Lineages report for GLAS

This report gives summaries of UK specific lineages sequenced by GLAS for week 2020-07-03. There are time lags due to batching, curation and analysis, the most recently sampled sequence is 2020-06-18. The analysis (eg time since last sample) is therefore undertaken from this date. 2069 sequences in the UK from the sequencing centre GLAS 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 129 and the maximum is 685

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

114 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 14 that remain: 7 are pending extinction, ie last seen three weeks ago. 4 lineages have gone quiet, ie haven't been seen this week. 1 has reactivated. 2 lineages have been continuously circulating.

The following table contains information about the ten largest lineages lineages and the number of sequences the dataset. Information about other lineages is found in the appendix, along with the raw data for all of the other figures.

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.

"Activity score" is calculated by taking the average gap between sampling for each lineage, and dividing it by the number of days since the lineage was last sampled. Therefore the higher the number, the more active the lineage is. If the score is above 1, then it has been sampled *more* recently than expected given its average gap size. We might interpret this as an increase in activity. If the score is below 1, it has been sampled *less* recently than expect given its average gap size, so we might interpret this as a decrease in activity.

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	Scotland	Date range	Total sequences	Global lineage	Time since last sample (days)	Activity score
UK5098	390	Mar-01,	390	B.1.p73, B.1	13	0.019
	(100.0%)	Jun-05				
UK36	266	Mar-20,	266	B.1	12	0.0245
	(100.0%)	Jun-06				
UK5	205	Feb-28,	205	B.1.1.p11, B.1.1.14,	3	0.1765
	(100.0%)	Jun-15		B.1.1.13, B.1.1.1, B.1.1		
UK40	153	Mar-13,	153	B.16, B	10	0.0572
	(100.0%)	Jun-08				
UK39	109	Mar-12,	109	A.2	20	0.0361
	(100.0%)	May-29				
UK5676	85	Mar-12,	85	B.2	22	0.0411
	(100.0%)	May-27				
UK2464	69	Mar-19,	69	B.1.p11	27	0.0349
	(100.0%)	May-22				
UK2913	68	Mar-19,	68	B.1.p11	30	0.0303
	(100.0%)	May-19				
UK668	66	Mar-22,	66	B.1	8	0.1538
	(100.0%)	Jun-10				
UK199	60	Mar-05,	60	B.1.p73, B.1.5, B.1	52	0.0173
	(100.0%)	Apr-27				

These data is represented in the figure one. 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.

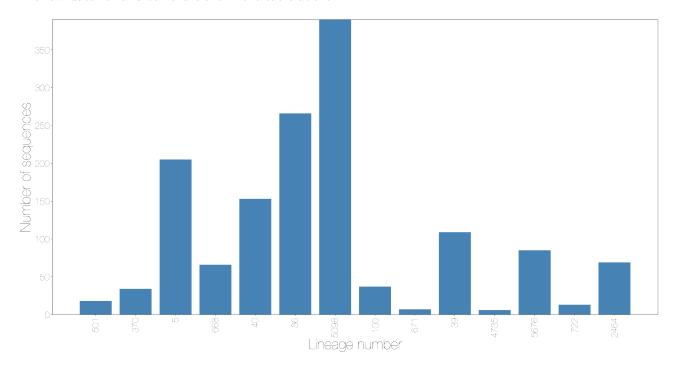


Figure 1: Number of sequences sampled in a lineage by country

Different sequencing centres have different delays in turn around from receipt of samples to submission of sequence data. This will affect all of the figures shown after this if lineages have geographical variation, as some regions have less up to date data.

The lag for this sequencing centre is 15 days

The relative growth and decline of the ten most sampled lineages in terms of number of counties they are present in is shown in figure three.

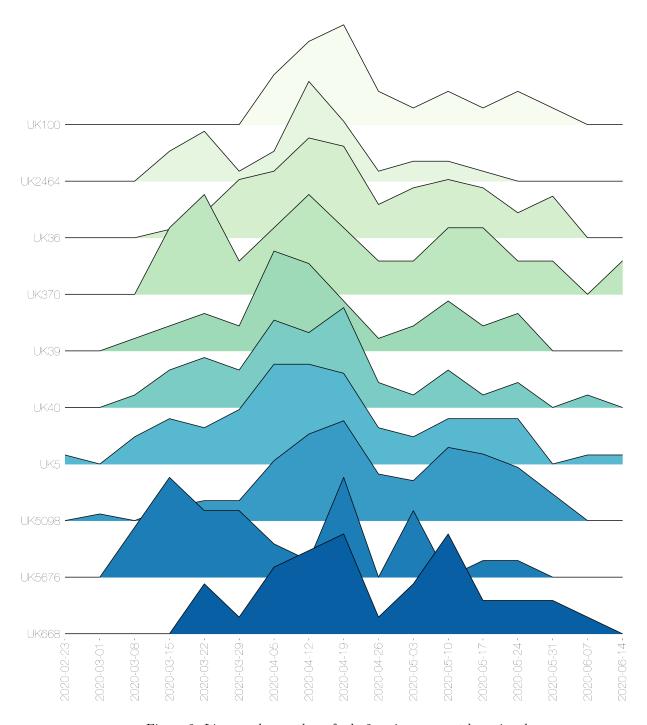


Figure 2: Lineages by number of adm2 regions present by epiweek

These lineages are shown on the timeline. 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.

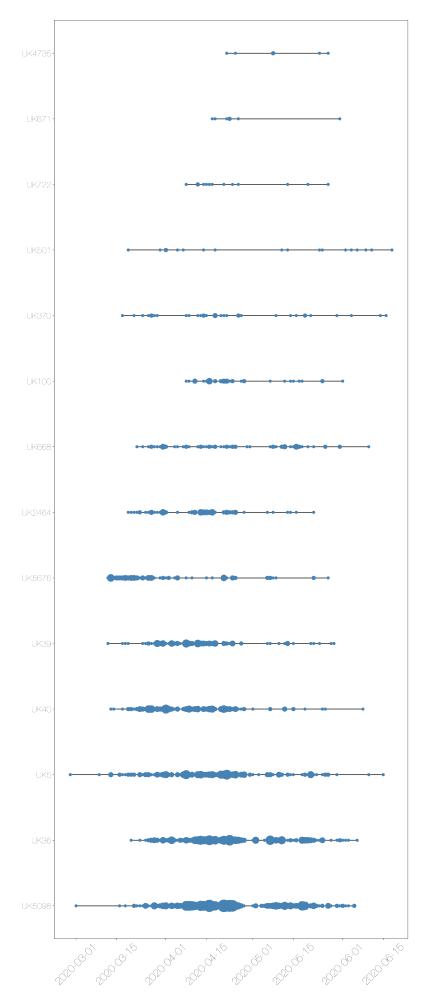


Figure 3: Timeline of lineages, sized by number of sequences from each country.

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

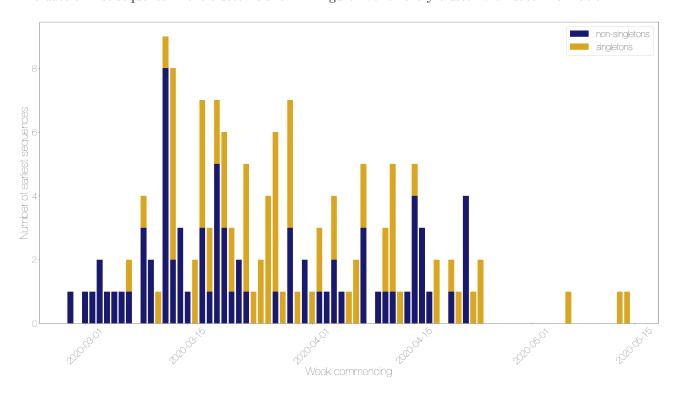


Figure 4: Lineage starts per week, split by singletons and non-singletons

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.

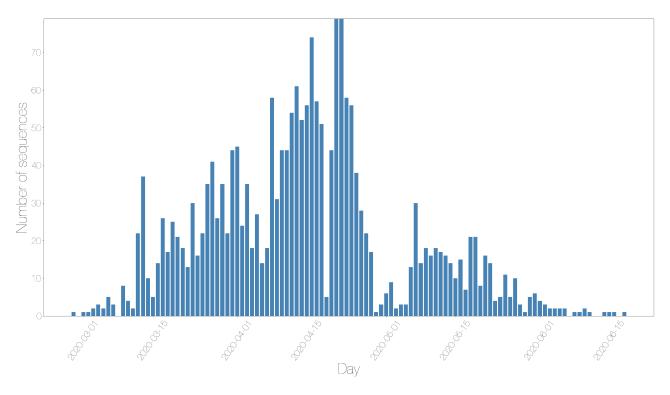


Figure 5: Sequences taken on each day by country

The map 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.

All sequences have been assigned clean adm2 data this week.

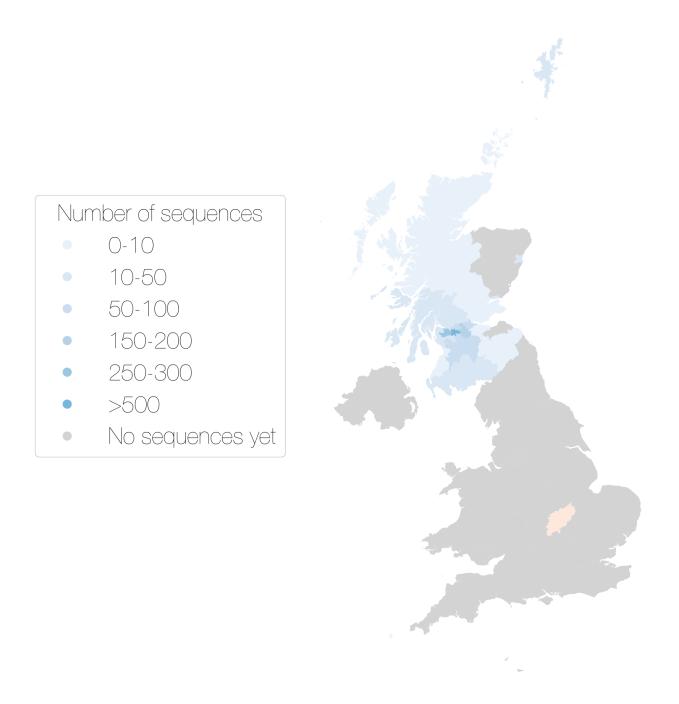


Figure 6: Map showing the number of sequences sampled by adm2 region

There are some sequences with locations that are not matched to real Admin2 regions, some manual curation required.

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

Appendix

Below are the raw data tables for each of the figures in the report.

 $\textbf{Table S1} \ \ \text{Description of all lineages that have been circulating in the last month, and have more than 5 sequences.}$

Lineage name	Scotland	Date range	Total sequences	Global lineage	Time since last sample (days)	Activity score
UK5098	390	Mar-01,	390	B.1.p73, B.1	13	0.019
	(100.0%)	Jun-05		- /		
UK36	266	Mar-20,	266	B.1	12	0.0245
TTT 7.2	(100.0%)	Jun-06	207	D. 1. 1. D. 1.1.		0.4505
UK5	205	Feb-28,	205	B.1.1.p11, B.1.1.14,	3	0.1765
UK40	(100.0%) 153	Jun-15 Mar-13,	153	B.1.1.13, B.1.1.1, B.1.1 B.16, B	10	0.0572
UIX40	(100.0%)	Jun-08	100	В.10, В	10	0.0572
UK39	109	Mar-12,	109	A.2	20	0.0361
	(100.0%)	May-29			-	
UK5676	85	Mar-12,	85	B.2	22	0.0411
	(100.0%)	May-27				
UK2464	69	Mar-19,	69	B.1.p11	27	0.0349
TTT/0010	(100.0%)	May-22	20	D 1 11	20	0.0000
UK2913	68 (100.0%)	Mar-19,	68	B.1.p11	30	0.0303
UK668	(100.0%)	May-19 Mar-22,	66	B.1	8	0.1538
011000	(100.0%)	Jun-10	00	Б.1	O	0.1000
UK199	60	Mar-05,	60	B.1.p73, B.1.5, B.1	52	0.0173
0 0 0	(100.0%)	Apr-27		,,	<u> </u>	0.02,0
UK42	42	Mar-10,	42	B.1.5, B.1.p73, B.1	31	0.0543
	(100.0%)	May-18				
UK100	37	Apr-08,	37	B.1.5, B.1	17	0.0882
	(100.0%)	Jun-01	2.4	D 1 1 10		4.0=00
UK370	34	Mar-17,	34	B.1.1.10	2	1.3788
UK87	(100.0%) 32	Jun-16 Mar-13,	32	B.1.70	55	0.0246
UKOI	(100.0%)	Apr-24	32	B.1.70	99	0.0240
UK15	27	Mar-12,	27	B.1.1	51	0.0354
0 0	(100.0%)	Apr-28			~-	0.000
UK107	24	Mar-09,	24	B.2.1	38	0.0721
	(100.0%)	May-11				
UK501	18	Mar-19,	18	B.1	0	active
TTT 7.4 .4	(100.0%)	Jun-18		T.	20	today
UK14	17	Mar-14,	17	В	69	0.0245
UK44	(100.0%) 16	Apr-10 Mar-17,	16	В	56	0.044
UIX44	(100.0%)	Apr-23	10	Б	50	0.044
UK502	16	Mar-06,	16	B.1.69	90	0.0104
	(100.0%)	Mar-20				
UK261	14	Mar-15,	14	A.3	69	0.029
	(100.0%)	Apr-10				
UK72	14	Mar-07,	14	В	77	0.026
TITZ 40	(100.0%)	Apr-02	1.4	A =	۳۵	0.0400
UK43	(100.0%)	Mar-23,	14	A.5	53	0.0493
UK722	(100.0%) 13	Apr-26 Apr-08,	13	B.1.5	22	0.1856
OIX122	(100.0%)	May-27	10	D.1.9	22	0.1000
UK58	13	Mar-12,	13	B.1	55	0.0652
	(100.0%)	Apr-24				
UK335	12	Apr-15,	12	B.1.1	28	0.1169
	(100.0%)	May-21				

Lineage name	Scotland	Date range	Total sequences	Global lineage	Time since last sample (days)	Activity score
UK2200	10	Mar-31,	10	B.1.5	45	0.084
	(100.0%)	May-04				
UK120	10	Mar-02,	10	В	63	0.0794
	(100.0%)	Apr-16				
UK167	9	Mar-12,	9	B.1	53	0.1061
	(100.0%)	Apr-26				
UK137	8	Mar-09,	8	B.1.1	81	0.0353
	(100.0%)	Mar-29	_			
UK671	7	Apr-17,	7	B.1.p73	18	0.4074
TTTT0-1	(100.0%)	May-31	_	D 4		
UK271	7	Apr-15,	7	B.1	53	0.0346
T.T.Z.O.F.O.F.	(100.0%)	Apr-26		D 1.1	40	0.000
UK2735	6	Apr-08,	6	B.1.1	62	0.029
T.T.Z.1.0.F	(100.0%)	Apr-17	0	D 1		0.0505
UK187	6	Apr-04,	6	B.1	55	0.0727
IIIZC01	(100.0%)	Apr-24	C	D 10	70	0.0460
UK601	6 (100.0%)	Mar-14,	6	B.10	78	0.0462
I IIZ 479E	(100.0%)	Apr-01	6	D 1 1	22	0.2100
UK4735	•	Apr-22,	О	B.1.1	22	0.3182
	(100.0%)	May-27				

 $\textbf{Table S2} \ \text{Raw data for figure two showing lags between the most recent sequence and current date for each sequencing centre}$

	Centre	Lag in days
0	GLAS	15

Table S3 Raw data for figure three showing the number of admin2 regions a lineage is present in over time

Week commencing	UK5098	UK36	UK5	UK40	UK39	UK5676	UK2464	UK668	UK100	UK370
2020-02-23	0	0	1	0	0	0	0	0	0	0
2020-03-01	1	0	0	0	0	0	0	0	0	0
2020-03-08	0	0	3	1	1	3	0	0	0	0
2020-03-15	2	1	5	3	2	6	3	0	0	2
2020-03-22	3	3	4	4	3	4	5	3	0	3
2020-03-29	3	7	6	3	2	4	1	1	0	1
2020-04-05	9	8	11	7	8	2	3	4	3	2
2020-04-12	13	12	11	6	7	1	10	5	5	3
2020-04-19	15	11	10	8	4	6	6	6	6	2
2020-04-26	7	4	4	2	1	0	1	1	2	1
2020-05-03	6	6	3	1	2	4	2	3	1	1
2020-05-10	11	7	5	3	4	0	2	6	2	2
2020-05-17	10	6	5	1	2	1	1	2	1	2
2020-05-24	8	3	5	2	3	1	0	2	2	1
2020-05-31	4	5	0	0	0	0	0	2	1	1
2020-06-07	0	0	1	1	0	0	0	1	0	0
2020-06-14	0	0	1	0	0	0	0	0	0	1

Table S4 is not appropriate for this report and so has been omitted.

 $\textbf{Table S5} \ \text{Raw data for figure five showing when lineages started per day, divided by singletons and non-singletons}$

Day	Number of singleton starts	Number of non-singleton starts	Total
2020-02-28	0	1	1
2020-03-01	0	1	1
2020-03-02	0	1	1
2020-03-03	0	2	2
2020-03-04	0	1	1
2020-03-05	0	1	1
2020-03-06	0	1	1
2020-03-07	1	1	2
2020-03-09	1	3	4
2020-03-10	0	2	2
2020-03-11	1	0	1
2020-03-12	1	8	9
2020-03-13	6	2	8
2020-03-14	0	3	3
2020-03-15	0	1	1
2020-03-16	2	0	2
2020-03-17	4	3	7
2020-03-18	$\frac{1}{2}$	1	3
2020-03-19	$\frac{1}{2}$	5	7
2020-03-20	3	$\ddot{3}$	6
2020-03-21	$\frac{\circ}{2}$	1	3
2020-03-22	0	2	2
2020-03-23	$\frac{3}{4}$	1	5
2020-03-24	1	0	1
2020-03-24	$\overset{1}{2}$	0	2
2020-03-26	$\frac{2}{4}$	0	4
2020-03-20	5	1	6
2020-03-21	1	0	1
2020-03-28	4	3	7
2020-03-29	1	0	1
2020-03-30	0	$\frac{0}{2}$	2
2020-03-31	1	0 = 0	1
2020-04-01	$\frac{1}{2}$	1	3
2020-04-02		1	3 1
2020-04-03	$0 \\ 2$	$\frac{1}{2}$	$\frac{1}{4}$
2020-04-04		1	1
2020-04-05	0 1	0	
	$\frac{1}{2}$		1
2020-04-07		0	2
2020-04-08	2	3	5
2020-04-10	0	1	1
2020-04-11	2	1	3
2020-04-12	4	1	5
2020-04-13	1	0	1
2020-04-14	0	1	1
2020-04-15	1	4	5
2020-04-16	0	3	3
2020-04-17	0	1	1
2020-04-18	2	0	2
2020-04-20	1	1	2
2020-04-21	1	0	1
2020-04-22	0	4	4
2020-04-23	1	0	1
2020-04-24	2	0	2
2020-05-06	1	0	1
2020-05-13	1	0	1
2020-05-14	1	0	1

 ${\bf Table~S6~{\rm Raw~data~for~figure~six~showing~the~number~of~sequences~taken~over~time.}$

Day	Scotland
2020-02-28	1
2020-02-28	1
2020-03-01	1
2020-03-03	2
2020-03-04	3
2020-03-05	2
2020-03-06	5
2020-03-07	3
2020-03-09	8
2020-03-10	4
2020-03-11	2
2020-03-12	22
2020-03-13	37
2020-03-14	10
2020-03-15	5
2020-03-16	14
2020-03-17	26
2020-03-18	17
2020-03-19	25
2020-03-20	21
2020-03-21	18
2020-03-22	13
2020-03-23	30
2020-03-24	16
2020-03-25	22
2020-03-26	35
2020-03-27	41
2020-03-28	26
2020-03-29	35
2020-03-30	22
2020-03-31	44
2020-04-01 2020-04-02	45 24
2020-04-02	$\frac{24}{35}$
2020-04-03	18
2020-04-04	$\frac{10}{27}$
2020-04-06	14
2020-04-07	18
2020-04-08	58
2020-04-09	31
2020-04-10	44
2020-04-11	44
2020-04-12	54
2020-04-13	61
2020-04-14	52
2020-04-15	56
2020-04-16	74
2020-04-17	57
2020-04-18	51
2020-04-19	5
2020-04-20	44
2020-04-21	79
2020-04-22	79
2020-04-23	58
2020-04-24	56
2020-04-25	38
2020-04-26	28

	C 41 1
Day	Scotland
2020-04-27	22
2020-04-28	17
2020-04-29	1
2020-04-30	3
2020-05-01	6
2020-05-02	9
2020-05-03	2
2020-05-04	3
2020-05-05	3
2020-05-06	13
2020-05-07	30
2020-05-08	14
2020-05-09	18
2020-05-10	16
2020-05-11	18
2020-05-12	17
2020-05-13	16
2020-05-14	14
2020-05-15	10
2020-05-16	15
2020-05-17	7
2020-05-18	21
2020-05-19	21
2020-05-20	8
2020-05-21	16
2020-05-22	14
2020-05-23	4
2020-05-24	5
2020-05-25	11
2020-05-26	5
2020-05-27	10
2020-05-28	3
2020-05-29	1
2020-05-30	5
2020-05-31	6
2020-06-01	4
2020-06-02	3
2020-06-03	2
2020-06-04	2
2020-06-05	2
2020-06-06	2
2020-06-08	1
2020-06-09	1
2020-06-10	2
2020-06-11	1
2020-06-14	1
2020-06-15	1
2020-06-16	1
2020-06-18	1

Table S7 Raw data for the figure seven with the number of sequences assigned to each admin2 region.

Admin2	Country	Number of sequences	Sequence group
ABERDEEN	Scotland	19	10-50
ARGYLL AND BUTE	Scotland	14	10-50
CLACKMANNANSHIRE	Scotland	1	1-10
DUMFRIES AND GALLOWAY	Scotland	42	10-50
DUNDEE	Scotland	11	10-50
EAST AYRSHIRE	Scotland	59	50-100
EAST DUNBARTONSHIRE	Scotland	73	50-100
EAST RENFREWSHIRE	Scotland	40	10-50
EDINBURGH	Scotland	9	1-10
EILEAN SIAR	Scotland	2	1-10
FALKIRK	Scotland	70	50-100
FIFE	Scotland	1	1-10
GLASGOW	Scotland	1006	>500
HIGHLAND	Scotland	9	1-10
INVERCLYDE	Scotland	42	10-50
NORTH AYRSHIRE	Scotland	18	10-50
NORTH LANARKSHIRE	Scotland	197	150-200
NORTHAMPTONSHIRE	England	1	1-10
ORKNEY ISLANDS	Scotland	1	1-10
PERTHSHIRE AND KINROSS	Scotland	2	1-10
RENFREWSHIRE	Scotland	286	250-300
SCOTTISH BORDERS	Scotland	1	1-10
SHETLAND ISLANDS	Scotland	14	10-50
SOUTH AYRSHIRE	Scotland	7	1-10
SOUTH LANARKSHIRE	Scotland	66	50-100
STIRLING	Scotland	16	10-50
WEST DUNBARTONSHIRE	Scotland	49	10-50
WEST LOTHIAN	Scotland	1	1-10