## **UK lineages summary report**

This report gives summaries of UK specific lineages sequenced by EDIN for week 2020-05-29. There are time lags due to batching, curation and analysis, the most recently sampled sequence is 2020-05-24. The analysis (eg time since last sample) is therefore undertaken from this date. 1049 sequences in the UK from the sequencing centre EDIN have been included in this analysis.

A few notes: the size of a lineage may be due to a low amount of transmission of this lineage, but it is likely also that it just hasn't been sampled as frequently, especially for newer lineages. It's also important to realise that these lineages are *estimates* of how we think the virus is spreading in the UK after being introduced from abroad, as the low evolutionary rate of the virus makes it difficult to separate lineages with certainty.

The minimum number of introductions is 6130 and the maximum is 9084

Sequences which were replicates or too error-prone were removed from this analysis.

365 are lineages which only contained five sequences or fewer, and so have been left out of visualisation in the interests of clarity

Furthermore, those sequences which haven't been sampled in the last month are not shown.

Of the 23 that remain: 10 are pending extinction, ie last seen three weeks ago. 5 lineages have gone quiet, ie haven't been seen this week. 4 lineages have reactivated. 4 lineages have been continuously circulating.

The following table contains information about lineages and the number of sequences the dataset, in reverse size order.

Each entry is the count of sequences from each lineage in each country, with the percentage of the total sequences from that lineage that this count represents.

The global lineages are correct as of the data release on 2020-05-19

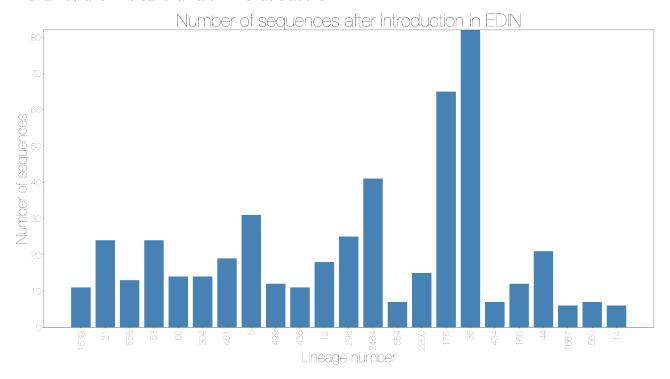
It is written to "summary\_files" as "lineage\_summary.tsv" for further use, and the full list of lineages is available in the same directory as "all\_lineages.csv"

Lineage			Total	Global	Time since last sample
name	Scotland	Date range	sequences	lineage	(days)
UK36	82	Mar-21,	82	B.1	20
	(100.0%)	May-04			
UK175	65	Mar-22,	65	B.1	20
	(100.0%)	May-04			
UK2464	41	Mar-20,	41	B.1.p11	12
	(100.0%)	May-12			
UK5	31	Mar-18,	31	B.1.1.1	6
	(100.0%)	May-18			
UK296	25	Apr-08,	25	B.1.5	11
	(100.0%)	May-13			
UK21	24	Mar-18,	24	B.1.40	1
	(100.0%)	May-23			
UK53	24	Apr-16,	24	B.1.1.4	3
	(100.0%)	May-21			
UK44	21	Mar-25,	21	В	23
	(100.0%)	May-01			
UK461	19	Apr-18,	19	B.1.5	5
	(100.0%)	May-19			

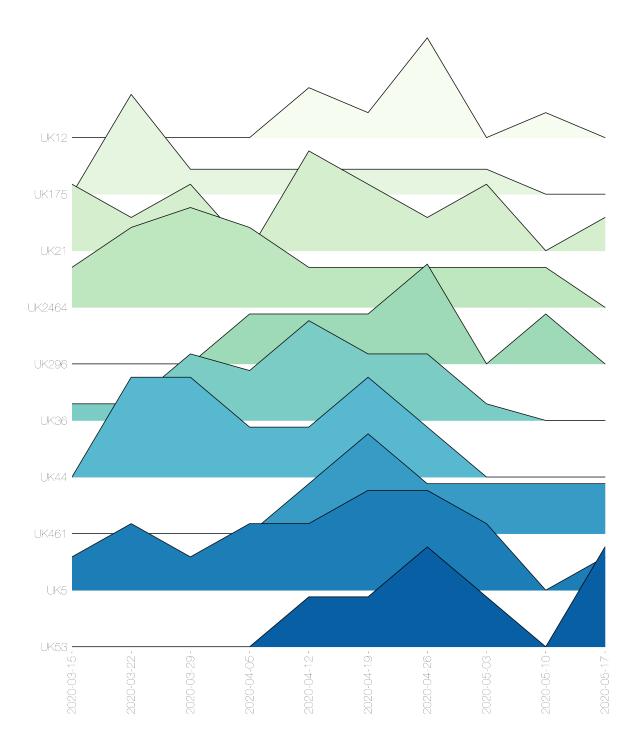
Lineage name	Scotland	Date range	Total sequences	Global lineage	Time since last sample (days)
			•		
UK150	19	Mar-21,	19	B.1.1.p12	32
1.114.4.0	(100.0%)	Apr-22	40	D.444	
UK12	18	Apr-12,	18	B.1.p11	11
	(100.0%)	May-13		5.505.5	
UK2200	15	Mar-25,	15	B.1.5.6, B.1.5	19
	(100.0%)	May-05			
UK66	14	Mar-28,	14	B.1.1.8	4
	(100.0%)	May-20			
UK43	14	Mar-26,	14	A.5	36
	(100.0%)	Apr-18			
UK156	14	Mar-18,	14	B.1.71	36
	(100.0%)	Apr-18			
UK304	14	Apr-16,	14	B.1.1.14	4
	(100.0%)	May-20			
UK558	13	Apr-24,	13	B.1.5	2
	(100.0%)	May-22			
UK160	12	Mar-31,	12	B.1.1	22
	(100.0%)	May-02			
UK499	12	Apr-24,	12	B.1.5	9
	(100.0%)	May-15			
UK562	11	Mar-27,	11	B.1	29
	(100.0%)	Apr-25			
UK436	11	Apr-13,	11	B.1.5	10
	(100.0%)	May-14			
UK1539	11	May-09,	11	B.1.5	0
	(100.0%)	May-24			
UK225	11	Mar-17,	11	B.2	49
	(100.0%)	Apr-05			
UK414	10	Apr-05,	10	B.1.5	32
	(100.0%)	Apr-22			
UK1548	9	Apr-13,	9	B.1.5.5, B.1.5	30
	(100.0%)	Apr-24			
UK560	7	Apr-15,	7	B.1.5	27
	(100.0%)	Apr-27			
UK554	7	Apr-23,	7	B.1.5	19
	(100.0%)	May-05			
UK434	7	Apr-20,	7	B.1.5	21
	(100.0%)	May-03	-		_,
UK14	6	Mar-21,	6	В	27
	(100.0%)	Apr-27	· ·	_	
UK555	6	Apr-13,	6	B.1.5	29
2	(100.0%)	Apr-25	O .	20	25
LIK1667	, ,		6	R 1 n9	26
J111001			0	5.1.90	20
I IK 133			e	R 1	29
OICIOO			0	D. 1	29
UK1667 UK133	6 (100.0%) 6 (100.0%)	Mar-31, Apr-28 Mar-22, Apr-25	6	B.1.p9 B.1	

These data is represented in the stacked bar chart below. Note that the number of sequences is likely to be due more to differing sampling efforts in different regions, rather than genuine differences in numbers of cases.

The raw data for this bar chart are in the table above.

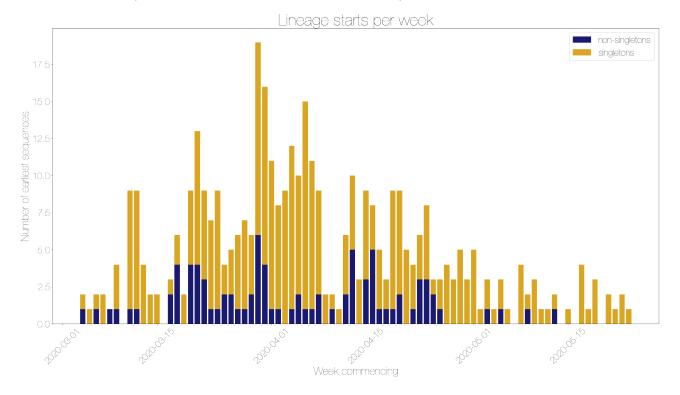


The relative growth and decline of the ten most sampled lineages in terms of number of counties they are present in is shown below. The raw data for the plot is shown below it, with each column representing a lineage, and the number of admin2 regions it is present in in each week.



Week commencing	UK36	UK175	UK2464	UK5	UK296	UK53	UK21	UK44	UK461	UK12
2020-03-15	1	0	2	1	0	0	2	0	0	0
2020-03-22	1	4	4	2	0	0	1	2	0	0
2020-03-29	4	1	5	1	0	0	2	2	0	0
2020-04-05	3	1	4	2	1	0	0	1	0	0
2020-04-12	6	1	2	2	1	1	3	1	1	2
2020-04-19	4	1	2	3	1	1	2	2	2	1
2020-04-26	4	1	2	3	2	2	1	1	1	4
2020-05-03	1	1	2	2	0	1	2	0	1	0
2020-05-10	0	0	2	0	1	0	0	0	1	1
2020-05-17	0	0	0	1	0	2	1	0	1	0

## The date of first sequence in the cluster is shown below for every cluster with date information.

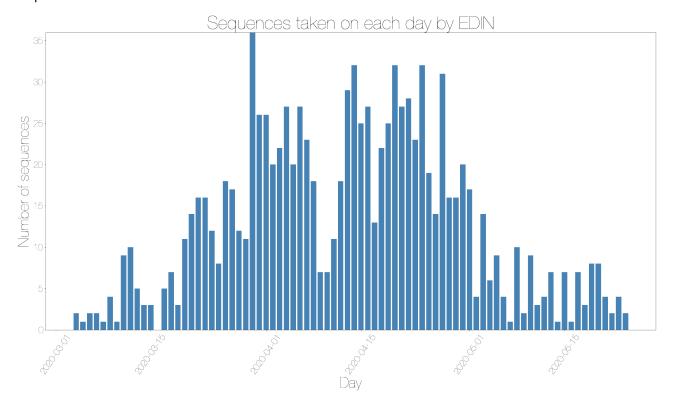


Day	Number of singleton starts	Number of non-singleton starts	Total
2020-03-04	1	1	2
2020-03-05	1	0	1
2020-03-06	1	1	2
2020-03-07	2	0	2
2020-03-08	0	1	1
2020-03-09	3	1	4
2020-03-11	8	1	9
2020-03-12	8	1	9
2020-03-13	4	0	4
2020-03-14	2	0	2
2020-03-15	2	0	2
2020-03-17	1	2	3
2020-03-18	2	4	6
2020-03-19	2	0	2
2020-03-20	5	4	9
2020-03-21	9	4	13
2020-03-22	6	3	9
2020-03-23	6	1	7
2020-03-24	8	1	9
2020-03-25	2	2	4
2020-03-26	3	2	5
2020-03-27	5	1	6
2020-03-28	6	1	7
2020-03-29	4	2	6
2020-03-30	13	6	19
2020-03-31	12	4	16

Day	Number of singleton starts	Number of non-singleton starts	Total
2020-04-01	10	1	11
2020-04-02	7	1	8
2020-04-03	9	0	9
2020-04-04	11	1	12
2020-04-05	8	2	10
2020-04-06	14	1	15
2020-04-07	10	1	11
2020-04-08	7	2	9
2020-04-09	2	0	2
2020-04-10	1	1	2
2020-04-11	1	0	1
2020-04-12	4	2	6
2020-04-13	5	5	10
2020-04-14	3	0	3
2020-04-15	6	3	9
2020-04-16	3	5	8
2020-04-17	4	1	5
2020-04-18	2	1	3
2020-04-19	8	1	9
2020-04-20	7	2	9
2020-04-21	5	0	5
2020-04-22	3	1	4
2020-04-23	3	3	6
2020-04-24	5	3	8
2020-04-25	1	2	3
2020-04-26	2	1	3
2020-04-27	4	0	4
2020-04-28	3	0	3
2020-04-29	5	0	5
2020-04-30	3	0	3
2020-05-01	5	0	5
2020-05-02	1	0	1
2020-05-03	2	1	3
2020-05-04	1	0	1
2020-05-05	2	1	3
2020-05-06	1	0	1
2020-05-08	4	0	4
2020-05-09	1	1	2
2020-05-10	3	0	3
2020-05-11	1	0	1
2020-05-12	1	0	1
2020-05-13	1	1	2
2020-05-15	1	0	1
2020-05-17	4	0	4
2020-05-18	1	0	1
2020-05-19	3	0	3
2020-05-21	2	0	2
2020-05-22 2020-05-23	1 2	0	1 2
ZUZU-UU-ZU	2	U	۷

Day	Number of singleton starts	Number of non-singleton starts	Total
2020-05-24	1	0	1

For comparison, here is a plot of the day that every sequence was taken, coloured by country. Note that sequences without dates were not included.

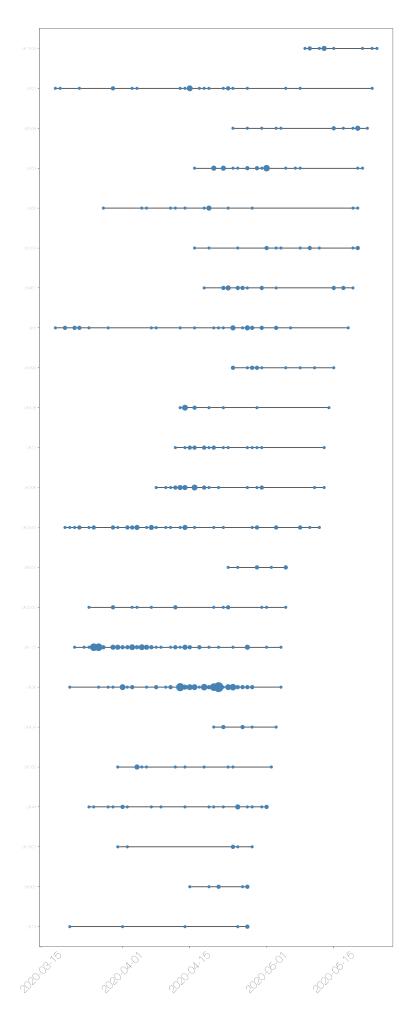


Day	Scotland
2020-03-04	2
2020-03-05	1
2020-03-06	2
2020-03-07	2
2020-03-08	1
2020-03-09	4
2020-03-10	1
2020-03-11	9
2020-03-12	10
2020-03-13	5
2020-03-14	3
2020-03-15	3
2020-03-17	5
2020-03-18	7
2020-03-19	3
2020-03-20	11
2020-03-21	14
2020-03-22	16
2020-03-23	16
2020-03-24	12

Day	Scotland
2020-03-25	8
2020-03-26	18
2020-03-27	17
2020-03-28	12
2020-03-29	11
2020-03-30	36
2020-03-31	26
2020-04-01	26
2020-04-02	20
2020-04-03	22
2020-04-04	27
2020-04-05	20
2020-04-06	27
2020-04-07	23
2020-04-08	18
2020-04-09	7
2020-04-09	7
2020-04-10	11
2020-04-11	18
2020-04-12	
	29
2020-04-14	32
2020-04-15	25
2020-04-16	27
2020-04-17	13
2020-04-18	22
2020-04-19	25
2020-04-20	32
2020-04-21	27
2020-04-22	28
2020-04-23	23
2020-04-24	32
2020-04-25	19
2020-04-26	14
2020-04-27	31
2020-04-28	16
2020-04-29	16
2020-04-30	20
2020-05-01	17
2020-05-02	4
2020-05-03	14
2020-05-04	6
2020-05-05	9
2020-05-06	4
2020-05-07	1
2020-05-08	10
2020-05-09	2
2020-05-10	9
2020-05-11	3
2020-05-12	4

Day	Scotland
2020-05-13	7
2020-05-14	1
2020-05-15	7
2020-05-16	1
2020-05-17	7
2020-05-18	3
2020-05-19	8
2020-05-20	8
2020-05-21	4
2020-05-22	2
2020-05-23	4
2020-05-24	2

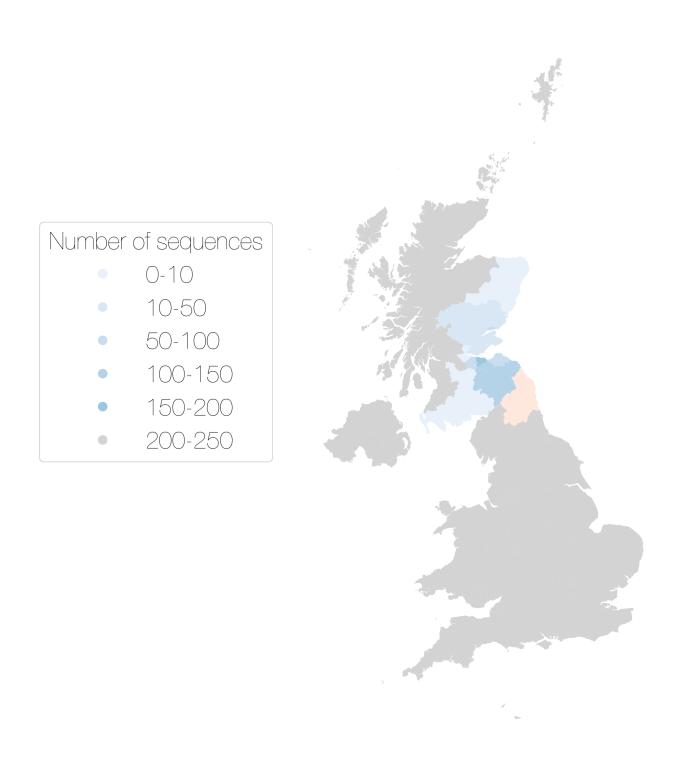
These lineages are shown on the timeline below. Each line represents the length of the cluster, from oldest to most recent sampling date. The dots are sized by the number of sequences taken on that date, and again are colour coded by country. The raw data has been written to a summary file.



The map below shows the number of sequences sampled in each admin2 region in the UK. The colour scale is the same for all four countries, but with different underlying base colours.

There are 70 sequences without enough geographical information to map from this centre.

COVID-19 sequences from each Admn2 region UK



Admin2	Country	Number of sequences	Sequence group
ABERDEEN	Scotland	1	1-10
ABERDEENSHIRE	Scotland	5	1-10
ANGUS	Scotland	13	10-50
CLACKMANNANSHIRE	Scotland	2	1-10
DUMFRIES AND GALLOWAY	Scotland	2	1-10
DUNDEE	Scotland	82	50-100
EAST LOTHIAN	Scotland	54	50-100
EDINBURGH	Scotland	402	400-500
FALKIRK	Scotland	4	1-10
FIFE	Scotland	42	10-50
GLASGOW	Scotland	1	1-10
MIDLOTHIAN	Scotland	127	100-150
NORTHUMBERLAND	England	1	1-10
PERTHSHIRE AND KINROSS	Scotland	18	10-50
SCOTTISH BORDERS	Scotland	128	100-150
SOUTH LANARKSHIRE	Scotland	4	1-10
WEST LOTHIAN	Scotland	92	50-100

Other results modules for UK lineage analysis can be added in here if required.