

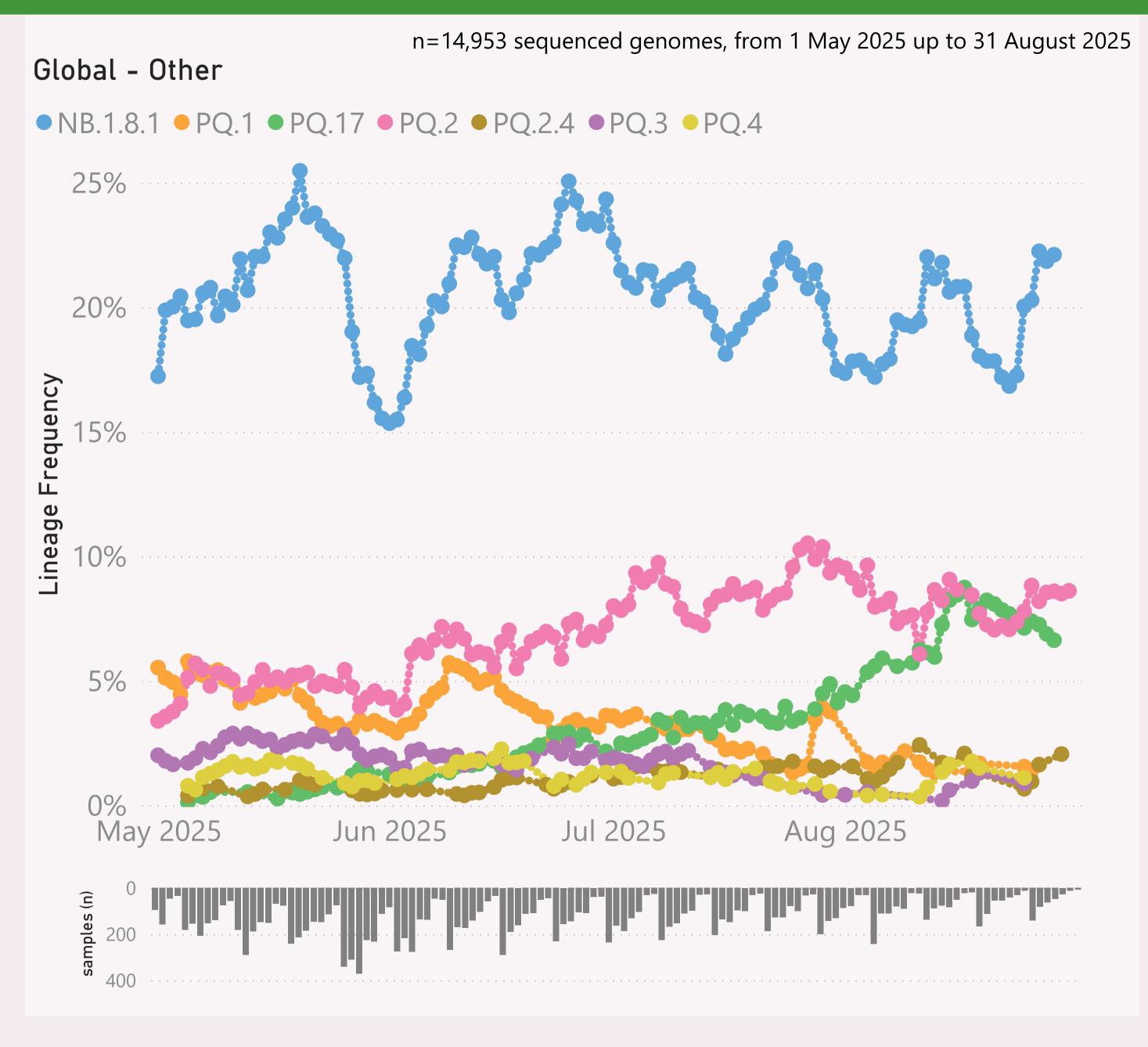
This "Global - Other" report aggregates the available data from countries besides those I regularly report on: Australia, NZ, the US, Canada and Europ.

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 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.

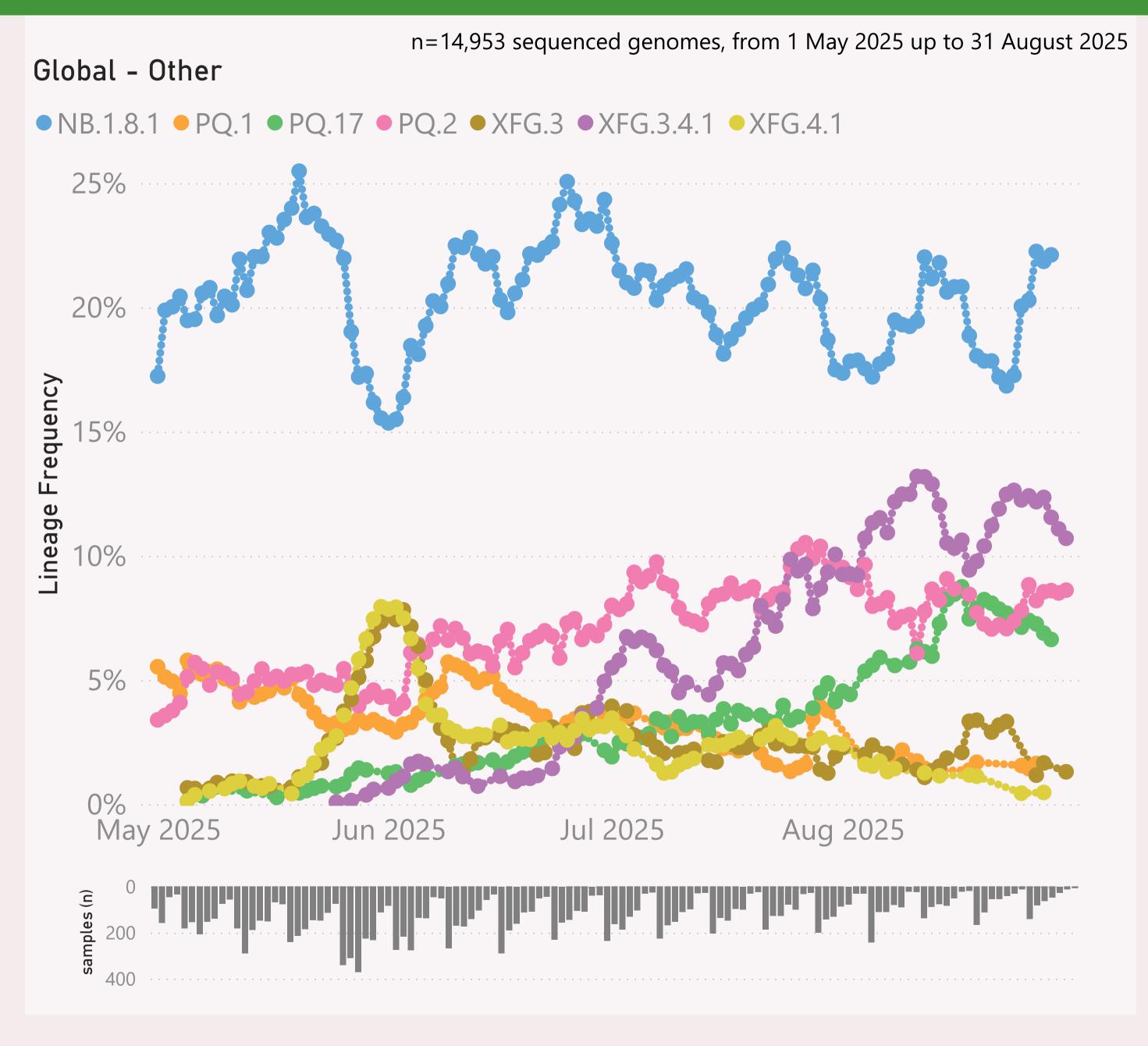


This page shows the frequency of the top 7 lineages, across recent months. The lineages are filtered for a "Lineage L2" group of interest, currently NB.1.8.1.* Nimbus.

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.

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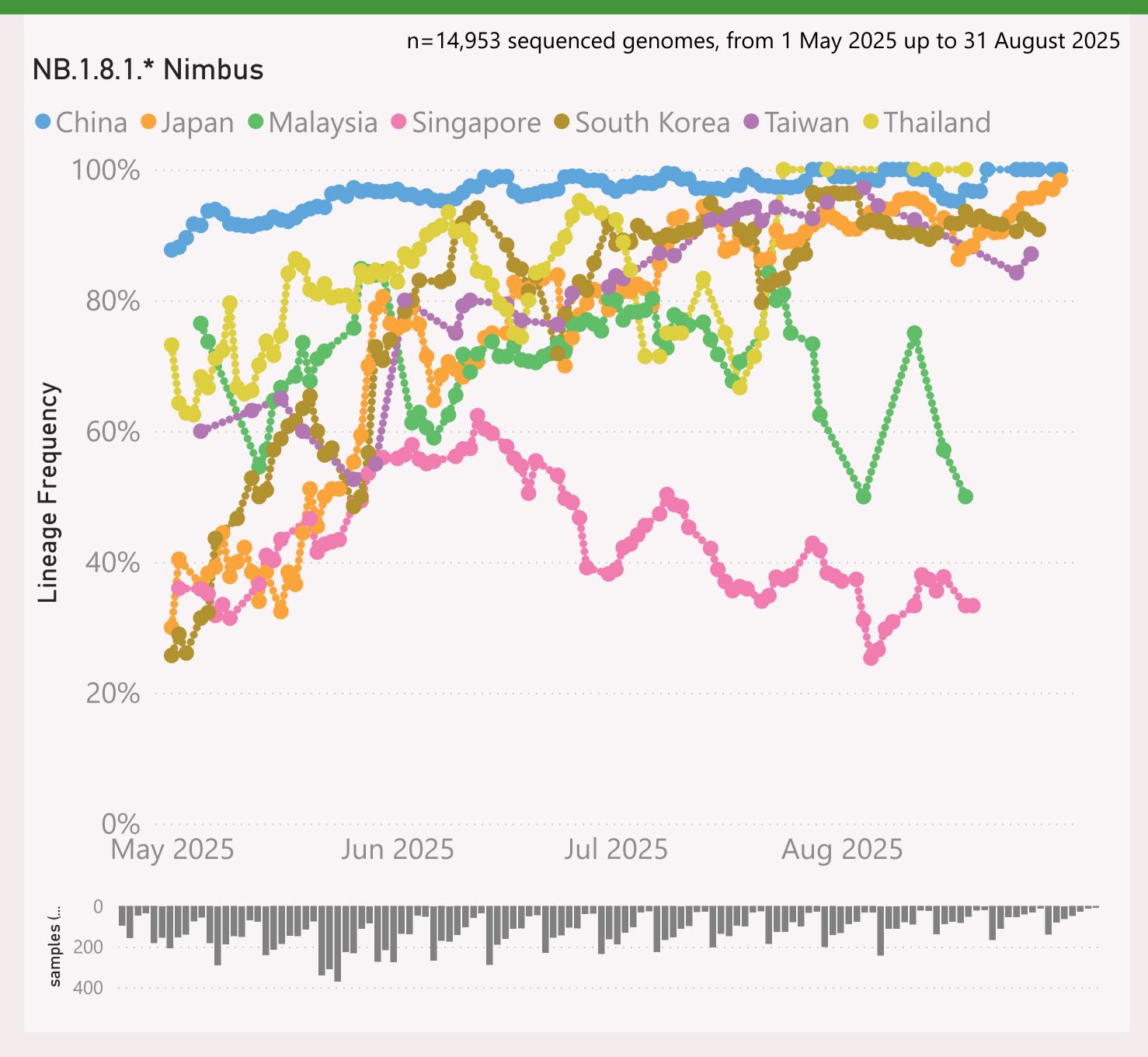


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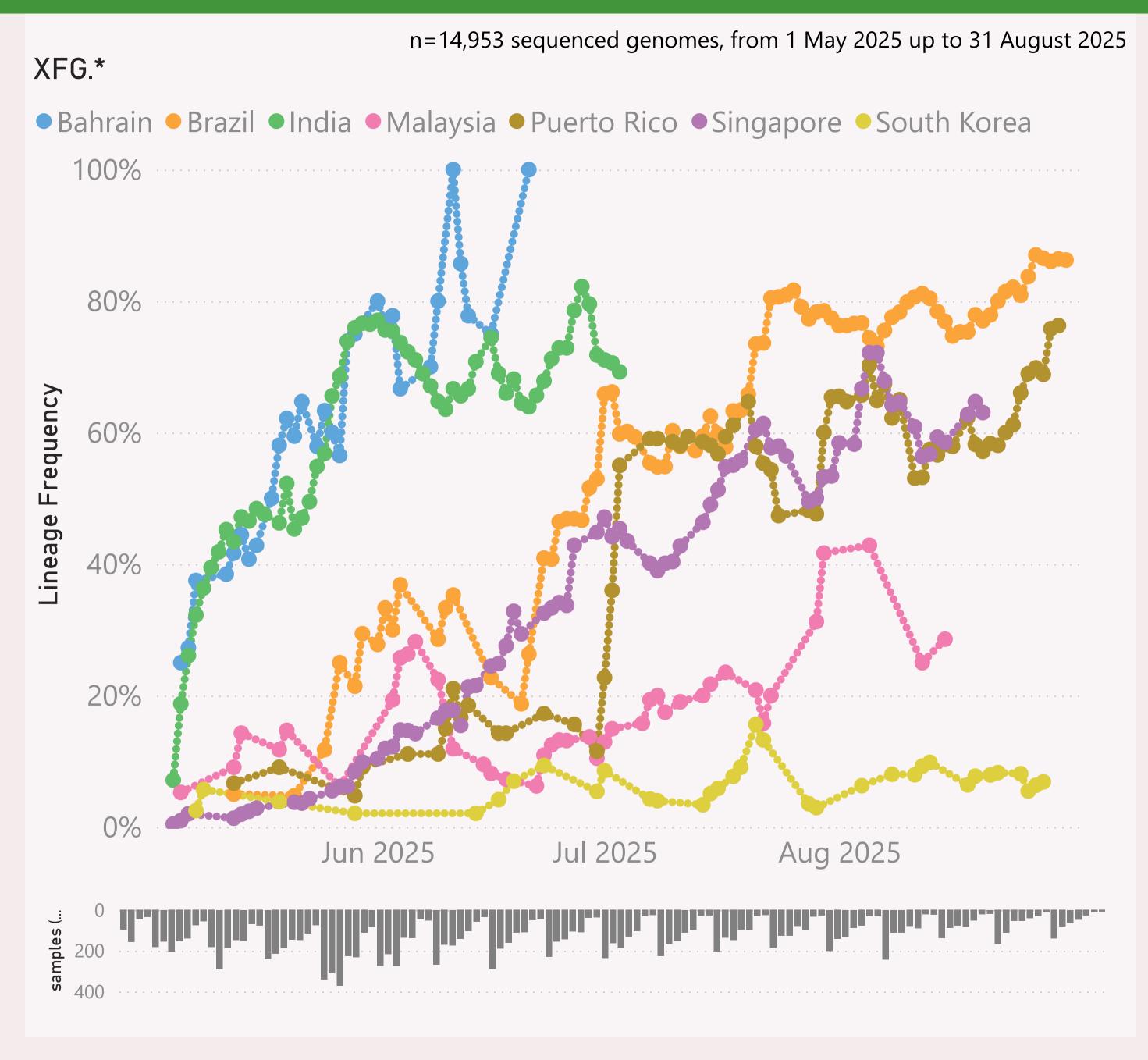
This page shows the frequency of a selected "Lineage L2" group of interest, for the 7 countries reporting the most samples over recent months.

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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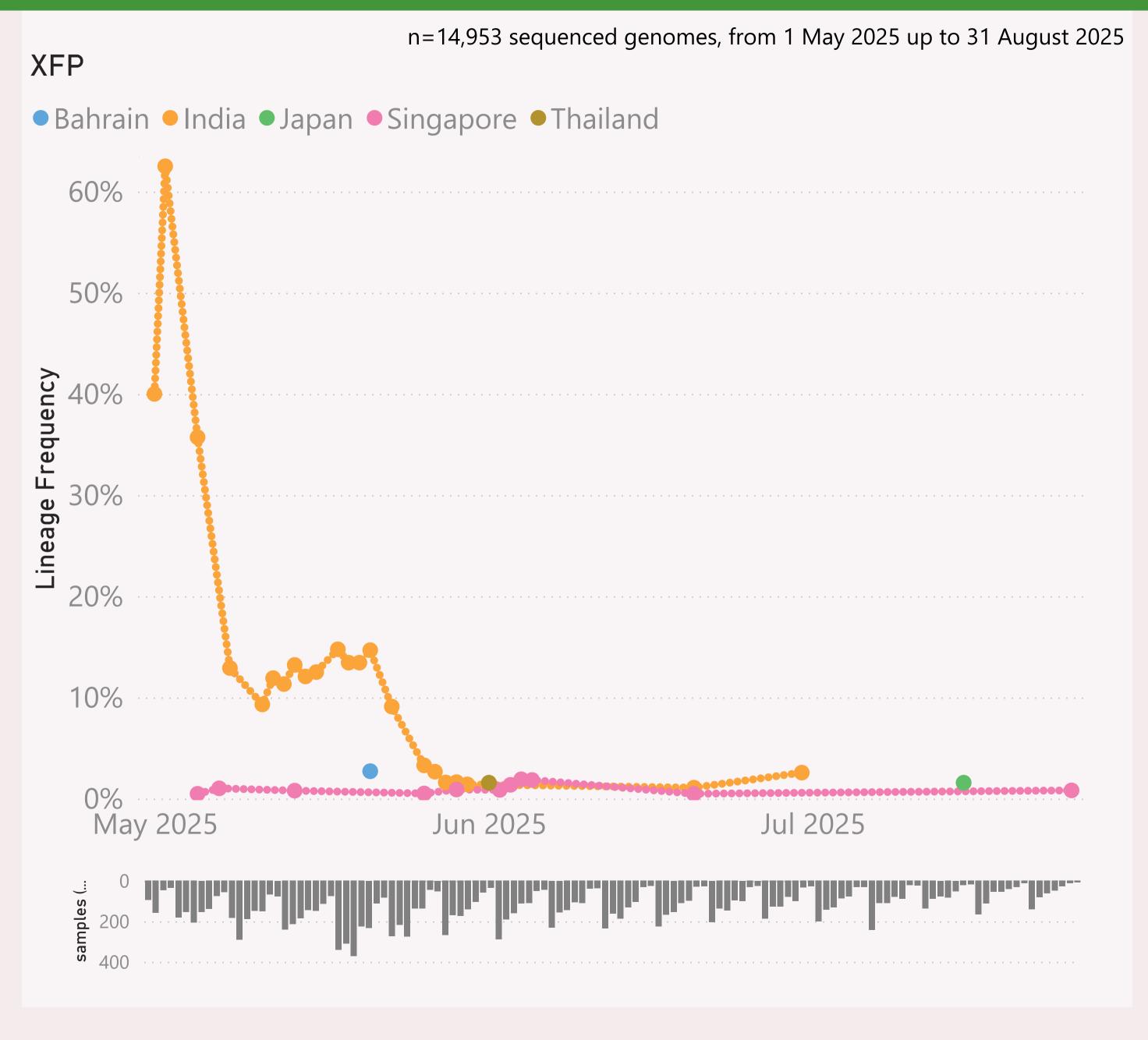
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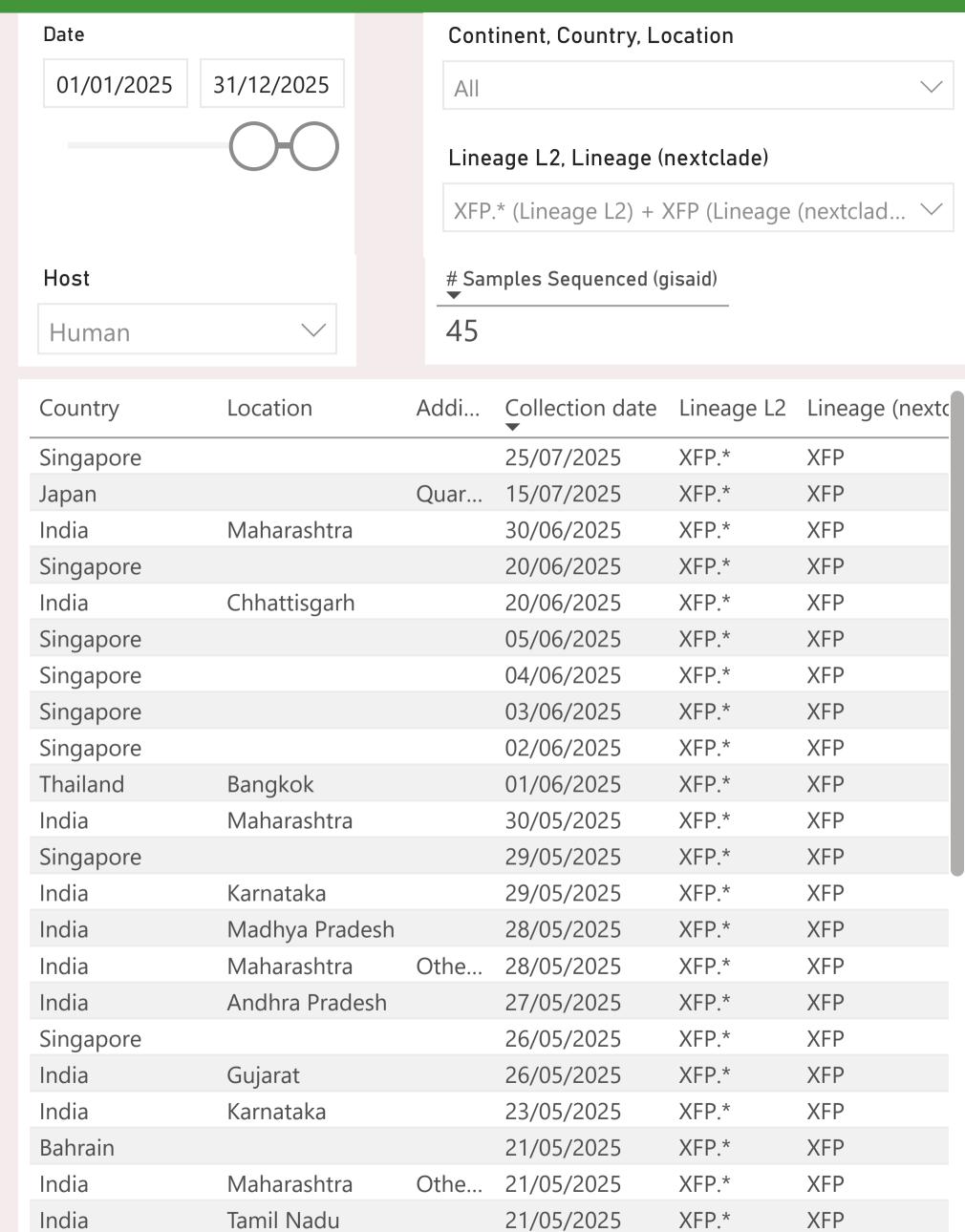
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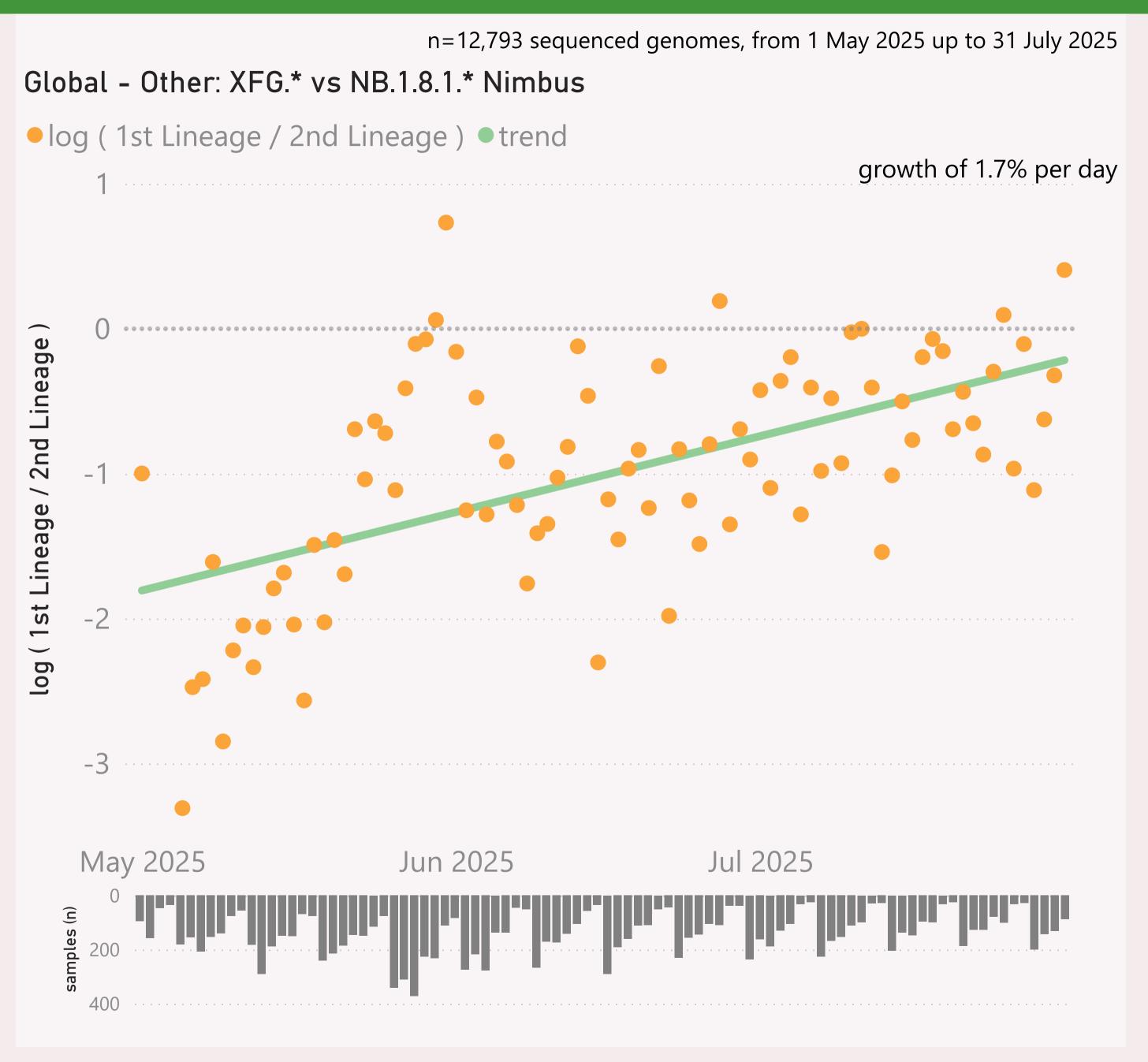
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India

Total

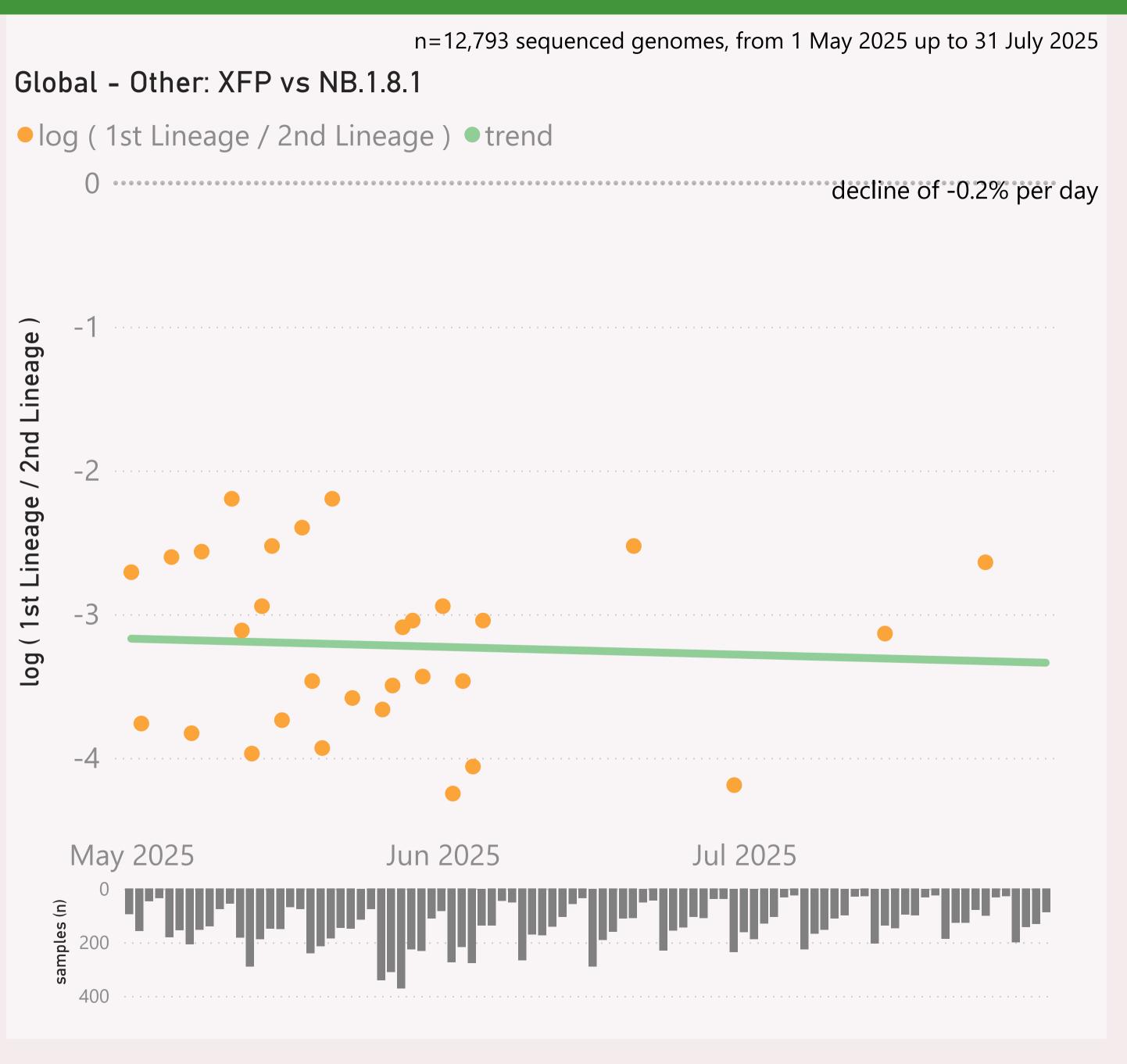


This page compares the relative frequency of 2 selected "Lineage L2" groups, over recent months. A challenging Lineage L2 is selected first, and compared to the incumbent.

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.

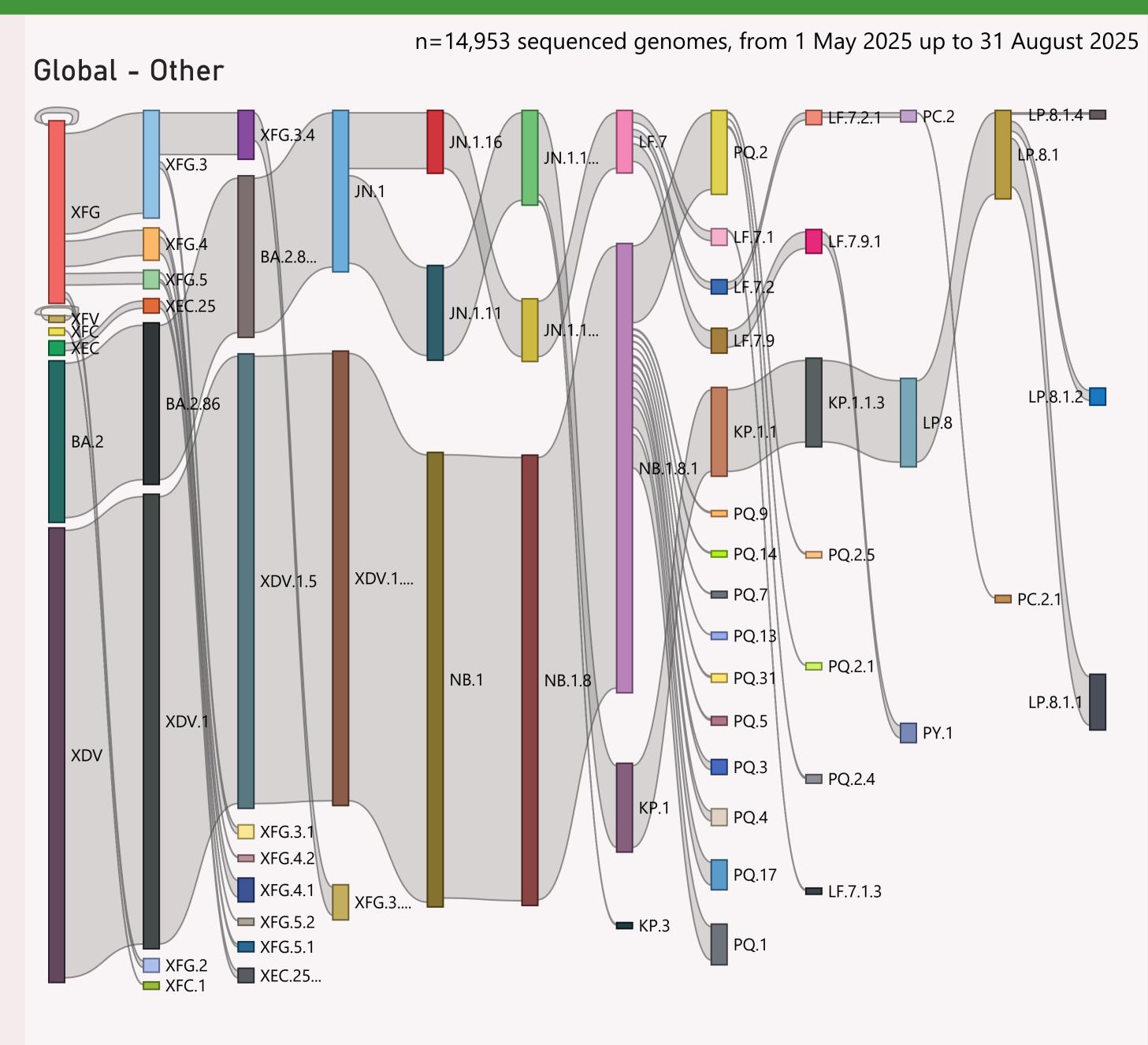


This page compares the relative frequency of 2 selected Lineages, over recent months. A challenging Lineage is selected first, and compared to the incumbent.

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
⊕ China	1,808	31/08/2025		06/09/2025	I
	862	28/08/2025	اللبلنيون .	06/09/2025	and the second of
⊕ Brazil	851	31/08/2025	وبالوالة أوارا والمستحدد والمستحدد	06/09/2025	
	760	31/08/2025	والموال المارية ووجود ورواد	06/09/2025	A Secretaria de la companya del companya del companya de la compan
	456	20/08/2025	an Ilda	25/08/2025	I i I i i i i i i i i i i i i i i i i i
	423	10/07/2025	attrate.	06/09/2025	- I .
⊕ Puerto Rico	324	30/08/2025	. Alatandi	06/09/2025	. 1
	307	12/08/2025	واللاهون	03/09/2025	11 .1 1
⊞ Taiwan	300	27/08/2025	.	06/09/2025	.1
	256	18/08/2025	and the second	06/09/2025	a tall
⊕ India	204	03/07/2025	hab.l.	17/08/2025	
	104	30/08/2025	and take of	06/09/2025	1. 1 1 1
	102	09/08/2025	the rate of a con-	06/09/2025	Here is
	85	18/08/2025	India	27/08/2025	Ι.
	54	29/07/2025	a. Hat.	06/09/2025	I
	48	25/08/2025	and taked	06/09/2025	
± Laos	47	29/07/2025	r II millioni	13/08/2025	and the
	46	01/08/2025	al. a	19/08/2025	- I - 1
⊞ Ecuador	46	27/08/2025	ata di ba	06/09/2025	
⊕ Qatar	46	12/07/2025	Maria	28/08/2025	
⊞ Egypt	38	31/08/2025	A section of	06/09/2025	
H Guam	38	18/08/2025	an albuma a calum	26/08/2025	n
⊕ Guatemala	36	08/08/2025	randa i	29/08/2025	
⊕ Brunei	33	08/07/2025	a alalat	06/09/2025	
⊕ Nepal	31	22/07/2025	. It	28/07/2025	
± Chile	30	10/08/2025	, Juli	05/09/2025	
⊞ Barbados	27	28/08/2025		06/09/2025	
	24	20/08/2025	i indi	06/09/2025	
Total	7,556	31/08/2025		06/09/2025	الأعديدات

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.