

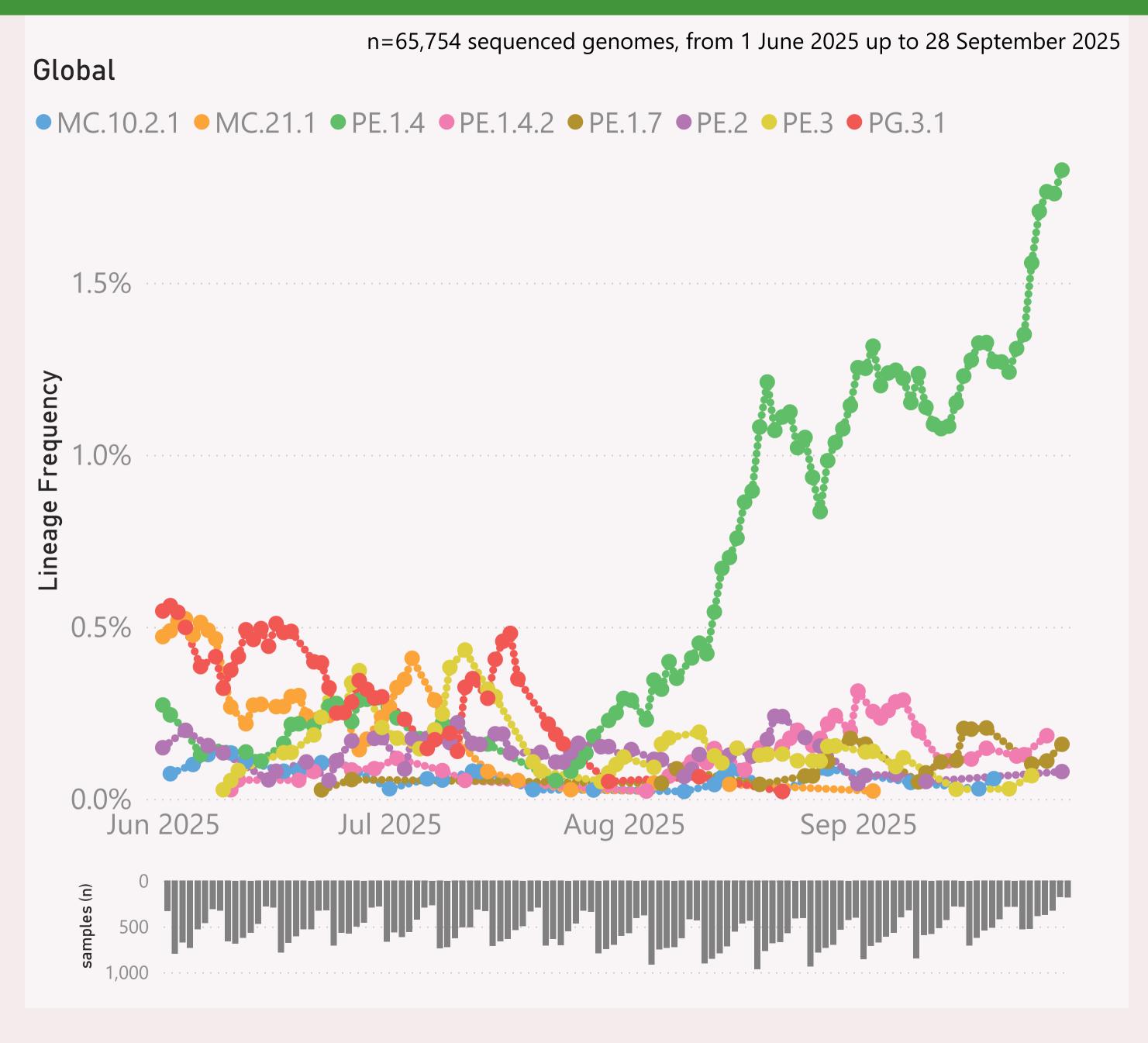
This page shows the frequency of the top 6 "L2" lineages, across recent months.

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.

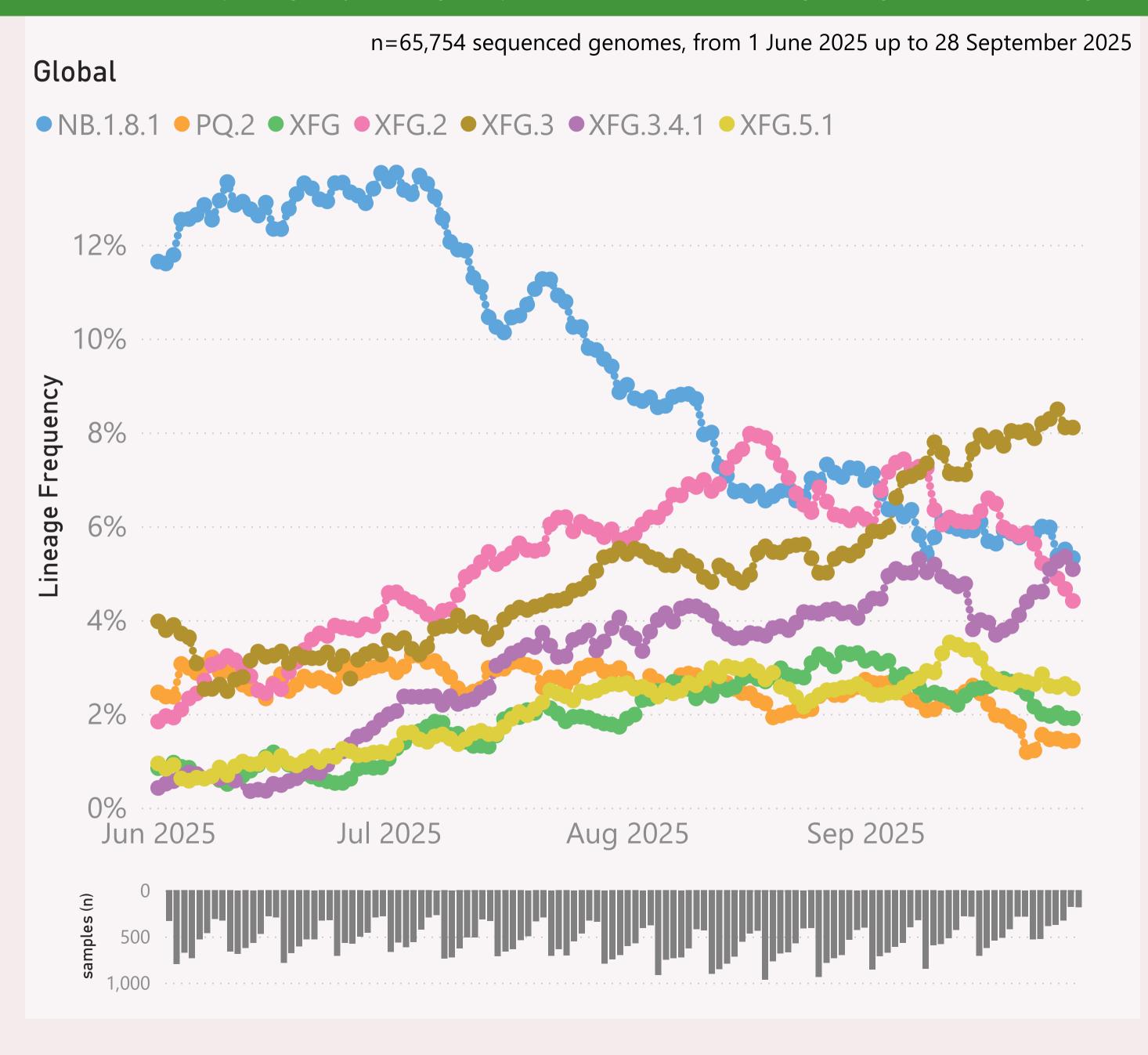


This page shows the frequency of the top 7 lineages, across recent months. The lineages are filtered for a "Lineage L2" group of interest.

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

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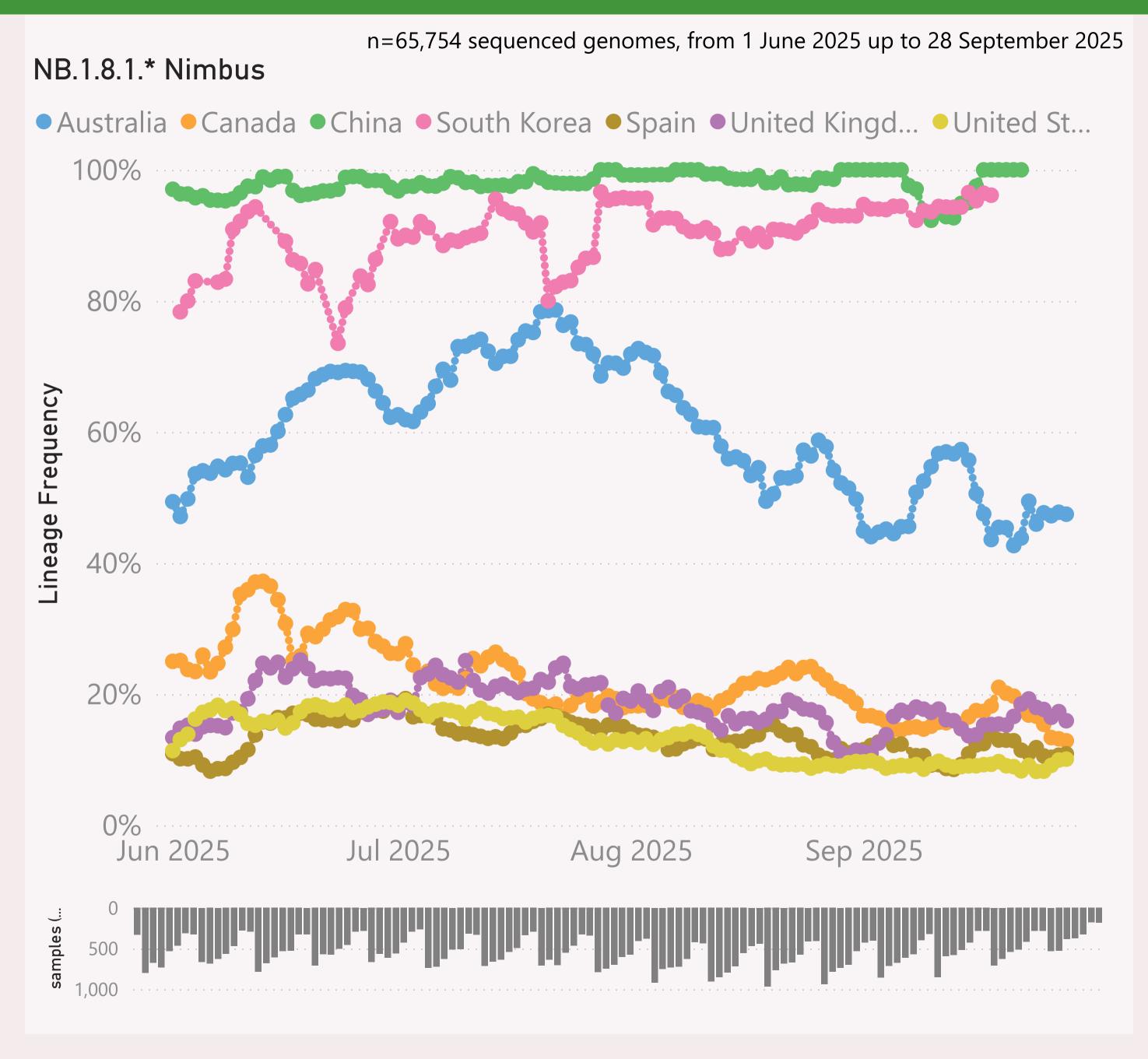


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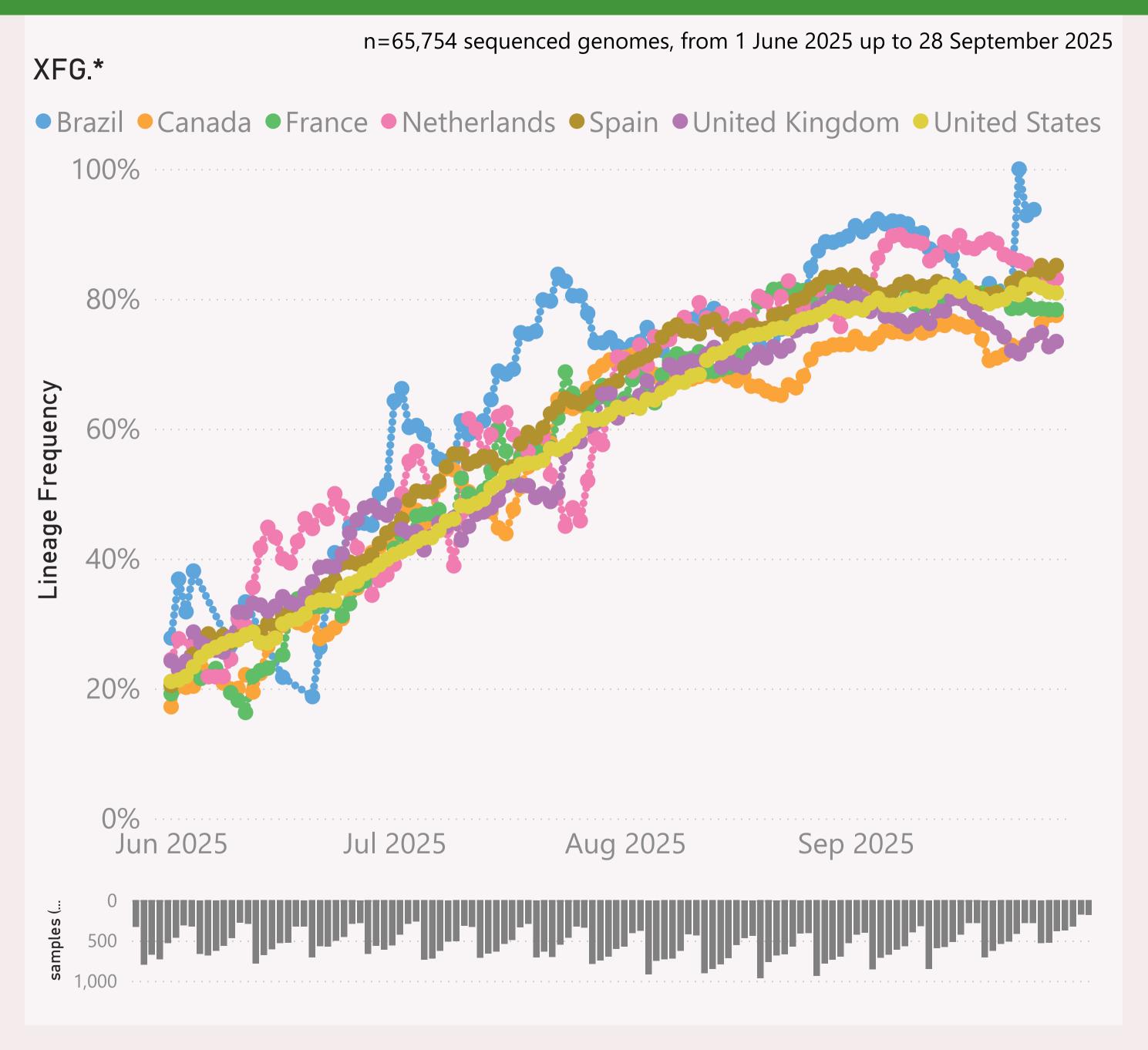
This page shows the frequency of a selected "Lineage L2" group of interest, for the 7 countries reporting the most samples over recent months.

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 "JN.1.\* +FLiRT" group includes the descendants of JN.1.\* with the mutations: F456L & R346T.

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, for that country.

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XFP

XFP

XFP

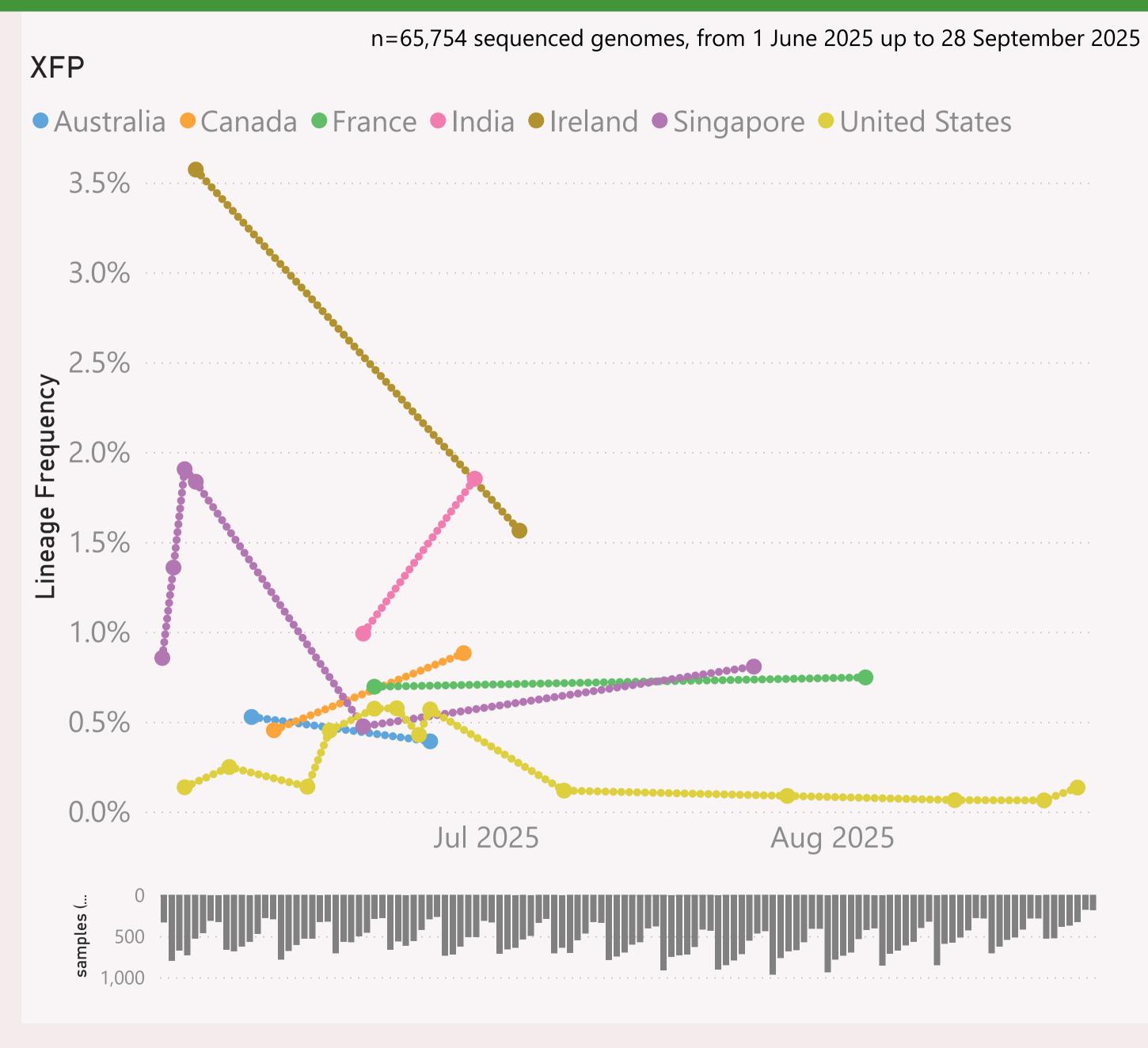
XFP

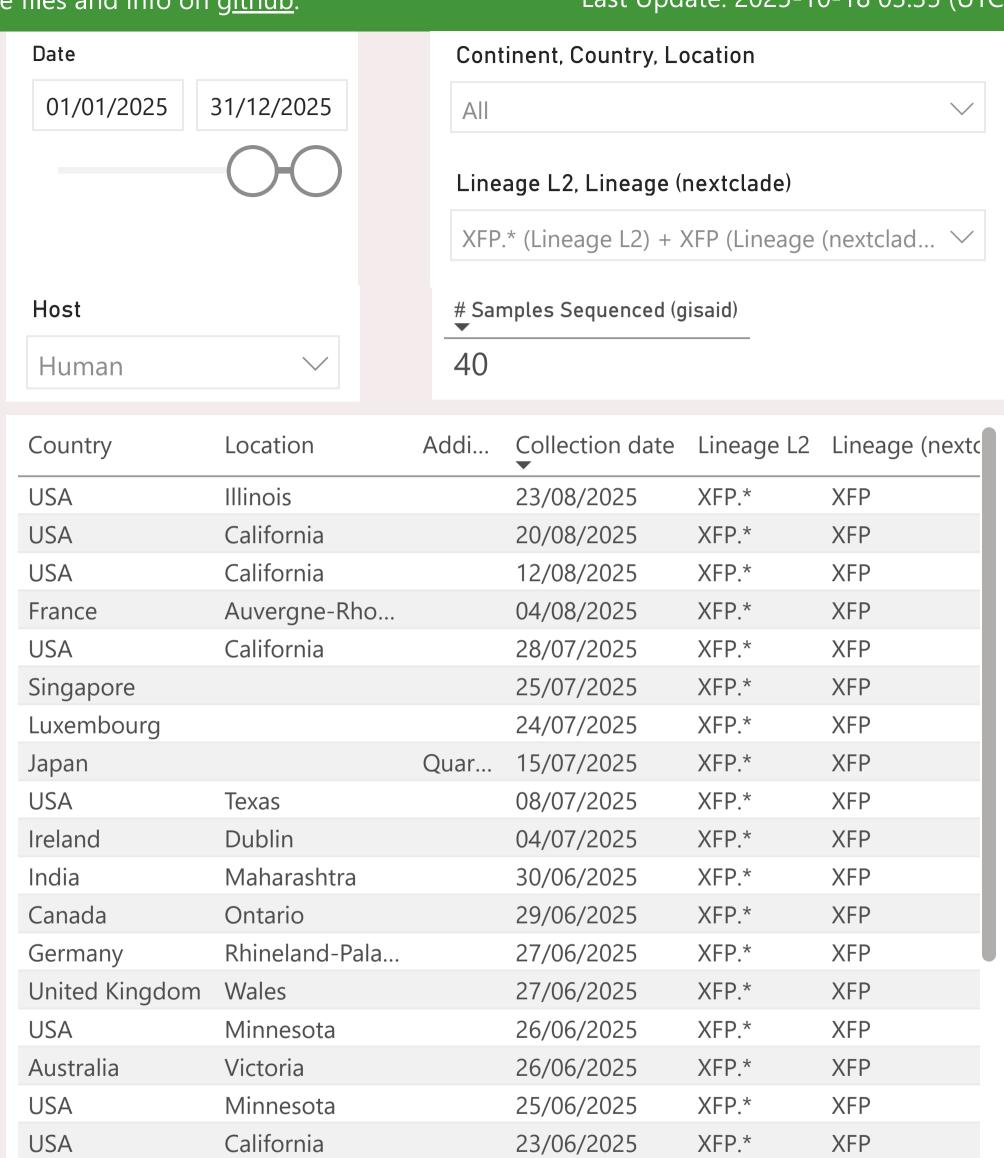
XFP.\*

XFP.\*

XFP.\*

XFP.\*





21/06/2025

21/06/2025

20/06/2025

20/06/2025

USA

France

India

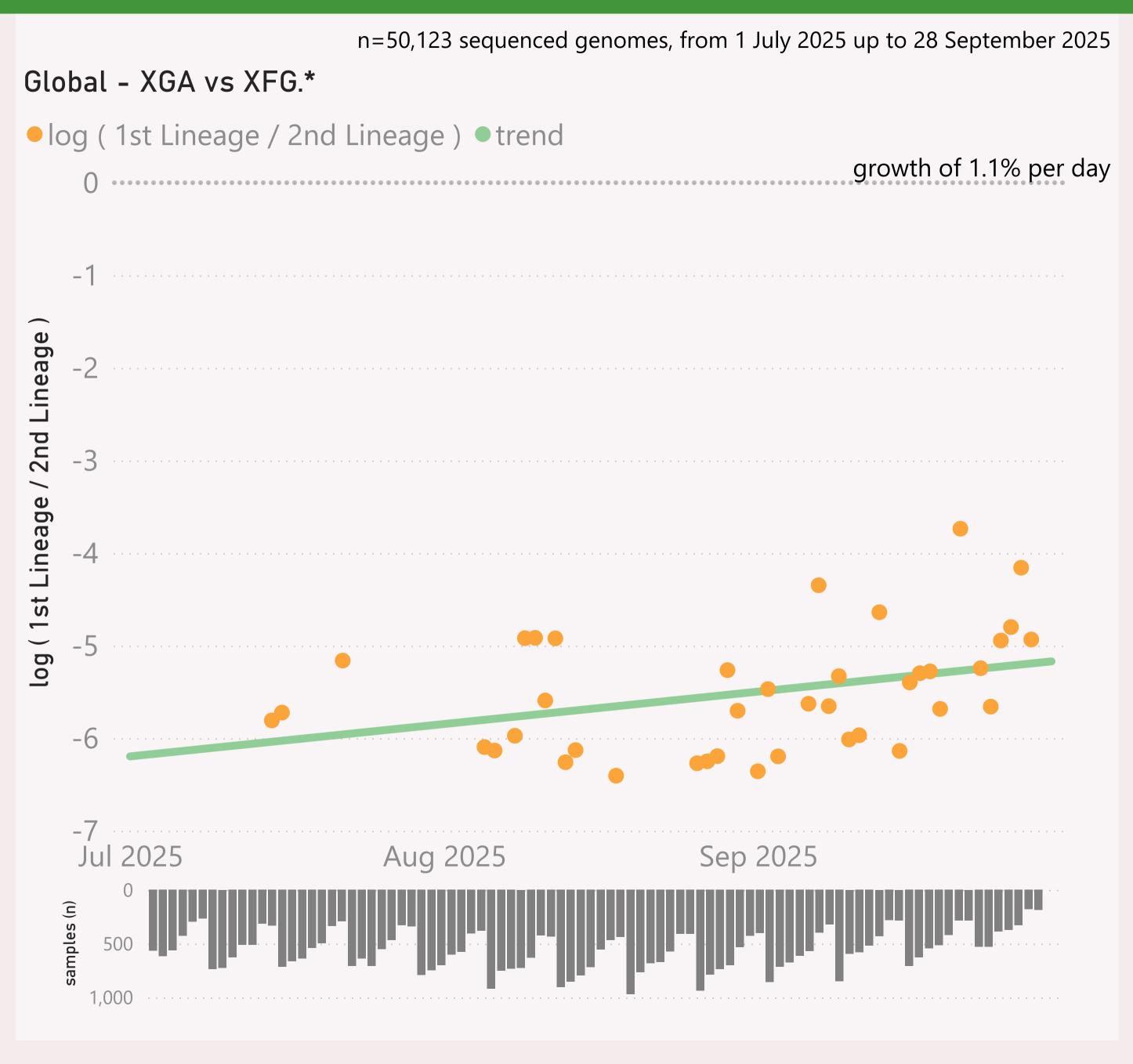
**Total** 

Singapore

Oregon

Provence-Alpe...

Chhattisgarh

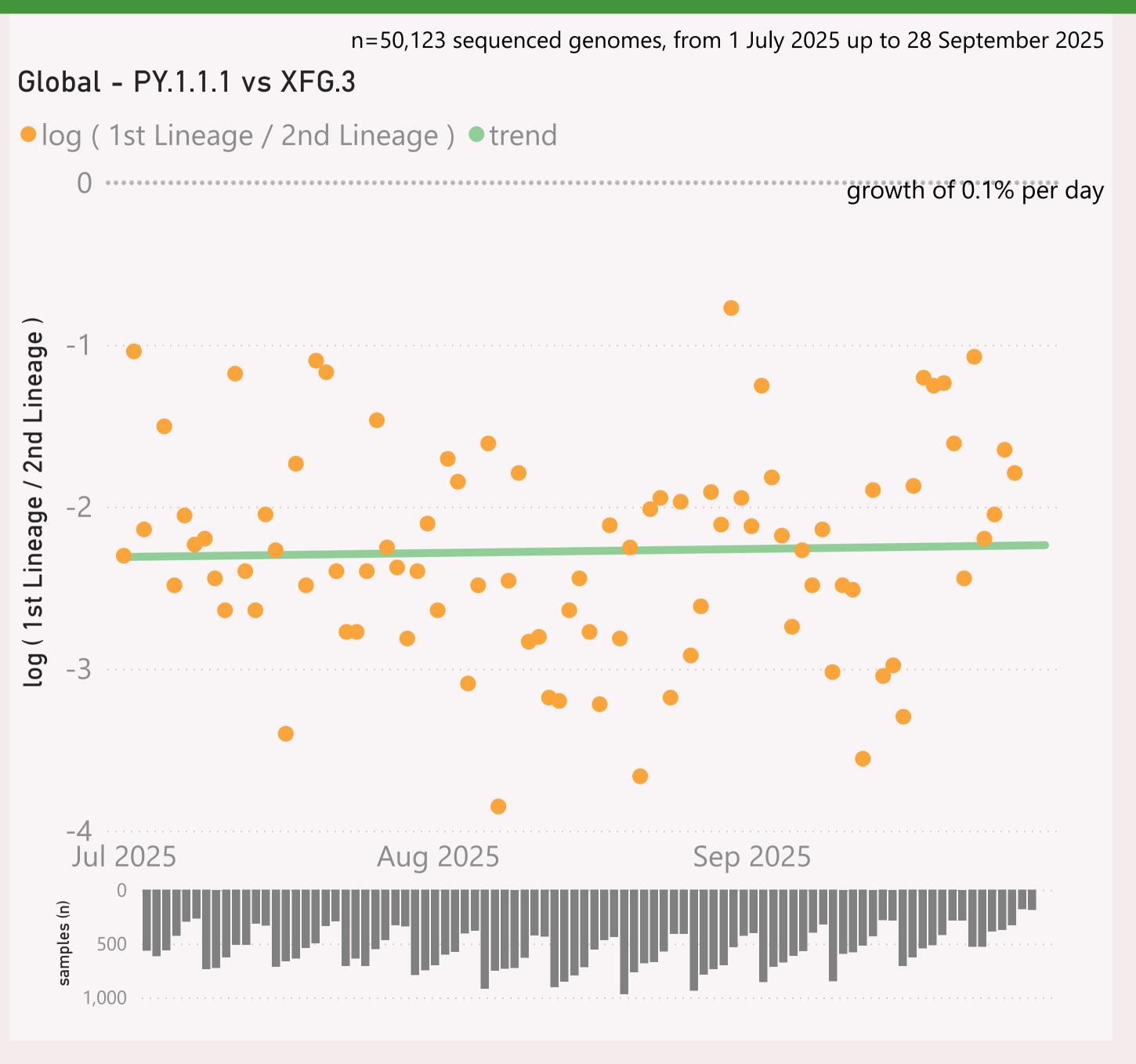


This page compares the relative frequency of 2 selected "Lineage L2" groups, over recent months. A challenging Lineage L2 is selected first, and compared to the incumbent.

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.

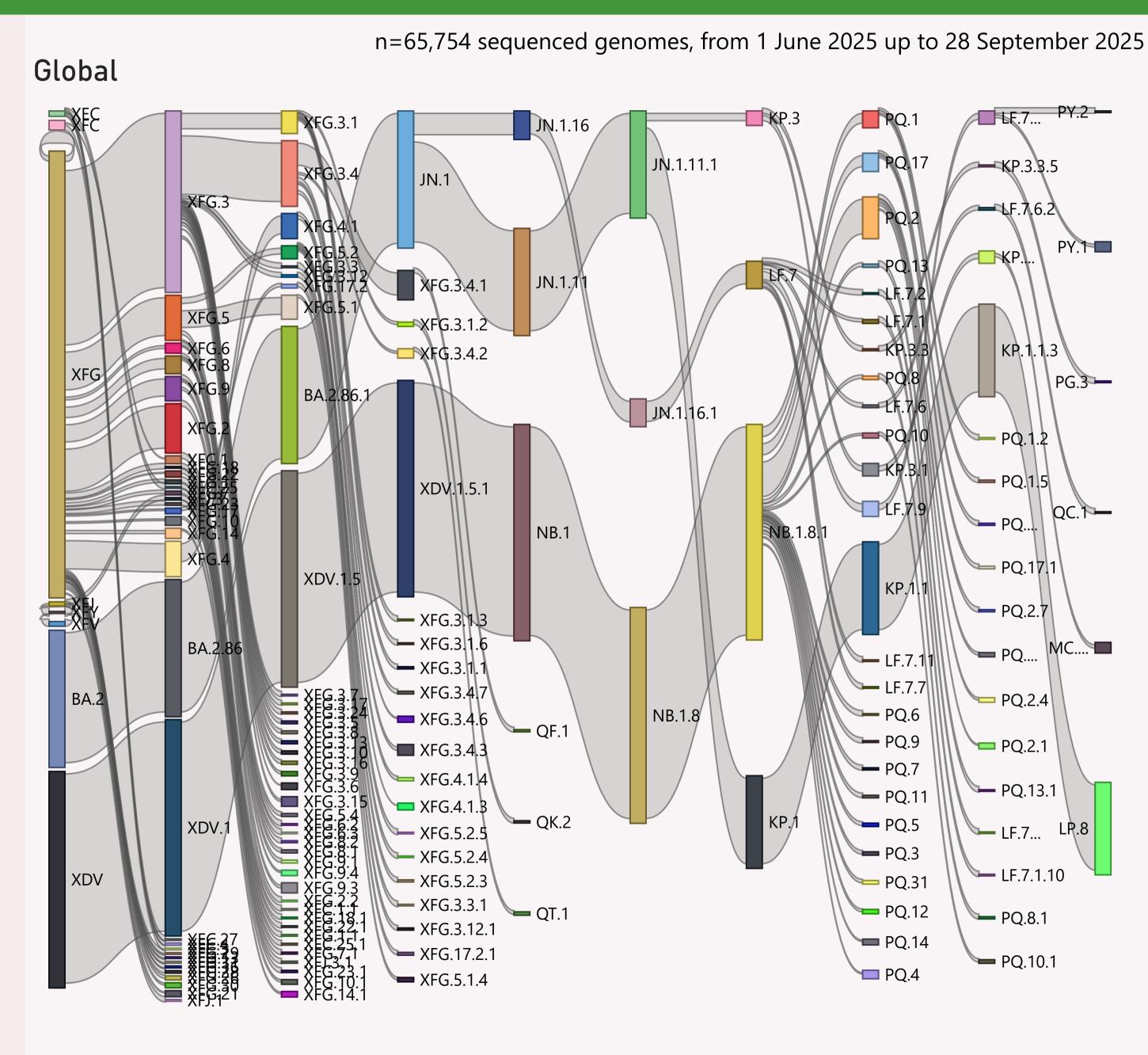


This page compares the relative frequency of 2 selected Lineages, over recent months. A challenging Lineage is selected first, and compared to the incumbent.

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.



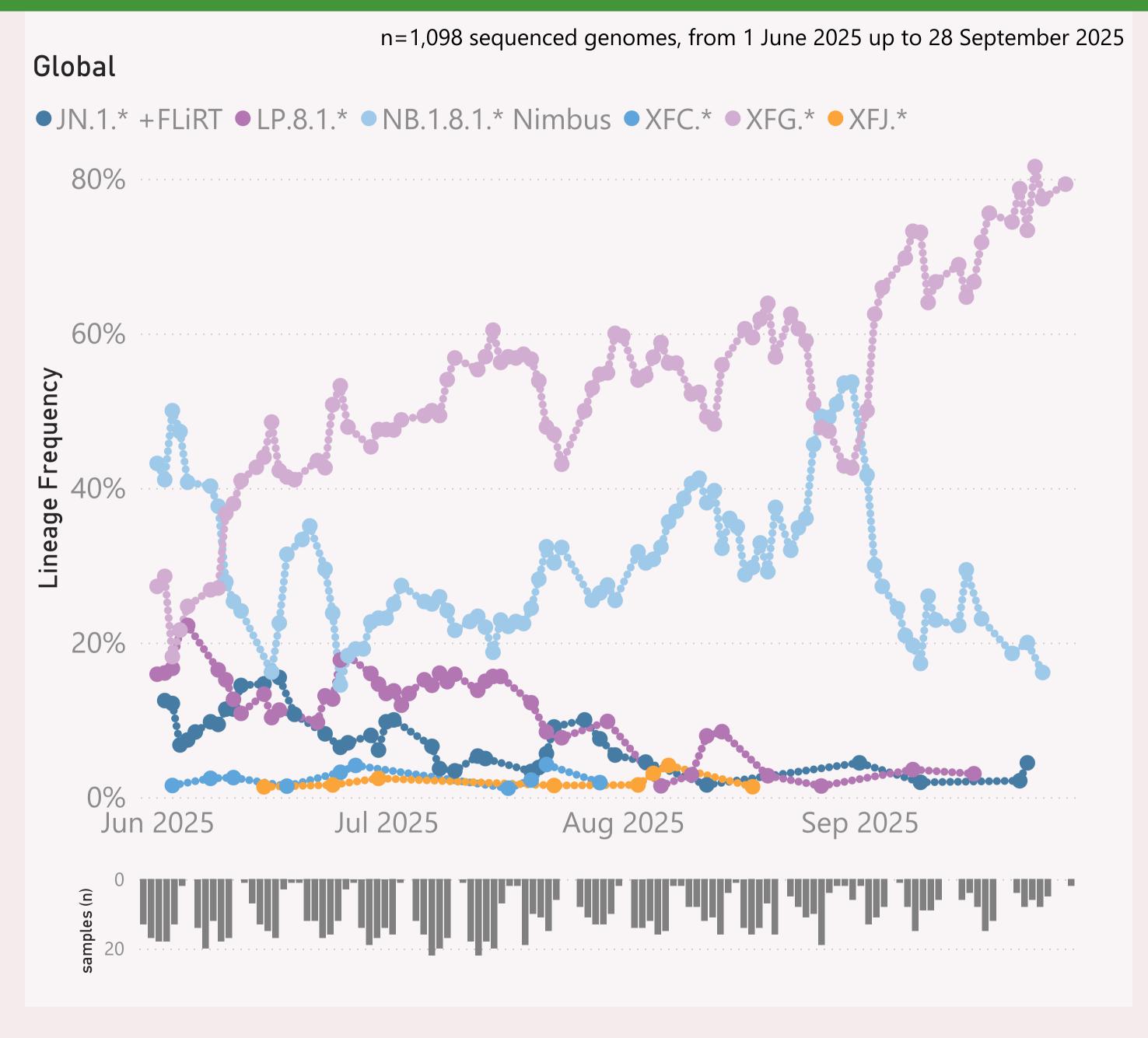
This page shows the hierarchy of the significant Lineages, over recent months.

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.



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

This is probably a more randomised sample than the "Global" aggregate of all samples submitted to GISAID, as those are dominated by the US and Canada

These samples are mainly collected from arrivals into the US and Japan.

## Data Submitted in the last 8 weeks

Country	# Samples Sequenced	Latest Collection date	by Collection date	Latest Submission date	by Submission date
	12,549	28/09/2025		08/10/2025	de de adame la calell
⊞ Spain	4,675	28/09/2025	A STATE OF THE PARTY OF THE PAR	08/10/2025	أبريور والباسور ويوايان
⊕ Canada	4,286	28/09/2025		08/10/2025	أعدا يمتنينا
	3,293	28/09/2025	datilikka	08/10/2025	المصلحات ما
	1,942	28/09/2025	بالماسينين والمستعدد	08/10/2025	أبا بشيرات
	1,720	28/09/2025	and the second second	08/10/2025	إحتناه مبايدا مخا
	1,306	18/09/2025	ـ المالم الماليات	08/10/2025	., ( ), ( ), [ ], [ ]
⊞ China	1,280	22/09/2025	and the second state of the second	08/10/2025	بالمالية بالمالية
⊕ Brazil	1,207	25/09/2025	and a substitution of	08/10/2025	and the filler and contributed
	881	28/09/2025	والأنطالي ومسور والمسا	08/10/2025	النابيا ا
⊞ Japan	733	28/09/2025	act association filtera	08/10/2025	and all and the architect
	645	28/09/2025	og hittisik	08/10/2025	ang madaasa a atal
	551	28/09/2025		08/10/2025	adit in a company
⊞ Ireland	545	28/09/2025	and distributions	08/10/2025	call and raid
⊕ Germany	542	28/09/2025	e lideklida ja are	08/10/2025	أرا بالمناسيا
⊞ Russia	438	13/09/2025		06/10/2025	الباز با
	425	28/09/2025	. antibalikara	08/10/2025	1   1   1
	380	22/09/2025	.aulj	08/10/2025	, , , , , , , , ,
	363	31/08/2025	adomith Ad	22/09/2025	
	304	25/09/2025	and the same	08/10/2025	
	243	03/08/2025	defalla.	29/08/2025	
⊞ Sweden	239	28/09/2025	adákhaga	08/10/2025	
	236	28/09/2025	وأفه بإنصاد ومرو	08/10/2025	
	234	15/09/2025	Committee of the Commit	08/10/2025	
⊕ Costa Rica	233	25/09/2025	المتعومالسالين	08/10/2025	
	180	11/09/2025	articula Marchaela (	02/10/2025	
⊞ Romania	172	22/09/2025	والطاوات بير	08/10/2025	1. i - 1
	167	30/08/2025	والمتعلقية الأطواف	08/10/2025	
Total	41,824	28/09/2025		08/10/2025	. I .n.lntnh.llt.nhl

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.