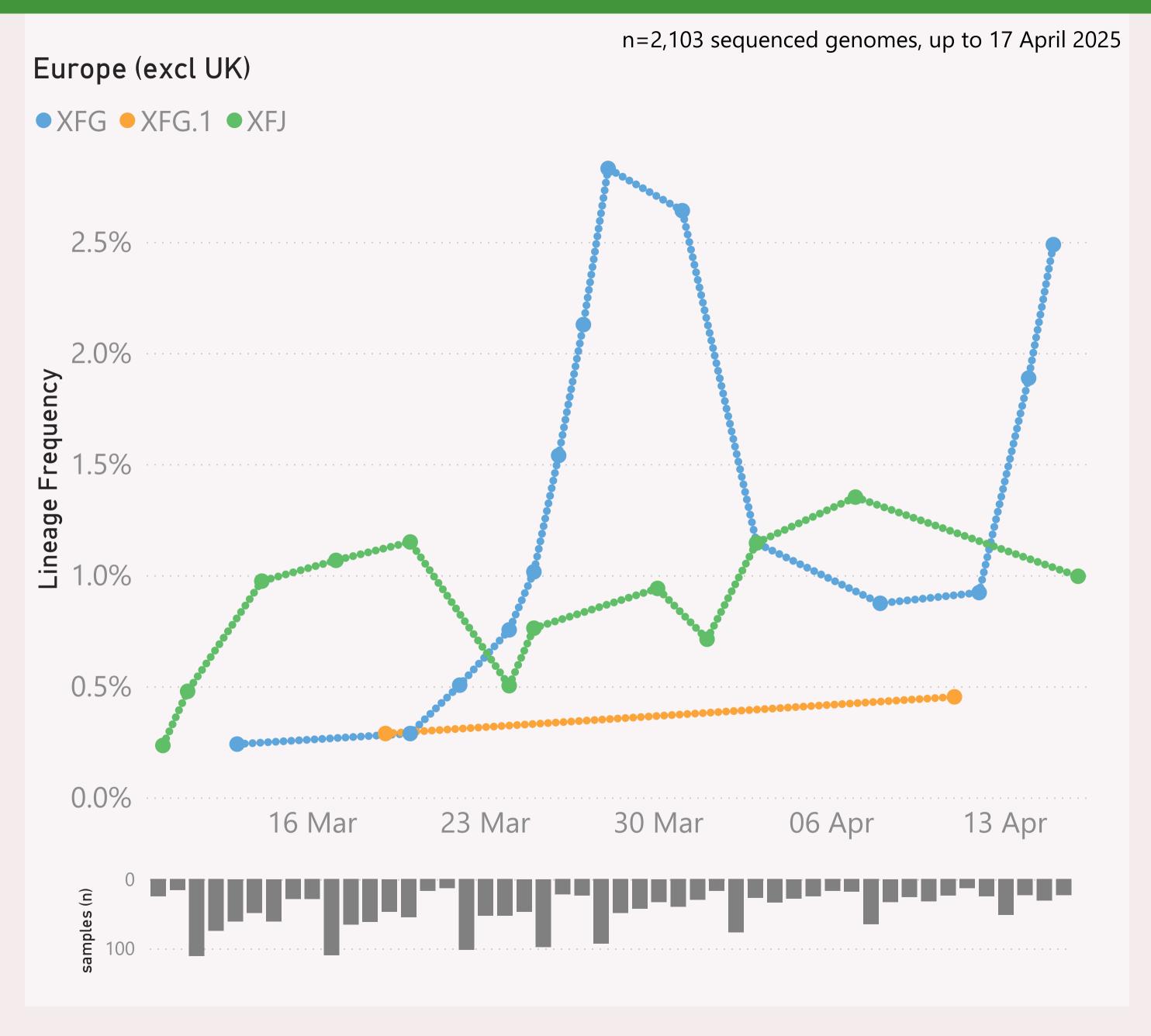
This page shows the frequency of the top 6 "L2" lineages, across recent months.

The detailed Lineage classifications are provided by Nextclade. I roll those up into "L2" groups, which roughly follow the WHO Variant definitions. For example, my "BA.2.86.\*" group includes BA.2.86 and all it's descendants, e.g. the JN.\* lineages.

The detailed Lineage classifications are quite numerous and dynamic, so the "Lineage L2" groups give a simpler and more stable basis for analysis and comparison.

The frequency shown at each point is based on the 7-day rolling average across all lineages.

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.

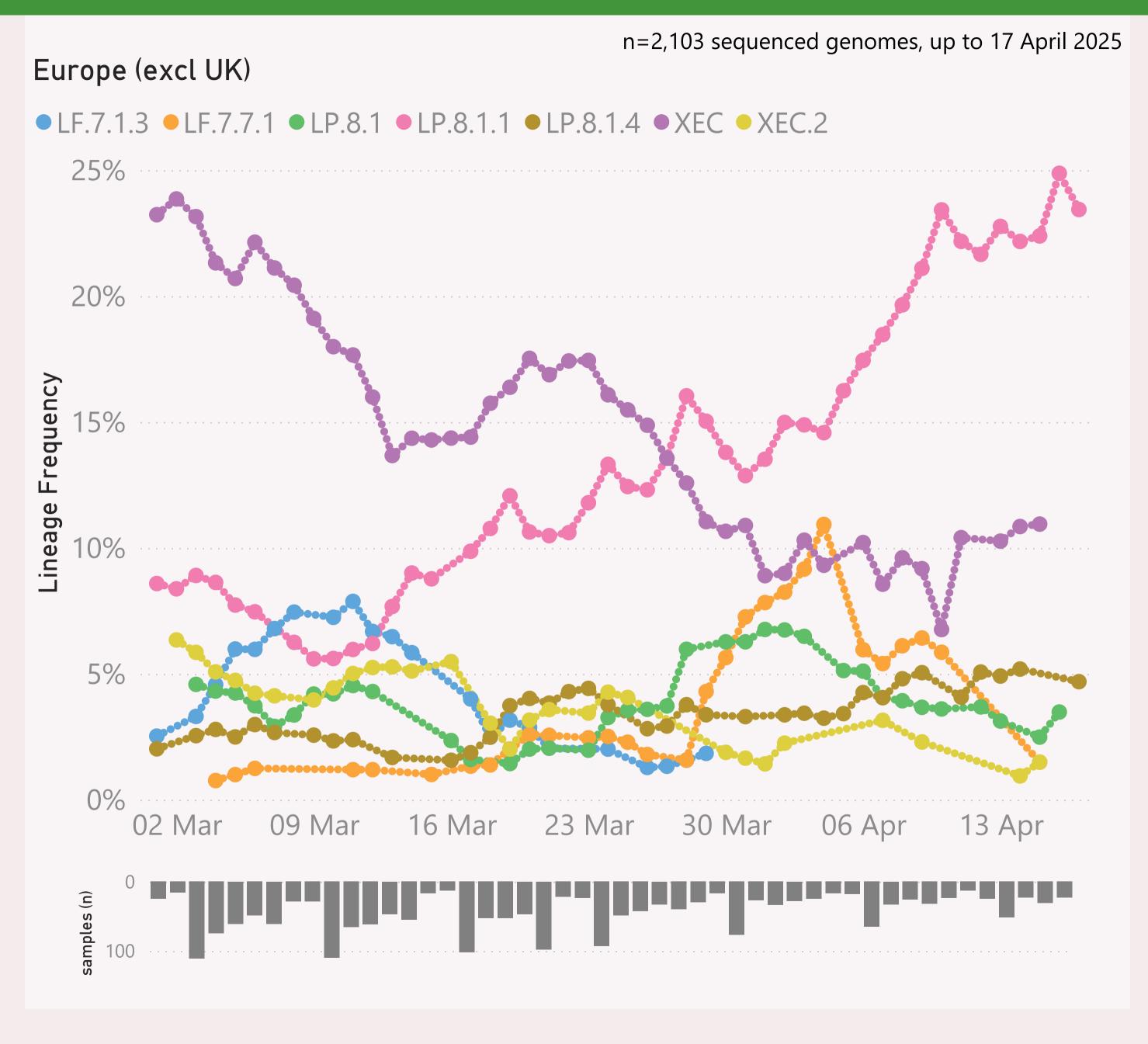


This page shows the frequency of the top 7 lineages, across recent months. The lineages are filtered for a "Lineage L2" group of interest, currently "XFG.\*" and "XFJ.\*".

The Lineage classifications are provided by Nextclade. The colour assignments are random.

The frequency shown at each point is based on the 7-day rolling average across all lineages.

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.



This page shows the frequency of the top 7 lineages, across recent months.

The Lineage classifications are provided by Nextclade. The colour assignments are random.

The frequency shown at each point is based on the 7-day rolling average across all lineages.

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.

XFG

XFG

XFG.1

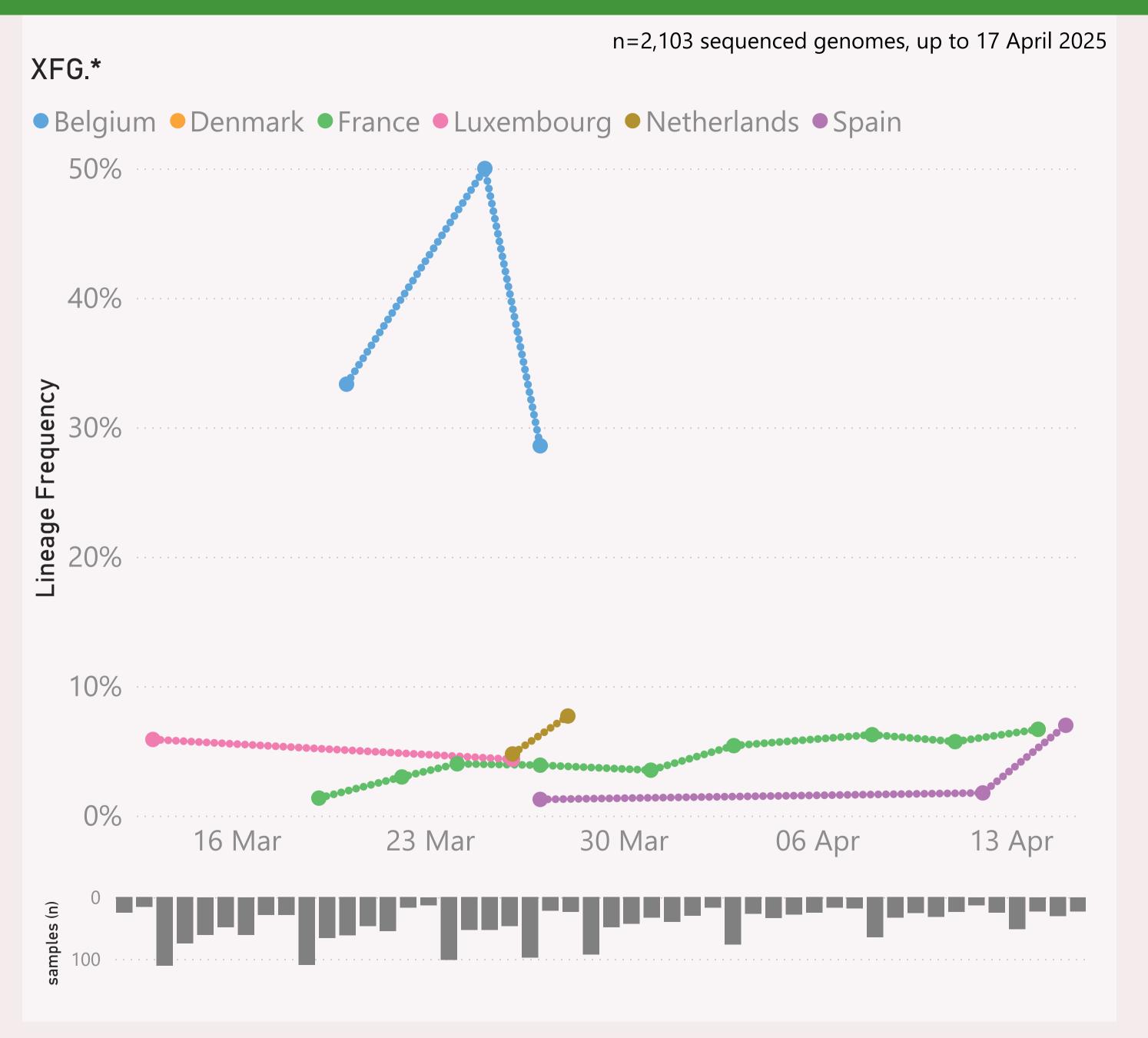
XFG

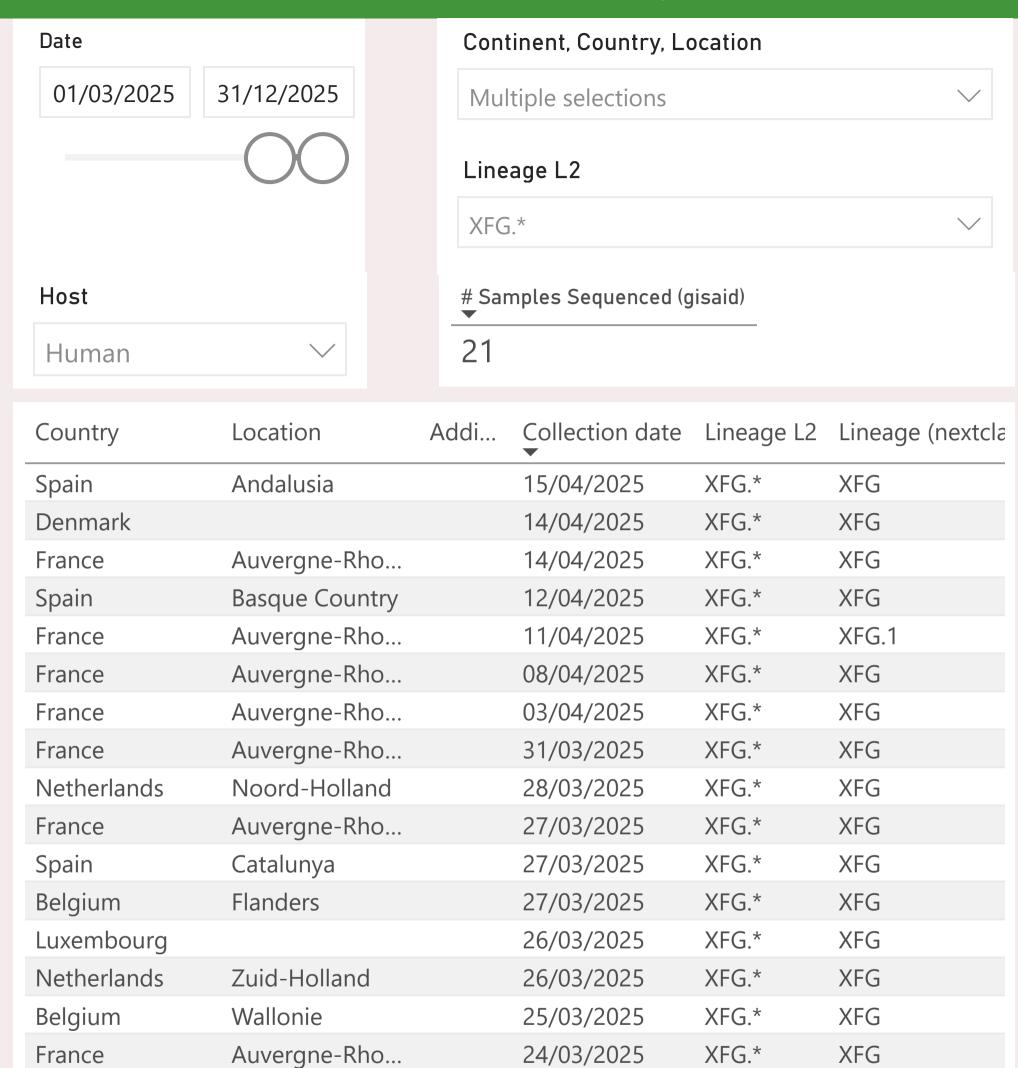
XFG.\*

XFG.\*

XFG.\*

XFG.\*





22/03/2025

20/03/2025

19/03/2025

13/03/2025

**Total** 

France

Belgium

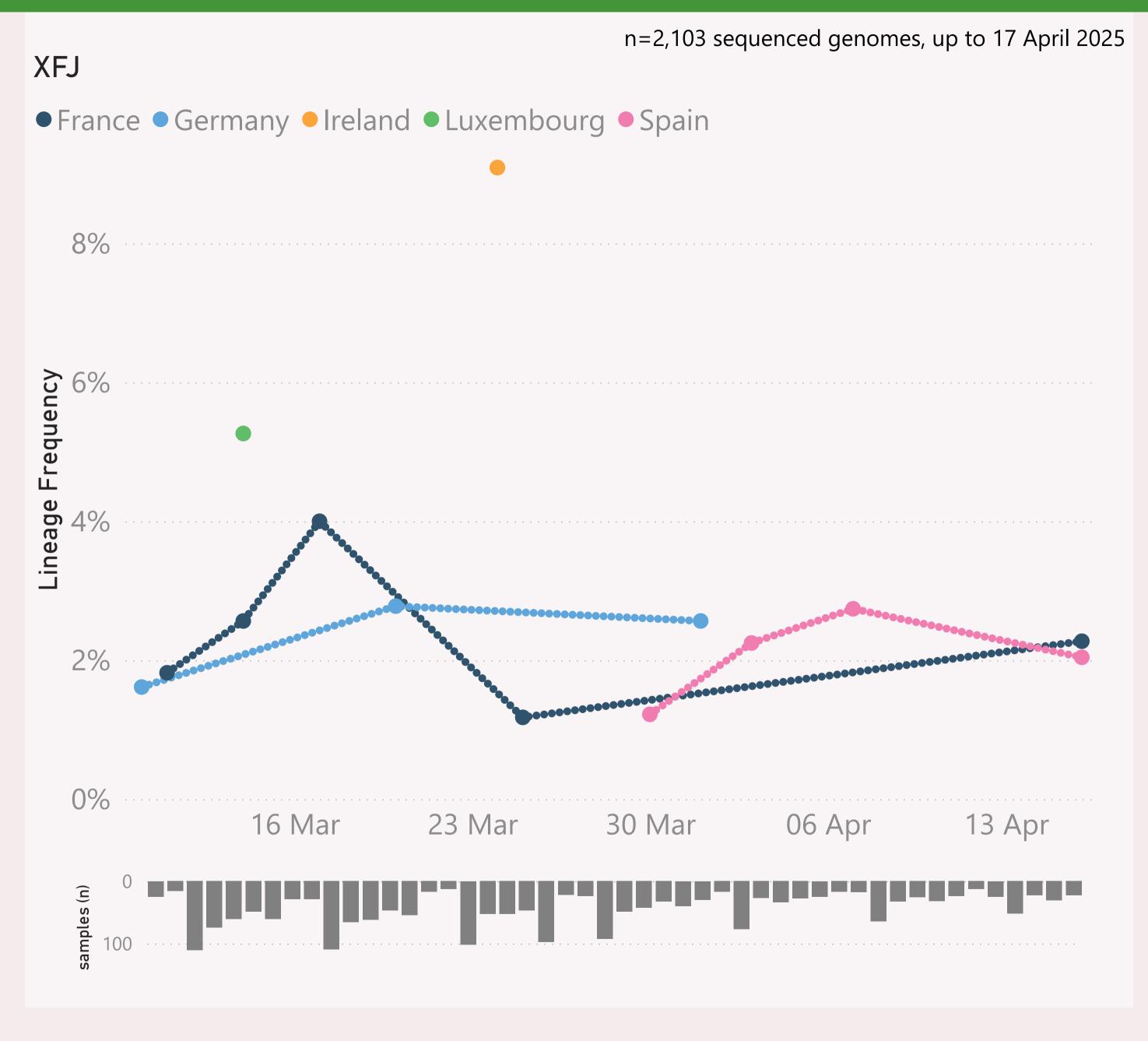
Luxembourg

France

Ile-de-France

Bruxelles

**Grand Est** 

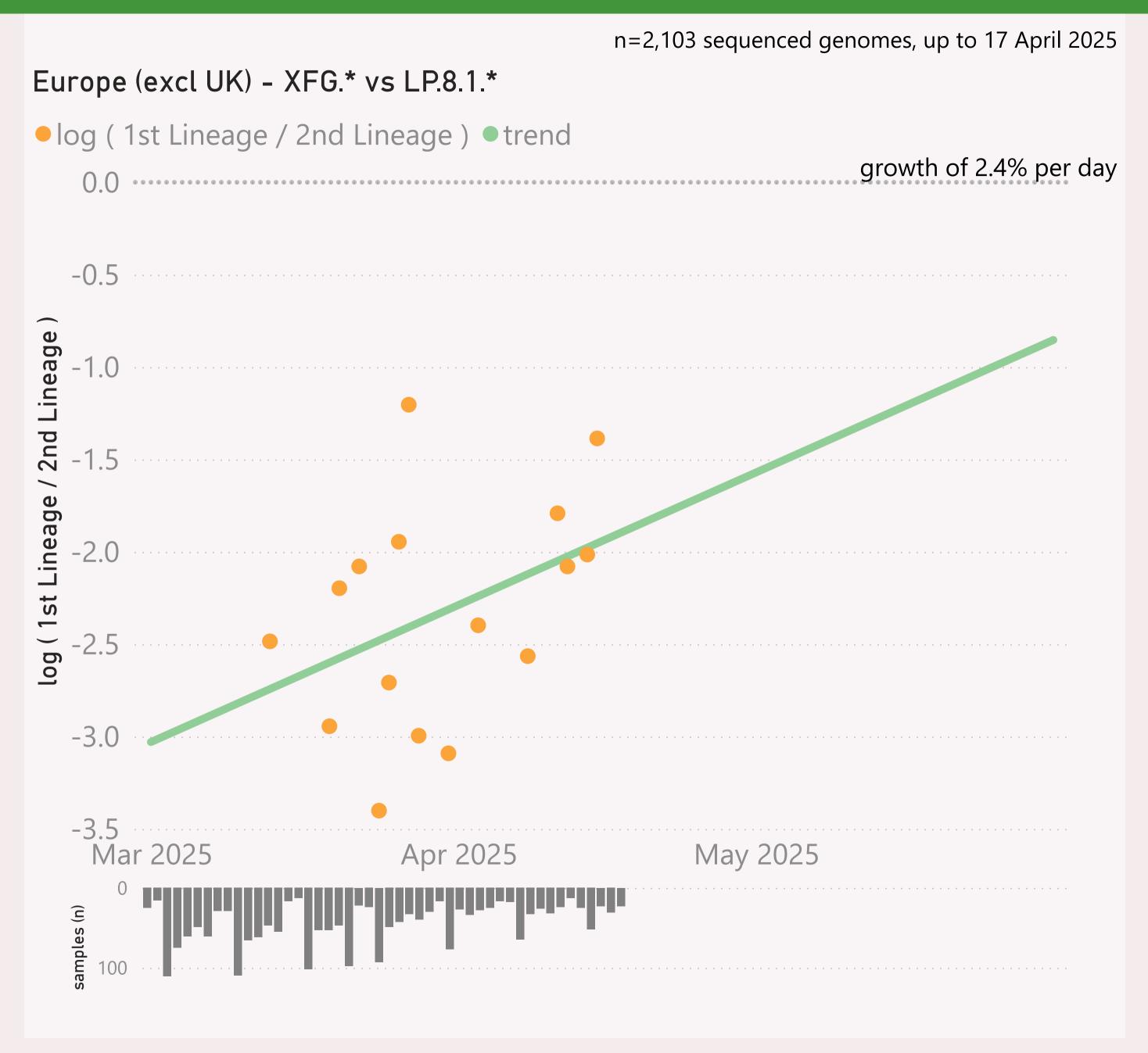


This page shows the frequency of a selected Lineage of interest, for the 7 countries reporting the most samples over recent months.

The Lineage classifications are provided by Nextclade.

The frequency shown at each point is based on the 7-day rolling average across all lineages, for that country.

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.

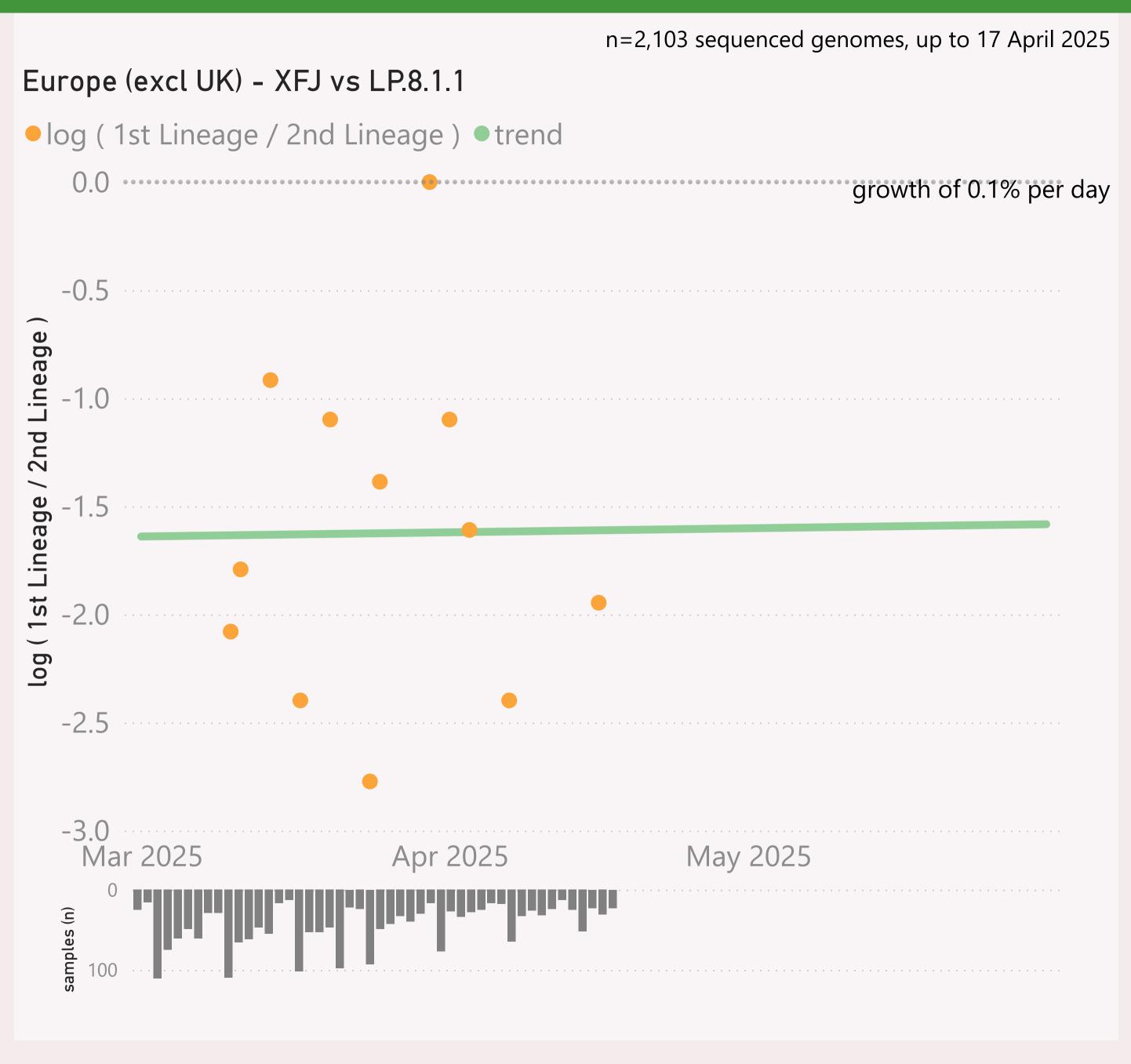


This page compares the relative frequency of 2 selected "Lineage L2" groups, over recent months. A challenging Lineage L2 is selected first, and compared to the incumbent.

The trend is shown as a green line and expressed as a daily growth % advantage. If the green line crosses over the 0.0 line, the date when that occurred or is predicted to occur will be shown. At that point the challenging Lineage L2 is considered to have "crossed over" or taken over dominance from the incumbent Lineage L2.

The Lineage classifications are provided by Nextclade. I add the "Lineage L2" groups, typically following common variant groupings, but occasionally being "creative".

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.

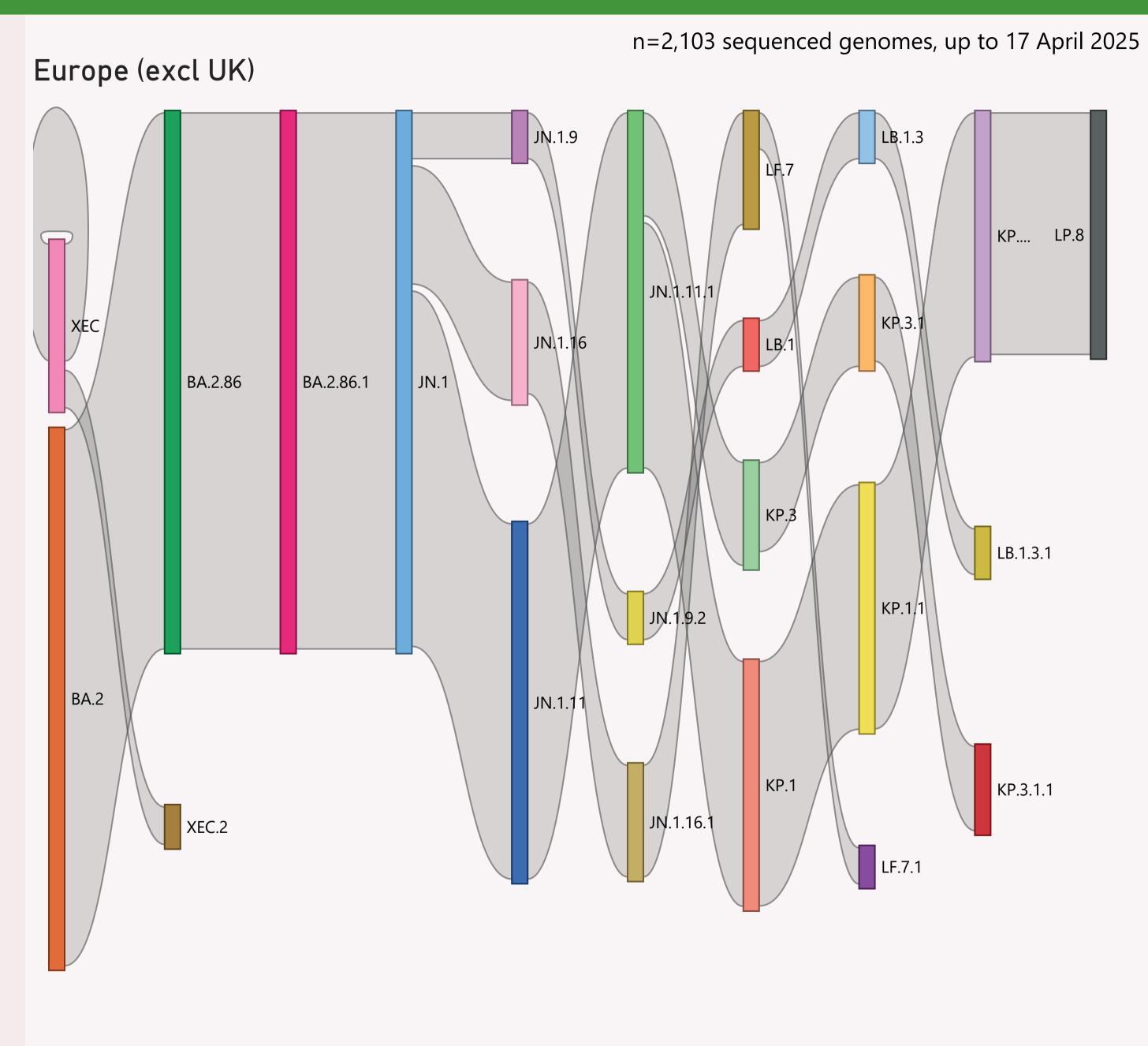


This page compares the relative frequency of 2 selected Lineages, over recent months. A challenging Lineage is selected first, and compared to the incumbent.

The trend is shown as a green line and expressed as a daily growth % advantage. If the green line crosses over the 0.0 line, the date when that occurred or is predicted to occur will be shown. At that point the challenging Lineage is considered to have "crossed over" or taken over dominance from the incumbent Lineage.

The Lineage classifications are provided by Nextclade.

The grey column chart across the bottom shows the volume of sequences available by date. As there can be long sample and data processing times, it is quite routine for recent dates to show lower sample sizes.



This page shows the hierarchy of the significant Lineages, over recent months.

The hierarchy can be read from left to right, starting with the earliest/highest Lineages being broken down into more detailed child Lineages.

The vertical height of each bar segment represents the relative volume of all the samples of that specific Lineage, as well as all it's descendants.

The full picture is typically quite busy, so insignificant Lineages (with few samples, or at the extreme top or bottom of the hierarchy) are not shown.

The Lineage classifications are provided by Nextclade.

## Data Submitted in the last 8 weeks

Country	# Samples Sequenced	Latest Collection date	by Collection date	Latest Submission date	by Submission date
⊕ France	1,252	17/04/2025		24/04/2025	
	956	17/04/2025		24/04/2025	ates and beautiful and
	390	01/04/2025	A bear	16/04/2025	
⊕ Germany	329	15/04/2025	i II.	24/04/2025	
⊕ Greece	320	04/04/2025	<b>Al</b> te.	15/04/2025	
	240	10/04/2025	ra	24/04/2025	La constantina de la constantina della constanti
	226	14/04/2025	1	24/04/2025	
⊕ Russia	192	29/03/2025		24/04/2025	
	181	17/04/2025	. 4	24/04/2025	
⊕ Ireland	136	17/04/2025		24/04/2025	a data ca
⊕ Finland	131	24/03/2025		09/04/2025	
Sweden	126	17/04/2025	441	24/04/2025	
	123	21/03/2025	ılı	04/04/2025	
⊕ Czechia	74	09/12/2024		24/03/2025	
⊕ Croatia	69	21/03/2025		10/04/2025	
⊕ Portugal	54	14/04/2025	41	24/04/2025	
⊞ Belgium	48	05/04/2025	ı İn	24/04/2025	
⊞ Slovenia	44	26/03/2025		24/04/2025	
	41	16/04/2025	7	24/04/2025	
	9	04/04/2025		24/04/2025	
	7	28/02/2025		31/03/2025	1.
⊞ Serbia	6	09/01/2025		04/04/2025	
	5	11/03/2025		22/04/2025	
	4	28/01/2025		25/03/2025	
<b>Romania</b>	2	14/03/2025		21/03/2025	
Total	4,965	17/04/2025	عالمها والمالية	24/04/2025	أعام الماعلية الماعلة

This page shows the volume and currency/timeliness of the genomic sequencing data shared via GISAID, over the last 8 weeks, for the countries sharing the most samples.

Each sample shared comes with a Collection date - when the PCR test for that sample was collected. The GISAID system also records a Submission date for each sample, which is typically the date that sample was uploaded.

The latest date of each type is shown, along with "sparkline"-style mini charts to give a flavour for the spread of recent data by Collection date and by Submission date.