

UK lineages summary report

This report gives summaries of UK specific lineages sequenced by LIVE for week 2020-05-22. There are time lags due to batching, curation and analysis, the most recently sampled sequence is 2020-05-05. The analysis (eg time since last sample) is therefore undertaken from this date. 536 sequences in the UK from the sequencing centre LIVE have been included in this analysis. 177 lineages have been recorded, 123 of which only contain one sequence.

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 5185 and the maximum is 8310

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

154 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: 6 are pending extinction, ie last seen three weeks ago. 7 lineages have gone quiet, ie haven't been seen this week. 3 lineages have reactivated. 7 lineages have been continuously circulating.

The following table contains information about lineages and the number of sequences in each country in the UK for each lineage, 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

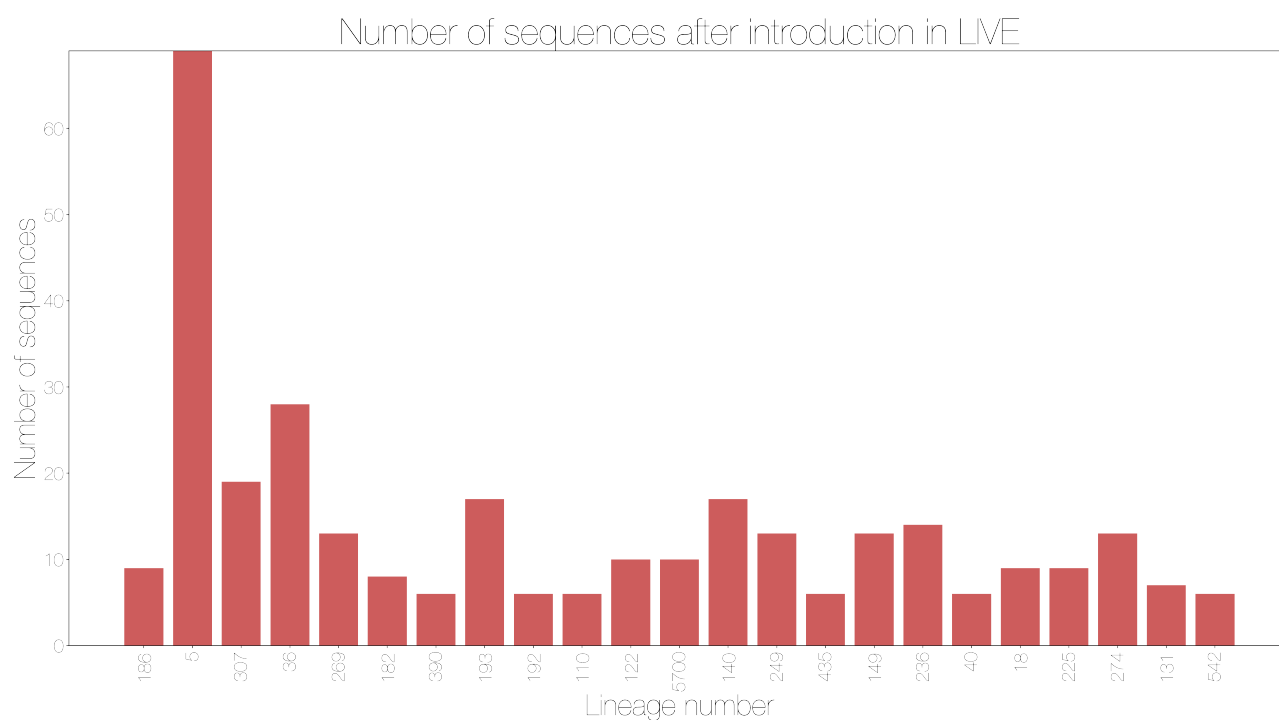
It is written to "summary_files" as "lineage_summary.tsv" for further use, and the full list of lineages is available in the same directory as "all_lineages.csv"

Lineage name	England	Date range	Total sequences	Global lineage	Time since last sample (days)
UK5	69 (100.0%)	Mar-21, May-05	69	B.1.1.1	0
UK36	28 (100.0%)	Mar-29, May-03	28	B.1	2
UK307	19 (100.0%)	Mar-28, May-04	19	B.1.1	1
UK193	17 (100.0%)	Apr-07, May-01	17	B.1.1	4
UK140	17 (100.0%)	Mar-23, Apr-25	17	B.1.1	10
UK236	14 (100.0%)	Mar-27, Apr-22	14	B.1.1	13
UK249	13 (100.0%)	Apr-01, Apr-25	13	B.1.1	10
UK274	13 (100.0%)	Mar-23, Apr-15	13	B.3	20

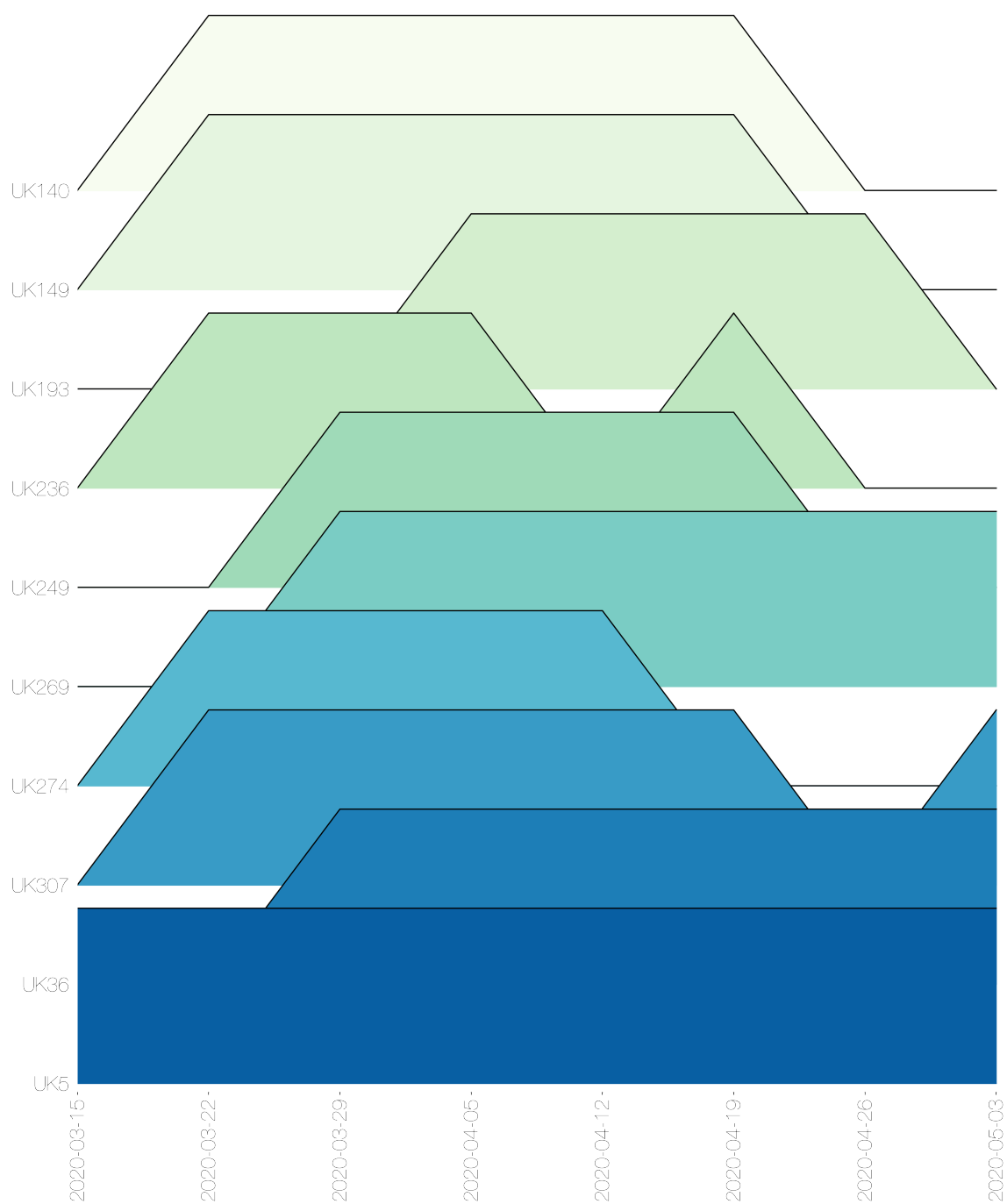
Lineage name	England	Date range	Total sequences	Global lineage	Time since last sample (days)
UK149	13 (100.0%)	Mar-23, Apr-22	13	B.1.1	13
UK269	13 (100.0%)	Apr-03, May-03	13	B.1.1	2
UK122	10 (100.0%)	Apr-16, Apr-28	10	B.1	7
UK5700	10 (100.0%)	Mar-24, Apr-27	10	B.1	8
UK186	9 (100.0%)	Apr-08, May-05	9	B	0
UK18	9 (100.0%)	Mar-27, Apr-19	9	B.1.1.7	16
UK225	9 (100.0%)	Mar-29, Apr-15	9	B.2, B.2.2	20
UK182	8 (100.0%)	Mar-29, May-02	8	B.1.1	3
UK131	7 (100.0%)	Mar-24, Apr-14	7	B.15	21
UK390	6 (100.0%)	Mar-29, May-01	6	B.1.5	4
UK110	6 (100.0%)	Mar-24, Apr-29	6	B.1	6
UK40	6 (100.0%)	Mar-31, Apr-20	6	B.16	15
UK542	6 (100.0%)	Apr-01, Apr-14	6	B.1	21
UK435	6 (100.0%)	Apr-03, Apr-23	6	B.1.5	12
UK192	6 (100.0%)	Apr-03, May-01	6	B.1.1	4

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.

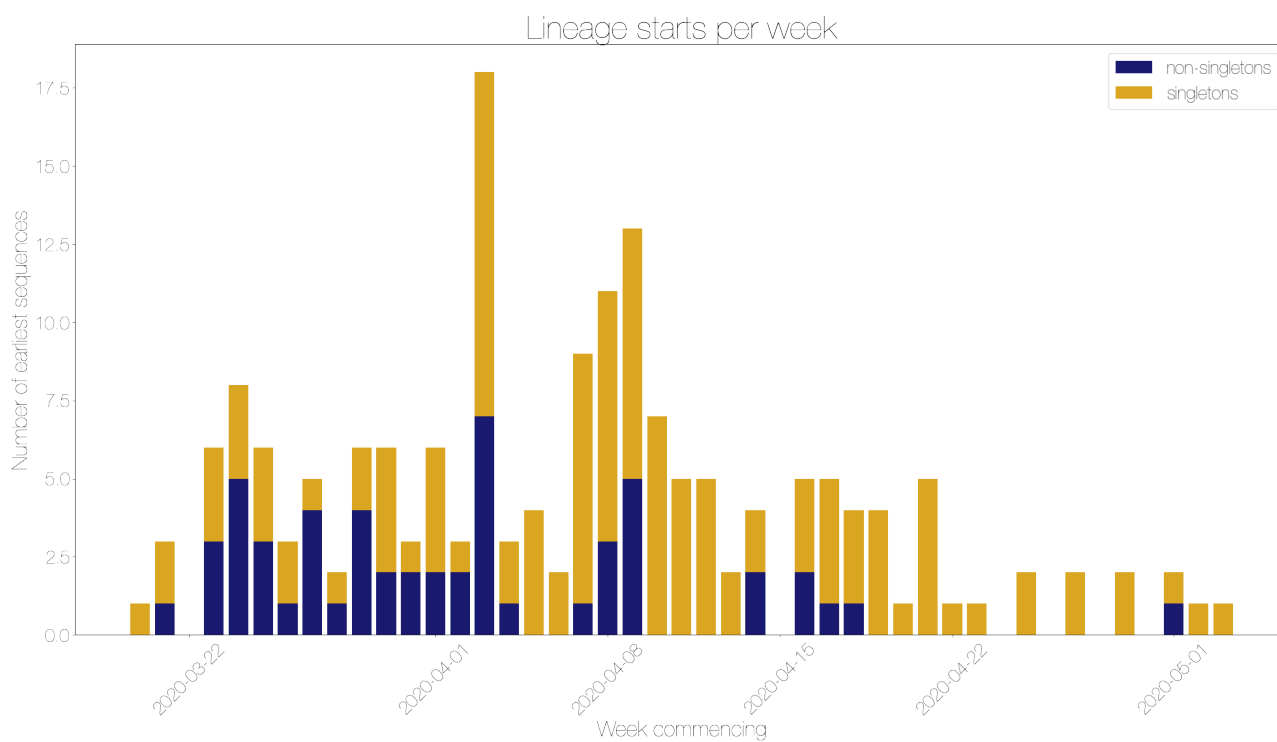


The relative growth and decline of the ten most sampled lineages in terms of number of counties they are present in is shown below. The raw data for the plot is shown below it, with each column representing a lineage, and the number of admin2 regions it is present in in each week.



Week commencing	UK5	UK36	UK307	UK140	UK193	UK236	UK274	UK149	UK249	UK269
2020-03-15	1	0	0	0	0	0	0	0	0	0
2020-03-22	1	0	1	1	0	1	1	1	0	0
2020-03-29	1	1	1	1	0	1	1	1	1	1
2020-04-05	1	1	1	1	1	1	1	1	1	1
2020-04-12	1	1	1	1	1	0	1	1	1	1
2020-04-19	1	1	1	1	1	1	0	1	1	1
2020-04-26	1	1	0	0	1	0	0	0	0	1
2020-05-03	1	1	1	0	0	0	0	0	0	1

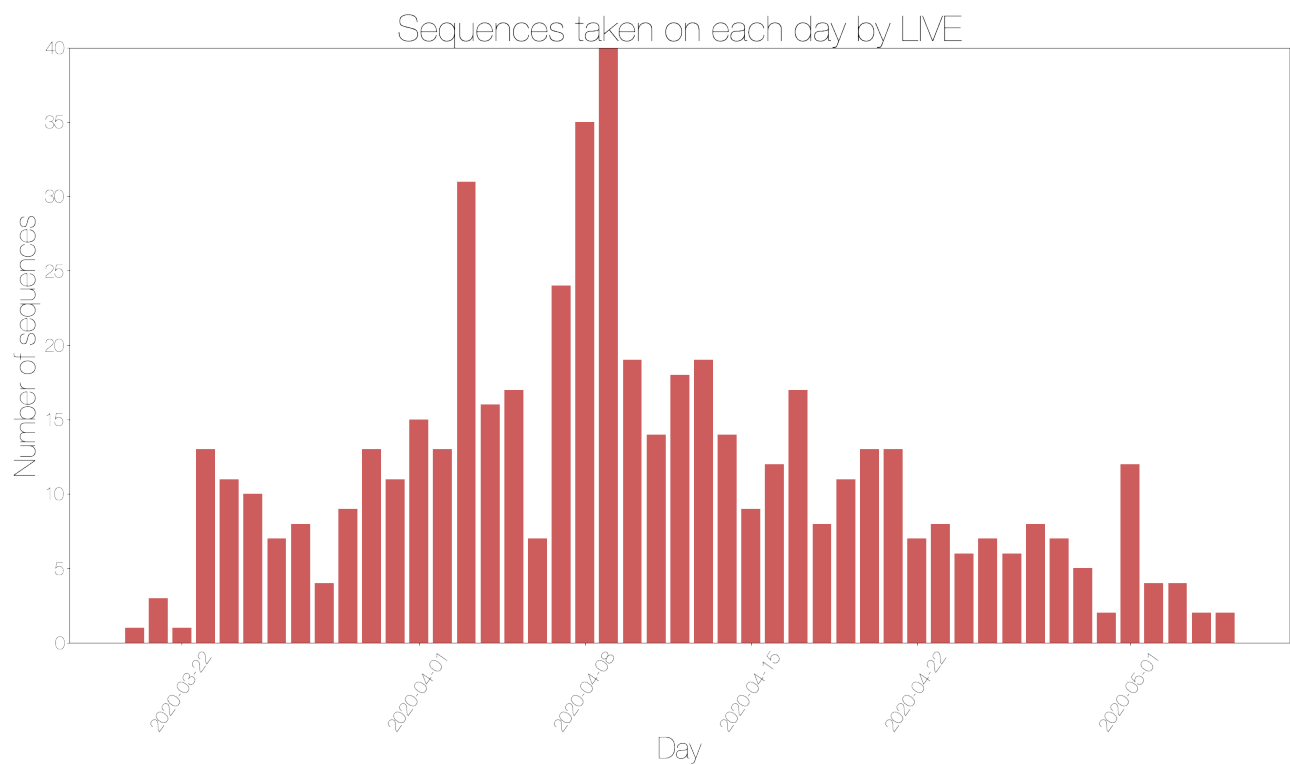
The date of first sequence in the cluster is shown below for every cluster with date information.



Day	Number of singleton starts	Number of non-singleton starts	Total
2020-03-20	1	0	1
2020-03-21	2	1	3
2020-03-23	3	3	6
2020-03-24	3	5	8
2020-03-25	3	3	6
2020-03-26	2	1	3
2020-03-27	1	4	5
2020-03-28	1	1	2
2020-03-29	2	4	6
2020-03-30	4	2	6
2020-03-31	1	2	3
2020-04-01	4	2	6
2020-04-02	1	2	3
2020-04-03	11	7	18
2020-04-04	2	1	3
2020-04-05	4	0	4
2020-04-06	2	0	2
2020-04-07	8	1	9
2020-04-08	8	3	11
2020-04-09	8	5	13
2020-04-10	7	0	7
2020-04-11	5	0	5
2020-04-12	5	0	5
2020-04-13	2	0	2
2020-04-14	2	2	4
2020-04-16	3	2	5
2020-04-17	4	1	5
2020-04-18	3	1	4

Day	Number of singleton starts	Number of non-singleton starts	Total
2020-04-19	4	0	4
2020-04-20	1	0	1
2020-04-21	5	0	5
2020-04-22	1	0	1
2020-04-23	1	0	1
2020-04-25	2	0	2
2020-04-27	2	0	2
2020-04-29	2	0	2
2020-05-01	1	1	2
2020-05-02	1	0	1
2020-05-03	1	0	1

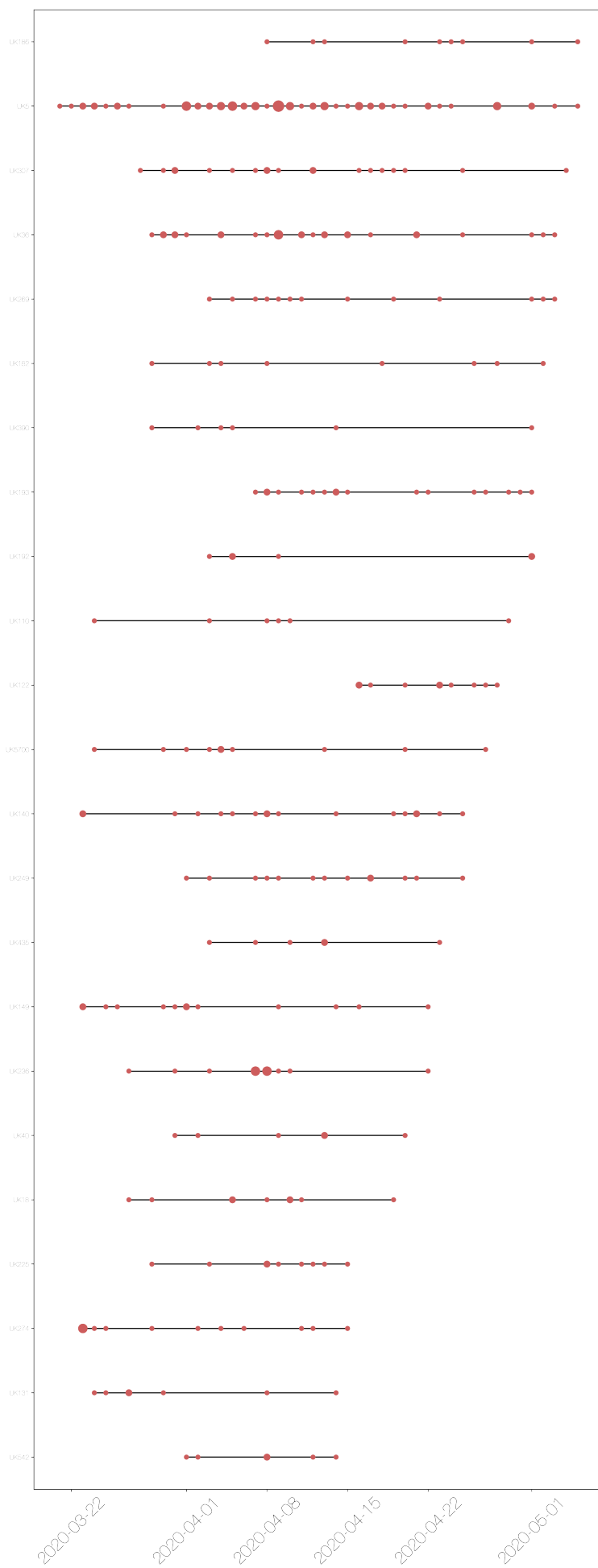
For comparison, here is a plot of the day that every sequence was taken, coloured by country. Note that sequences without dates were not included.



Day	England
2020-03-20	1
2020-03-21	3
2020-03-22	1
2020-03-23	13
2020-03-24	11
2020-03-25	10
2020-03-26	7
2020-03-27	8
2020-03-28	4
2020-03-29	9

Day	England
2020-03-30	13
2020-03-31	11
2020-04-01	15
2020-04-02	13
2020-04-03	31
2020-04-04	16
2020-04-05	17
2020-04-06	7
2020-04-07	24
2020-04-08	35
2020-04-09	40
2020-04-10	19
2020-04-11	14
2020-04-12	18
2020-04-13	19
2020-04-14	14
2020-04-15	9
2020-04-16	12
2020-04-17	17
2020-04-18	8
2020-04-19	11
2020-04-20	13
2020-04-21	13
2020-04-22	7
2020-04-23	8
2020-04-24	6
2020-04-25	7
2020-04-26	6
2020-04-27	8
2020-04-28	7
2020-04-29	5
2020-04-30	2
2020-05-01	12
2020-05-02	4
2020-05-03	4
2020-05-04	2
2020-05-05	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 no sequences with geographical information for this sequencing centre

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