Lineages report for Scotland

This report gives summaries of lineages sampled in Scotland for week 2020-06-19. There are time lags due to batching, curation and analysis, the most recently sampled sequence is 2020-06-08. The analysis (eg time since last sample) is therefore undertaken from this date. 3232 sequences from Scotland have been included in this analysis. 225 lineages have been recorded, 114 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 132 and the maximum is 1169

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

195 are lineages which were sampled less than five times in Scotland, 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 30 that remain: 10 are pending extinction, ie last seen three weeks ago. 7 lineages have gone quiet, ie haven't been seen this week. 5 lineages have reactivated. 8 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	Date range	Number of sequences	Global lineage	Time since last sample (days)	Activity score
			Global inleage	Sample (days)	50010
UK36	Mar-20,	353	B.1	5	0.0371
	Jun-03				
UK5098	Mar-16,	332	B.1.p73	12	0.0178
	May-27				
UK5	Feb-28,	302	B.1.1.p12, B.1.1.13, B.1.1.1,	0	active
	Jun-08		B.1.1.14, B.1.1, B.1.1.p11		today
UK42	Mar-01,	237	B.1.p73, B.1.71, B.1.p11, B.1,	4	0.0233
	Jun-04		B.1.5		
UK109	Mar-12,	229	B.1.5.5, B.1.5	0	active
	Jun-08				today
UK40	Mar-13,	164	B.16, B	13	$0.03\overline{3}3$
	May-26		,		
UK199	Mar-05,	151	B.1, B.1.p73, B.1.5	0	active
	Jun-08		, ,		today
UK5676	Mar-12,	137	B.2	17	0.011
	May-22				
UK2464	Mar-19,	125	B.1.p11	1	0.1848
	Jun-07		r	_	
UK39	Mar-12,	107	A.2	15	0.0451
	May-24	201		10	

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.

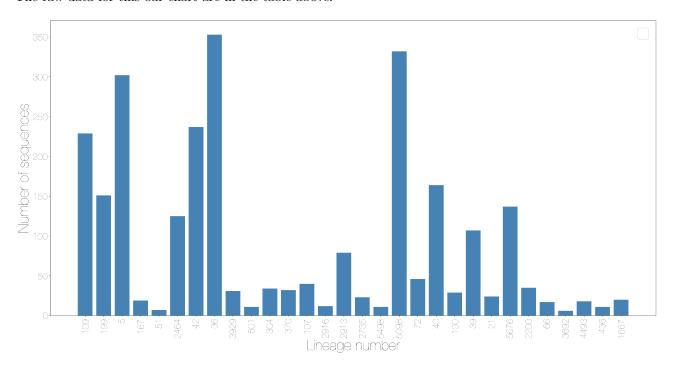


Figure 1: Number of sequences sampled in a lineage by country

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.

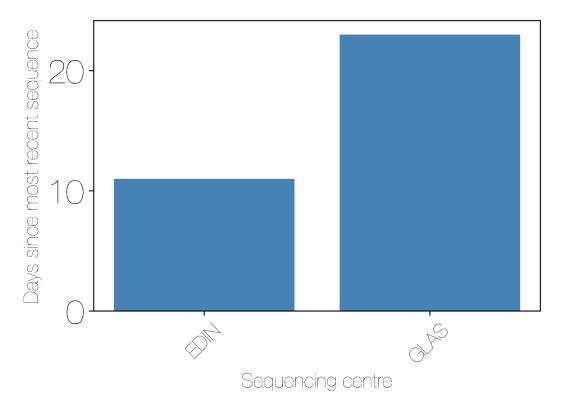


Figure 2: Lag since the most recent sequence from each sequencing centre to most current date

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.

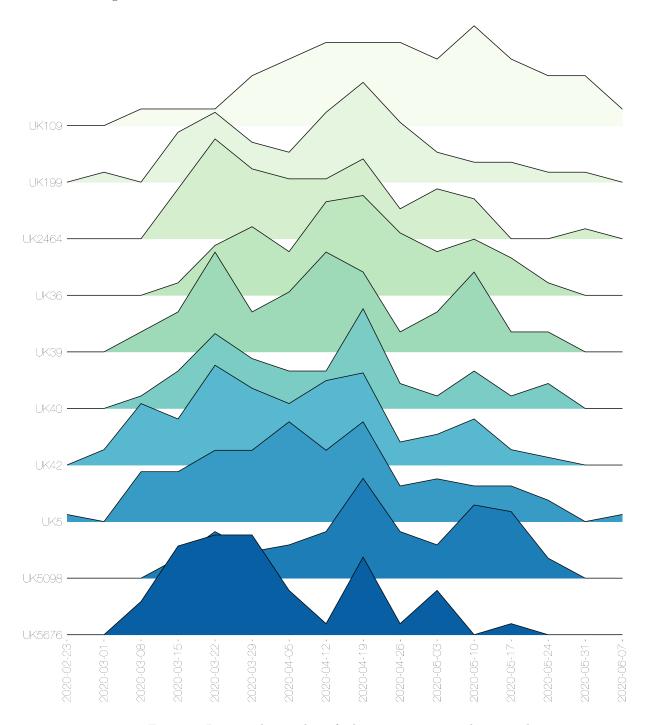


Figure 3: Lineages by number of adm2 regions present by epiweek

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.

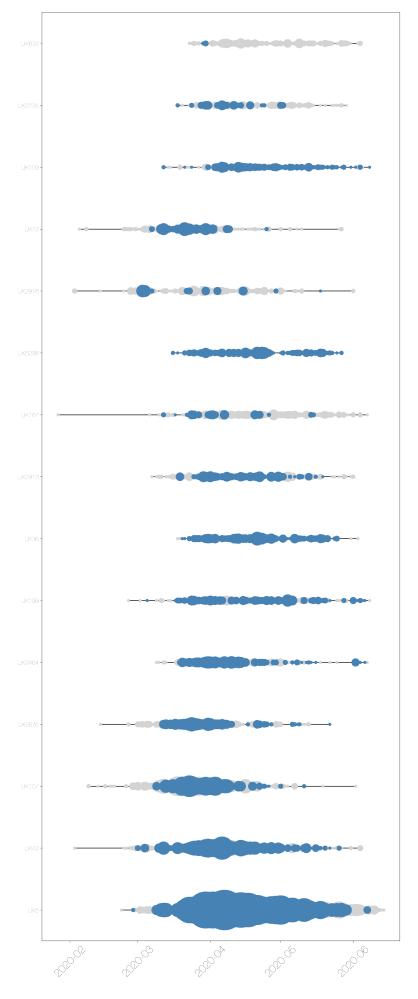


Figure 4: Timeline of lineages, sized by number of sequences from each country.

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

NB the lineage may have started anywhere in the UK, but has been recorded at least once in Scotland

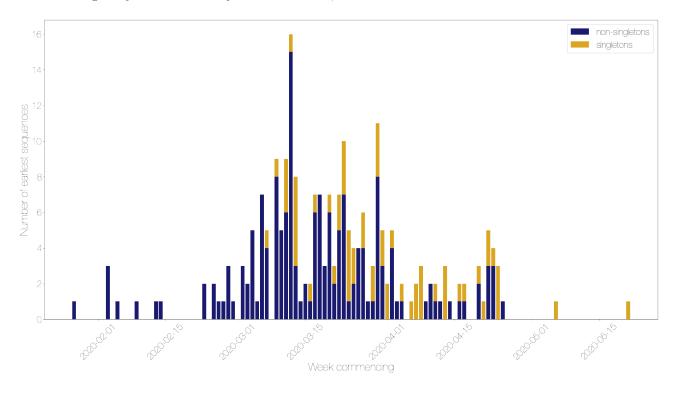


Figure 5: Lineage starts per week, split by singletons and non-singletons

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.

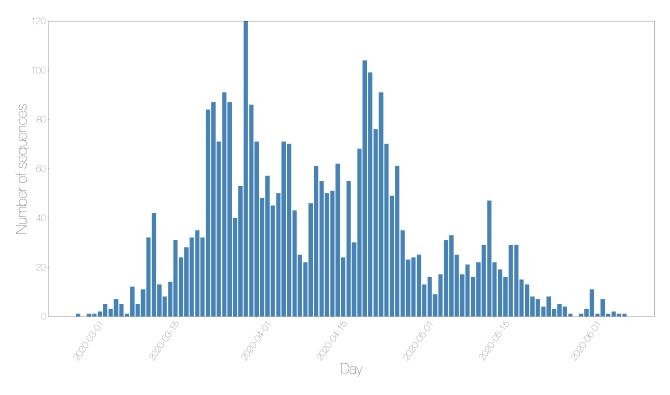


Figure 6: Sequences taken on each day by country

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.

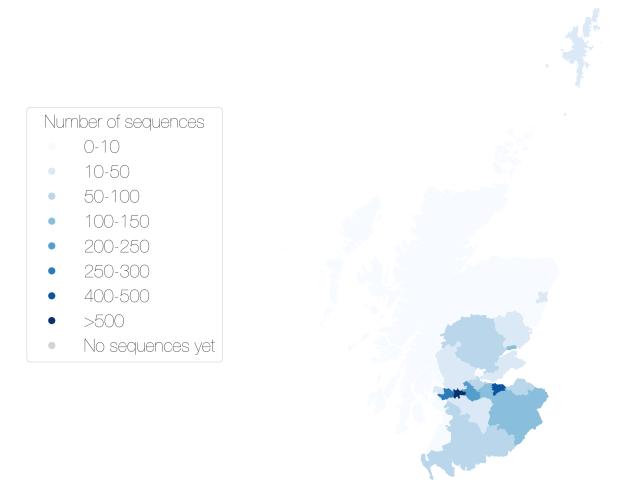


Figure 7: Map showing the number of sequences sampled by adm2 region

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	Date	Number of sequences	Global lineage	Time since last sample (days)	Activity score
name	range				
UK36	Mar-20,	353	B.1	5	0.0371
UK5098	Jun-03 Mar-16,	332	B.1.p73	12	0.0178
0110050	May-27	002	D.1.pro	12	0.0110
UK5	Feb-28,	302	B.1.1.p12, B.1.1.13, B.1.1.1,	0	active
	Jun-08		B.1.1.14, B.1.1, B.1.1.p11		today
UK42	Mar-01,	237	B.1.p73, B.1.71, B.1.p11, B.1,	4	0.0233
	Jun-04		B.1.5	_	
UK109	Mar-12,	229	B.1.5.5, B.1.5	0	active
UK40	Jun-08 Mar-13,	164	B.16, B	13	$\begin{array}{c} \text{today} \\ 0.0333 \end{array}$
UK40	May-26	164	В.10, В	10	0.0555
UK199	Mar-05,	151	B.1, B.1.p73, B.1.5	0	active
0 === 0 0	Jun-08		_ · - , _ · - · - · · · · · · · · · · · · · · ·	· ·	today
UK5676	Mar-12,	137	B.2	17	$0.01\overset{\circ}{1}$
	May-22				
UK2464	Mar-19,	125	B.1.p11	1	0.1848
TTT - 0.0	Jun-07				
UK39	Mar-12,	107	A.2	15	0.0451
UK2913	May-24	70	D 1 n11	7	0.0297
UK2915	Mar-19, Jun-01	79	B.1.p11	7	0.0297
UK72	Mar-07,	46	В	12	0.029
	May-27		_		0.000
UK107	Mar-09,	40	B.2.1	6	0.0148
	Jun-02				
UK198	Mar-17,	37	B.1, B.1.5	32	0.027
T.T.T. 2.2.2.2	May-07	25	Dara Dar	10	0.0450
UK2200	Mar-23,	35	B.1.5.6, B.1.5	19	0.0459
UK304	May-20 Apr-16,	34	B.1.1.14	6	0.2374
011304	Jun-02	94	D.1.1.14	O	0.2014
UK370	Mar-17,	32	B.1.1.10	6	0.0843
	$\operatorname{Jun-02}^{'}$				
UK3929	Mar-06,	31	B.1.1.4, B.1.1	5	0.0848
	Jun-03				
UK100	Mar-30,	29	B.1, B.1.5	14	0.1429
T1171 F	May-25	00	D 1 1	99	0.0105
UK15	Mar-07, May-06	28	B.1.1	33	0.0125
UK43	Mar-23,	28	A.5	43	0.0374
01110	Apr-26	20		10	0.0011
UK44	Mar-17,	28	В	50	0.0236
	Apr-19				
UK14	Mar-14,	28	В	42	0.0378
111701	Apr-27	2.4	D 1 40	4.0	0.1500
UK21	Mar-18,	24	B.1.40	16	0.1793
UK87	May-23 Mar-13,	23	B.1.70	45	0.0424
01701	Apr-24	23	D.1.10	40	0.0424
	-	23	B.1.1	10	0.0293
UK2735	Mar-18,	23	D.1.1	10	0.0295

Lineage	Date	Number of		Time since last	Activity
name	range	sequences	Global lineage	sample (days)	score
UK1667	Mar-31,	20	B.1.p9	25	0.0783
	May-14				
UK167	Mar-12,	19	B.1	1	0.3376
	Jun-07				
UK4493	Apr-23,	18	B.1	20	0.0765
	May-19				
UK502	Mar-06,	18	B.1.69	70	0.0202
	Mar-30				
UK66	Mar-28,	17	B.1.1.8	19	0.0304
	May-20				
UK120	Mar-02,	14	В	34	0.0416
	May-05				
UK261	Mar-15,	12	A.3	61	0.0358
	Apr-08				
UK2916	Mar-03,	12	B.1	7	0.0525
	Jun-01				
UK436	Apr-13,	11	B.1.5	25	0.1636
	May-14				
UK137	Mar-09,	11	B.1.1	69	0.0266
	Mar-31				
UK501	Mar-19,	11	B.1	6	0.1991
TTTT= 100	Jun-02		D 0 D		
UK5498	Mar-12,	11	B.2, B	11	0.0909
TTT 2004	May-28		P. 10	25	0.04.00
UK601	Mar-14,	8	B.10	35	0.0168
TITZ# 40	May-04		D 0.1	70	0.000
UK548	Mar-14,	8	B.2.1	70	0.0327
TITZEOA	Mar-30	0	D	90	0.0414
UK594	Apr-20,	8	В	38	0.0414
T.II.Z.1.0. D	May-01	7	D 1	20	0.0010
UK187	Mar-23,	7	B.1	39	0.0212
11170 7 1	Apr-30	7	D 1	49	0.0707
UK271	Apr-15, Apr-26	7	B.1	43	0.0797
THZEO	•	7	D 1	60	0.0290
UK58	Mar-12, Apr-09	7	B.1	60	0.0389
UK151	Mar-23,	7	B.1	45	0.1185
017191	Apr-24	1	D.1	40	0.1165
UK133	Mar-22,	7	B.1	44	0.0966
017133	Apr-25	'	Б.1	44	0.0900
UK51	Mar-26,	7	B.1.36	1	0.9873
01101	Jun-07	•	2.1.00	1	0.0010
UK3692	Mar-12,	6	B.1.1	20	0.1268
0110002	May-19	3		20	0.1200
UK240	Mar-22,	6	B.2	31	0.0259
	May-08	· ·	•	01	00
	<i>J</i>				

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

_	Centre	Lag in days
	Contre	Lag III days
0	EDIN	11
1	GLAS	23

 $\textbf{Table S3} \ \text{Raw data for figure three showing the number of admin2 regions a lineage is present in over time}$

Week commencing	UK36	UK5098	UK5	UK42	UK109	UK40	UK199	UK5676	UK2464	UK39
2020-02-23	0	0	1	0	0	0	0	0	0	0
2020-03-01	0	0	0	2	0	0	1	0	0	0
2020-03-08	0	0	7	8	1	1	0	3	0	1
2020-03-15	2	3	7	6	1	3	5	8	5	2
2020-03-22	8	7	10	13	1	6	7	9	10	5
2020-03-29	11	4	10	10	3	4	4	9	7	2
2020-04-05	7	5	14	8	4	3	3	4	6	3
2020-04-12	15	7	10	11	5	3	7	1	6	5
2020-04-19	16	15	14	12	5	8	10	7	8	4
2020-04-26	10	7	5	3	5	2	6	1	3	1
2020-05-03	7	5	6	4	4	1	3	4	5	2
2020-05-10	9	11	5	6	6	3	2	0	4	4
2020-05-17	6	10	5	2	4	1	2	1	0	1
2020-05-24	2	3	3	1	3	2	1	0	0	1
2020-05-31	0	0	0	0	3	0	1	0	1	0
2020-06-07	0	0	1	0	1	0	0	0	0	0

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

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

	NT 1 0 1 1		
Day	Number of singleton starts	Number of non-singleton starts	Total
2020-01-27	0	1	1
2020-02-03	0	3	3
2020-02-05	0	1	1
2020-02-09	0	1	1
2020-02-13	0	1	1
2020-02-14	0	1	1
2020-02-23	0	2	2
2020-02-25	0	2	2
2020-02-26	0	1	1
2020-02-27	0	1	1
2020-02-28	0	3	3
2020-02-29	0	1	1
2020-03-02	0	3	3
2020-03-03	0	2	2
2020-03-04	0	5	5
2020-03-05	0	1	1
2020-03-06	0	7	7
2020-03-07	1	4	5
2020-03-09	1	8	9
2020-03-10	0	5	5
2020-03-11	3	6	9
2020-03-12	1	15	16
2020-03-13	5	3	8
2020-03-14	0	1	1
2020-03-15	0	2	2
2020-03-16	1	1	2
2020-03-17	1	6	7
2020-03-18	0	7	7
2020-03-19	0	3	3
2020-03-20	1	6	7
2020-03-21	1	2	3
2020-03-22	2	5	7
2020-03-23	3	7	10
2020-03-24	4	1	5
2020-03-25	2	2	4
2020-03-26	0	4	4
2020-03-27	2	4	6
2020-03-28	0	1	1
2020-03-29	2	1	3
2020-03-30	3	8	11
2020-03-31	2	3	5
2020-04-01	2	0	2
2020-04-02	1	4	5
2020-04-03	0	1	1
2020-04-04	1	1	2
2020-04-06	1	0	1
2020-04-07	2	0	2
2020-04-08	3	0	3
2020-04-09	0	1	1
2020-04-10	0	2	2
2020-04-11	1	1	2
2020-04-12	0	1	1
2020-04-13	3	0	3
2020-04-14	0	1	1
2020-04-16	1	1	2
2020-04-17	1	1	2
2020-04-20	1	2	3

Day	Number of singleton starts	Number of non-singleton starts	Total
2020-04-21	1	0	1
2020-04-22	2	3	5
2020-04-23	1	3	4
2020-04-24	3	0	3
2020-04-25	0	1	1
2020-05-06	1	0	1
2020-05-21	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
2020-03-02	
	2
2020-03-04	5
2020-03-05	3
2020-03-06	7
2020-03-07	5
2020-03-08	1
2020-03-09	12
2020-03-10	5
2020-03-11	11
2020-03-12	32
2020-03-13	42
2020-03-14	13
2020-03-15	8
2020-03-16	14
2020-03-17	31
2020-03-18	24
2020-03-19	28
2020-03-20	32
2020-03-21	35
2020-03-22	32
2020-03-23	84
2020-03-24	87
2020-03-25	71
2020-03-26	91
2020-03-27	87
2020-03-28	40
2020-03-29	53
2020-03-30	120
2020-03-31	86
2020-04-01	71
2020-04-02	48
2020-04-03	57
2020-04-04	45
2020-04-05	50
2020-04-06	71
2020-04-07	70
2020-04-08	43
2020-04-09	25
2020-04-10	22
2020-04-11	46
2020-04-12	61
2020-04-13	55
2020-04-14	50
2020-04-15	51
2020-04-16	62
2020-04-17	24
2020-04-18	55
2020-04-19	30
2020-04-20	68
2020-04-21	104
2020-04-22	99
2020-04-23	76
2020-04-24	91
2020-04-25	70

Day	Scotland
2020-04-26	49
2020-04-20	61
2020-04-21	35
2020-04-29	23
2020-04-30	24
2020-05-01	25
2020-05-02	13
2020-05-03	16
2020-05-04	9
2020-05-05	17
2020-05-06	31
2020-05-07	33
2020-05-08	25
2020-05-09	17
2020-05-10	21
2020-05-11	16
2020-05-12	22
2020-05-13	29
2020-05-14	47
2020-05-15	22
2020-05-16	19
2020-05-17	16
2020-05-18	29
2020-05-19	29
2020-05-20	15
2020-05-21	13
2020-05-22	8
2020-05-23	7
2020 - 05 - 24	4
2020 - 05 - 25	8
2020-05-26	3
2020-05-27	5
2020-05-28	4
2020-05-29	1
2020 - 05 - 31	1
2020-06-01	3
2020-06-02	11
2020-06-03	1
2020-06-04	7
2020-06-05	1
2020-06-06	2
2020-06-07	1
2020-06-08	1

 $\textbf{Table S7} \ \text{Raw data for the figure seven with the number of sequences assigned to each admin 2 region.}$

Admin2	Country	Number of sequences	Sequence group
		-	
ABERDEEN	Scotland	22	10-50
ABERDEENSHIRE	Scotland	5	1-10
ANGUS	Scotland	38	10-50
ARGYLL AND BUTE	Scotland	6	1-10
CLACKMANNANSHIRE	Scotland	2	1-10
DUMFRIES AND GALLOWAY	Scotland	68	50-100
DUNDEE	Scotland	140	100-150
EAST AYRSHIRE	Scotland	84	50-100
EAST DUNBARTONSHIRE	Scotland	36	10-50
EAST LOTHIAN	Scotland	55	50-100
EAST RENFREWSHIRE	Scotland	21	10-50
EDINBURGH	Scotland	456	400-500
EILEAN SIAR	Scotland	2	1-10
FALKIRK	Scotland	92	50-100
FIFE	Scotland	45	10-50
GLASGOW	Scotland	988	> 500
HIGHLAND	Scotland	9	1-10
INVERCLYDE	Scotland	15	10-50
MIDLOTHIAN	Scotland	134	100-150
MORAY	Scotland	6	1-10
NORTH AYRSHIRE	Scotland	5	1-10
NORTH LANARKSHIRE	Scotland	200	200-250
ORKNEY ISLANDS	Scotland	1	1-10
PERTHSHIRE AND KINROSS	Scotland	57	50-100
RENFREWSHIRE	Scotland	272	250-300
SCOTTISH BORDERS	Scotland	143	100-150
SHETLAND ISLANDS	Scotland	14	10-50
SOUTH AYRSHIRE	Scotland	3	1-10
SOUTH LANARKSHIRE	Scotland	28	10-50
STIRLING	Scotland	11	10-50
WEST DUNBARTONSHIRE	Scotland	20	10-50
WEST LOTHIAN	Scotland	119	100-150