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Sexually Transmitted Infections**Identifying and interpreting spatial temporal variation in diagnoses of infectious syphilis amongst men, England: 2009 to 2013**

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Identifying and interpreting spatial temporal variation in diagnoses of infectious syphilis amongst men, England: 2009 to 2013

Authors: Jakob Petersen(1), Maurizio Gibin(2), Bersabeh Sile(3), Ian Simms(3)

1. Institute for Social and Economic Research, University of Essex, Colchester, Essex, CO43SQ, UK, +44(01206)873683, jpeterb@essex.ac.uk
2. European Commission, Joint Research Centre, Isidra, Italy
3. HIV & STI Dept, Health Protection Services, Public Health England, London, UK

Key words

EPIDEMIOLOGY (GENERAL), HEALTH SERV RESEARCH, SURVEILLANCE, SYPHILIS

Abstract

Objectives

Spatial clusters and variations in the trajectory of local epidemics were explored in relation to sexual orientation, demographic factors, stage of syphilis infection and HIV serostatus.

Methods

Kulldorff's scan statistics (SaTScan) was used to distinguish endemic and temporary clusters using a two-stage analysis.

Results

Endemic areas were found in London, Manchester, Brighton and Blackpool. Up to 40% of diagnoses were found within an 11km radius of central London. Of men diagnosed with syphilis in London 80% were men who have sex with men (MSM). Annual incidence in London increased from 24 (23-26 95% CI) cases per 100000 male population in 2009 to 36 (34-38) in 2013. In comparison with clusters, endemic areas were characterised by: a significantly higher ($p<0.05$) proportion of MSM (83% compared to 73%), an increased HIV positivity (41% v 15%), age 35 to 44 (34% v 23%), lower proportion of patients born in the UK (50% v 79%), and a lower proportion of primary stage infection (40% v 47%). Space-time clusters outside endemic areas occurred in urban and rural areas, diagnoses fluctuating below 10 per month. Exponential increases in diagnoses resembling point source outbreaks were seen at two locations.

Conclusion

Control of syphilis in endemic areas has proved elusive and clusters present unique intervention opportunities. Investigating the diversity of local epidemics provides information that can be used to predict outbreak structure, plan and evaluate sexual health services, and guide public health investigation, hypothesis generation and research.

Introduction

Spatial clustering has been a key feature of the English infectious syphilis epidemic since it re-emerged in 1997 [1,2]. Increasing incidence has been characterised by outbreaks and foci, infection quickly becoming endemic in London, Manchester and Brighton [1]. Between 2009 and 2013, diagnoses in men increased by 18% (from 2507 to 2970). Most diagnoses were seen in men who have sex with men (MSM), diagnoses in this group increasing by 41% from 1697 to 2393 during this time. In contrast diagnoses seen in women decreased by 19% (345 to 279)[3]. The epidemic has been focused on infection in white MSM aged 25 to 34 many of whom have HIV co-infection and high numbers of sexual partners [1,2]. Localised clusters have followed a similar pattern although some have centred around young, socially vulnerable heterosexuals [2,4]. Contextual information collected from these clusters has provided insight into the complex local and national public health challenges associated with the syphilis epidemic but detection has relied on the vigilance of local sexual health and public health professionals. Effective control relies on early identification and case management. The aim of this study was to explore spatial clusters and variations in the trajectory of local epidemics in relation to sexual orientation, demographic factors, stage of syphilis infections and HIV serostatus.

Methods

Data sources

Diagnoses of infectious (primary, secondary, early latent) syphilis were extracted from the GUMCADv2 surveillance dataset for the calendar years 2009 to 2013 inclusive [5]. This anonymised dataset included the date of syphilis diagnoses and stage of infection as well as patient age, sexual orientation, ethnicity, country of birth and census tract of residence (2001 Middle Layer Super Output Area or MSA01). Patients were excluded if their address was not within England (n=780, 5% of diagnoses). Population weights were used to convert the mid-year population estimates for men between 2009 and 2013 (denominator spatial unit) from the Census 2011 to the Census 2001 geography (numerator spatial unit) [6,7]. Data were geocoded to the Office of National Statistics 2011 built-up area subdivisions [8].

Statistical methods

A detailed description of the statistical methods is provided in the Web Appendix. Briefly, Kulldorff's scan statistics method (SaTScan 9.3) was used [9,10]. The methodology emulates a surveillance scenario in which monthly scans for emerging space-time clusters could be evaluated as potential outbreaks. SaTScan creates a large number of circles which assess potential clusters ranging from single census tracts to circles containing 50% of the base population. The expected number of cases was calculated from the denominators assuming a Poisson distribution. The most likely cluster was found by comparing the likelihood ratio (observed/expected) within each circle with that outside. Significance levels and recurrence intervals of the spatial clusters were obtained from Monte-Carlo replications. Covariates were not included since denominator data for MSM were not routinely available. Figure 3 was created by smoothing and mapping data using kernel density estimation [11].

Space-time cluster analysis - SaTScan

A two tier analytical strategy was adopted to prevent endemic areas from 'swamping' smaller outbreaks. Endemic areas were identified using spatial cluster analyses carried out for five calendar years in a row [12]. Data from endemic areas were then removed and prospective space-time cluster analysis was carried out for each month between January 2010 and December 2013 scanning across

1 a 6 months' time window. Since syphilis outbreaks can be sustained for several months the temporal
2 window was selected at 1 to 6 months. For areas where clusters were detected time trends were
3 smoothed using the Epanechnikov kernel function and plotted together with the results of the
4 monthly prospective space-time analyses.
5
6

7 **Results**

8 *Comparison of the characteristics of diagnoses made in London with the Rest of England*

9
10 Between 2009 and 2013, 12521 diagnoses were reported, 6082 (49%) of which were seen in London
11 (Table 1). Incidence in London was three times that seen in England overall. The proportion of men
12 diagnosed with syphilis who were MSM varied from 76% to 87% in London compared to 70% to 81%
13 in England. Diagnoses in patients born abroad accounted for nearly half of all diagnoses in London
14 (43 to 51%) whereas for England diagnoses were higher in UK born men (59 to 65%). Modal age (30
15 years) was the same for men born abroad whilst the age distribution of UK born men was bimodal;
16 peaks being seen in the late 20s and early 40s within and outwith London respectively (Figure 1).
17 Bimodality was observed in both the MSM and non-MSM group in initial analyses. Diagnoses in men
18 of white ethnic identity accounted for 67% to 69% of all diagnoses in London and 75 to 77% in
19 England. Diagnoses in black men varied from 10% to 14% in London and 6 to 8% in England. In
20 England, the majority of diagnoses of infectious syphilis consisted of primary infection, whilst there
21 was a move to a more equal distribution between primary, secondary and early latent in London
22 between 2009 and 2013. The proportion of syphilis diagnoses co-infected with HIV increased from
23 34% to 47% between 2009 and 2013. For England co-infection increased from 26% to 37% over the
24 same period.
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Table 1 Characteristics of diagnoses of infectious syphilis in men, London and England: 2009 to 2013.

| | London | | | | | England Total | | | | |
|---------------------------|-----------|-----------|-----------|---------|---------|---------------|---------|-------|----------|-----------|
| | 2009 | 2010 | 2011 | 2012 | 2013 | 2009 | 2010 | 2011 | 2012 | 2013 |
| <i>Diagnoses (n)</i> | 968 | 975 | 1253 | 1337 | 1549 | 2251 | 2223 | 2527 | 2640 | 2880 |
| <i>Sexual orientation</i> | | | | | | | | | | |
| MSM (%) | 77.1 | 76 | 85.9 | 86.6 | 87.3 | 69.8 | 70.4 | 78.2 | 79.5 | 81 |
| Other males (%) | 22.9 | 24 | 14.1 | 13.4 | 12.7 | 30.2 | 29.6 | 21.8 | 20.5 | 19 |
| <i>Age (mean, years)</i> | 36.8 | 37.8 | 37.3 | 36.8 | 37.9 | 36.7 | 37.0 | 36.4 | 36.6 | 37.2 |
| <i>Age group (years)</i> | | | | | | | | | | |
| <25 (%) | 9.9 | 8.4 | 7.7 | 9.8 | 6.7 | 13.6 | 13.0 | 14 | 14.4 | 11.8 |
| 25-34 (%) | 34 | 31.5 | 33.7 | 34.9 | 33.5 | 30.7 | 31.5 | 33.1 | 33.1 | 33.1 |
| 35-44 (%) | 35.5 | 34.1 | 37 | 32.2 | 34.3 | 32.1 | 30.6 | 31.1 | 27.9 | 29 |
| 45+ (%) | 20.5 | 25.9 | 21.6 | 23 | 25.5 | 23.4 | 24.5 | 21.5 | 24.4 | 25.9 |
| <i>Ethnicity</i> | | | | | | | | | | |
| White (%) | 67.9 | 68.9 | 69.2 | 67.2 | 67.2 | 76.7 | 76.2 | 76.2 | 75.4 | 74.8 |
| Mixed (%) | 4.3 | 4.1 | 4.9 | 5.2 | 5 | 2.7 | 2.7 | 3.4 | 3.4 | 3.7 |
| Asian (%) | 4.3 | 4.4 | 4.5 | 5.2 | 4.8 | 4.1 | 3.9 | 4.2 | 4.7 | 4.4 |
| Black (%) | 10.6 | 13.6 | 10.2 | 11.1 | 11.7 | 6 | 7.8 | 6.8 | 6.8 | 7.2 |
| Other (%)* | 12.8 | 8.9 | 11.1 | 11.3 | 11.3 | 10.5 | 9.4 | 9.4 | 9.7 | 9.9 |
| <i>Birthplace</i> | | | | | | | | | | |
| UK (%) | 42.5 | 45 | 42.2 | 41.5 | 40.7 | 64.7 | 65.4 | 62.2 | 61 | 58.9 |
| Abroad (%) | 44 | 43 | 50.3 | 49.4 | 50.7 | 25.9 | 26.6 | 32.3 | 31.9 | 33.1 |
| Other (%) | 13.5 | 12 | 7.5 | 9.1 | 8.7 | 9.4 | 8 | 5.5 | 7.1 | 8 |
| <i>Infection stage</i> | | | | | | | | | | |
| Primary (%) | 51.2 | 41.4 | 36.8 | 36.1 | 42.3 | 47 | 41.8 | 42.2 | 40.6 | 43 |
| Secondary (%) | 23.2 | 30.4 | 33.9 | 34.6 | 27.2 | 27.3 | 30.6 | 29.8 | 31.4 | 27.7 |
| Early latent (%) | 25.5 | 28.2 | 29.3 | 29.2 | 30.5 | 25.7 | 27.6 | 27.9 | 28 | 29.3 |
| <i>HIV serostatus</i> | | | | | | | | | | |
| Positive (%) | 34.4 | 41.1 | 44 | 44 | 46.7 | 26.4 | 29.1 | 30.8 | 33.8 | 36.5 |
| <i>Annual incidence</i> | | | | | | | | | | |
| Patients (n) | 943 | 945 | 1192 | 1283 | 1491 | 2207 | 2174 | 2449 | 2560 | 2799 |
| /100k male pop | 24.1 | 23.8 | 29.4 | 31.3 | 35.8 | 8.6 | 8.4 | 9.4 | 9.7 | 10.5 |
| 95% CI | 22.6;25.7 | 22.3;25.3 | 27.8;31.1 | 29.6;33 | 34;37.7 | 8.3;9 | 8.1;8.8 | 9;9.7 | 9.4;10.1 | 10.2;10.9 |

* Other includes all other ethnic groups and records for which is information was missing.

Table 2 Characteristics of diagnoses in endemic and cluster areas, men: 2009 to 2013

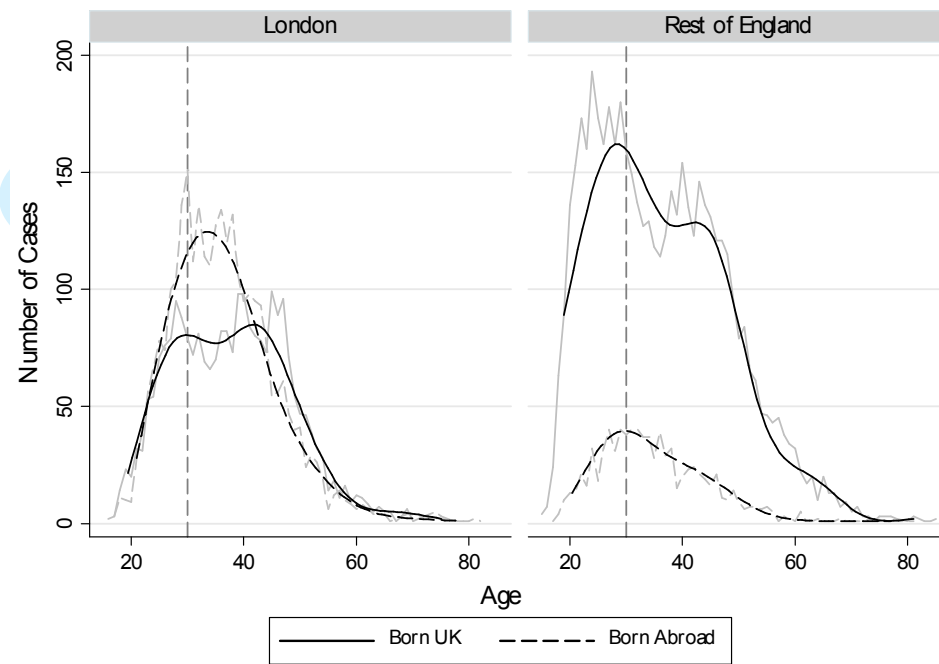
| | Cases (n) | MSM (%) | HIV-Positive (%) | Mean age (years) | <25 years (%) | 25-34 years (%) | 35-44 years (%) | 45+ years (%) | Born UK (%) | Primary syphilis (%) | Secondary syphilis (%) | Early latent (%) |
|----------------------------|--------------|------------|------------------|------------------|---------------|-----------------|-----------------|---------------|-------------|----------------------|------------------------|------------------|
| Endemic areas | 7514 | 83 | 41 | 37.0 | 10 | 34 | 34 | 23 | 50 | 40 | 30 | 30 |
| London | 6082 | 83 | 43 | 37.3 | 8 | 34 | 35 | 23 | 42 | 41 | 30 | 29 |
| Greater Manchester | 1040 | 81 | 34 | 34.8 | 16 | 37 | 29 | 17 | 84 | 37 | 25 | 37 |
| Brighton | 252 | 70 | 45 | 39.4 | 10 | 24 | 35 | 31 | 73 | 33 | 35 | 32 |
| Blackpool | 140 | 85 | 21 | 36.0 | 22 | 26 | 26 | 25 | 94 | 34 | 24 | 43 |
| Space-time clusters | 1146 | 73* | 15* | 36.7 | 19* | 36 | 23* | 22 | 79* | 47* | 30 | 23* |
| West Yorkshire | 384 | 74 | 19* | 35.3 | 16* | 38 | 24* | 21 | 82* | 49* | 27 | 24 |
| Tyneside | 232 | 74 | 9* | 32.2 | 26* | 38 | 25 | 11* | 72* | 35 | 32 | 33 |
| Bournemouth | 125 | 83 | 18* | 38.8 | 8 | 36 | 30 | 25 | 77* | 46 | 34 | 20 |
| Bristol | 74 | 82 | 11* | 34.8 | 26* | 34 | 15* | 24 | 80* | 73* | 19 | 8* |
| Nottingham | 72 | 51* | 8* | 36.3 | 17 | 33 | 24 | 26 | 85* | 43 | 38 | 19 |
| Herefordshire | 57 | 81 | 16* | 39.4 | 19 | 19 | 26 | 35 | 81* | 70* | 26 | 4* |
| Southampton | 54 | 76 | 22* | 37.5 | 17 | 39 | 19 | 24 | 65* | 67* | 17 | 17 |
| Kingston-upon- Hull | 54 | 50* | 11* | 33.3 | 30* | 28 | 17* | 26 | 82* | 48 | 35 | 17 |
| Plymouth | 49 | 76 | 14* | 38.1 | 22* | 27 | 8* | 43* | 96* | 18* | 51* | 31 |
| Southend | 45 | 69* | 22 | 34.2 | 16 | 47 | 16* | 16 | 73 | 47 | 27 | 27 |
| Rest of England | 3861 | 65* | 18* | 36.7 | 19* | 29* | 25* | 27* | 81* | 47* | 29 | 24* |
| Total | 12521 | 76 | 32 | 36.8 | 13 | 32 | 30 | 24 | 62 | 43 | 29 | 28 |

■ value above average

* proportion of non-endemic area significantly different from endemic area sub-total (95% level).

Characteristics of space-time endemic areas and clusters

Analysis of endemic areas and clusters revealed considerable heterogeneity (Table 2; Figures 2 and 3). Up to 40% of all diagnoses were resident within a 11km radius in Central London. Manchester, the West Yorkshire conurbation, Brighton, Tyneside, Blackpool and Bournemouth included 8% (1040), 3% (384), 2% (252), 2% (232), 1% (140) and 1% (125) of diagnoses, respectively. Sixty percent of diagnoses were seen in endemic areas whereas 9% and 31% seen in space-time clusters and the rest of England respectively (Table 2). There was a strong annual increase in diagnoses in London and a rising trend in Brighton whereas diagnoses in Manchester fell (Figure 4). Other endemic areas saw a lower but consistent burden of diagnoses: Brighton and Blackpool experienced weeks during which no diagnoses were made.

Figure 1 Diagnoses of infectious syphilis in men by birthplace & region: 2009 to 2013*

*Vertical line = age 30 years (mode).

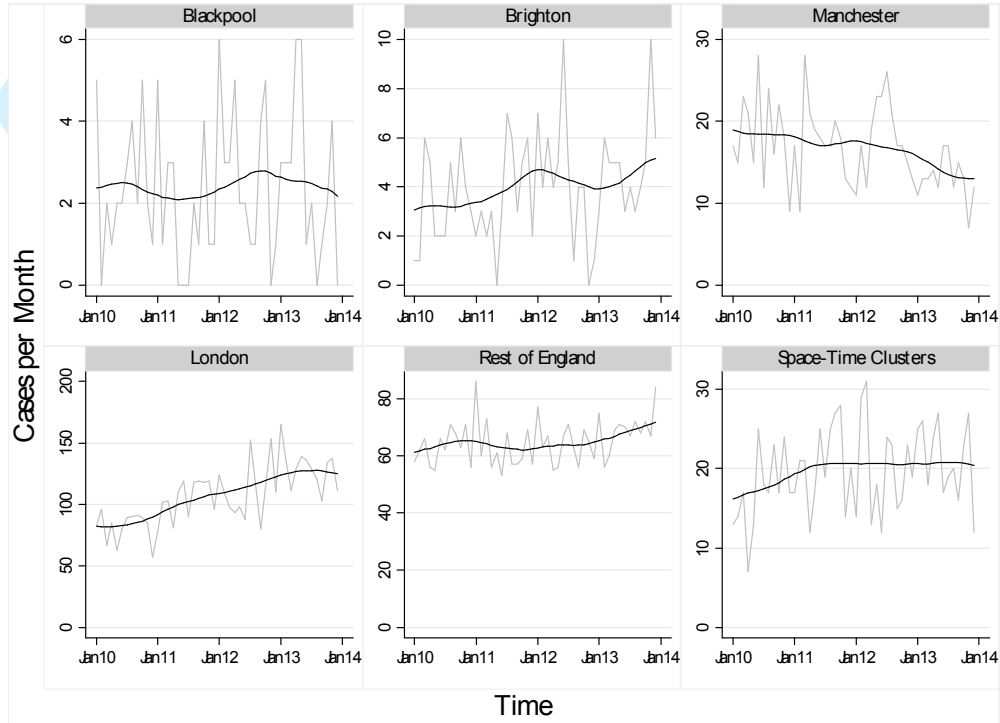
Figure 2 Space-time clusters of male infectious syphilis outbreaks: 2010 to 2013

[Map – England – attached]

Figure 3 Case density map of male syphilis cases in central London: 2009 to 2013*

[Map – London - attached]

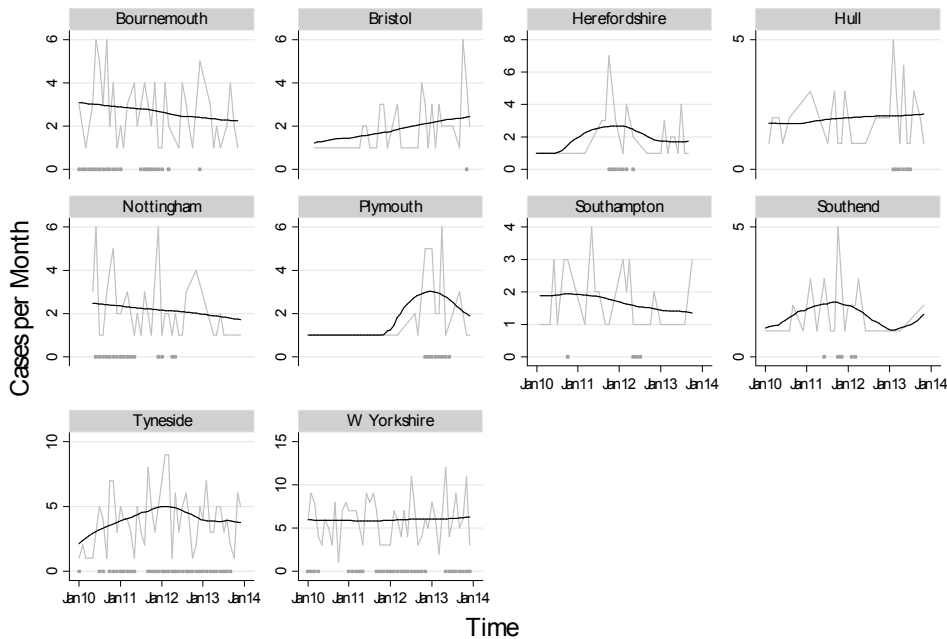
Figure 4 Monthly cases of infectious syphilis in males by endemic areas and the rest of England: 2010 to 2013*†



* Solid black lines represent the observations smoothed using Epanechnikov kernel function.

† The range on the y-axes varies between panels.

Figure 5 Monthly cases of infectious syphilis in men, clusters: 2010 to 2013*†



* Area units were defined by ONS's built-up area polygons for urban areas and by local authority for rural areas.

† Solid black lines represent the observations smoothed using Epanechnikov kernel function.

.... Months where 1 to 6 months space-time clusters were detected.

1
2 In comparison with endemic areas, space-time clusters only included 1146 diagnoses (9%). Clusters
3 occurred in urban and rural locations diagnoses fluctuating below 10 cases per month (Figure 5, the
4 small squares on the baseline indicate when space-time clusters were identified). The exponential
5 increase in diagnoses seen in Herefordshire and Plymouth resemble point source outbreaks. Clusters
6 were significantly associated with a higher proportion of diagnoses made in men born in the UK and
7 lower proportions of MSM, a lower proportion of HIV co-infection, and a higher proportion of <25
8 year olds, and a lower proportion of 35 to 44 year olds (Table 2).
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12 On completion of the two tier analysis, 3861 diagnoses (31%) could not be classified into an endemic
13 or cluster areas. Overall the characteristics of the sporadic cases were similar to those diagnoses
14 made in cluster areas.
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17 Discussion

18 Spatial-temporal variation is a critical feature of the infectious syphilis epidemic seen in men in
19 England. London bears the heaviest burden of diagnoses in terms of incidence, density and absolute
20 numbers. The majority of diagnoses were seen within endemic areas but these four locations
21 differed in terms of the characteristics of each local epidemic. In contrast, space-time clusters only
22 represent a tenth of diagnoses, developed over several months and like endemic areas showed
23 substantial variation in the stage of infection, HIV-positivity, sexual orientation, age, and place of
24 birth.
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29 The main weakness of the investigation was the lack of denominator data for the MSM population
30 which in England is focused on London, Manchester, Brighton and Blackpool. This is likely to have
31 attenuated the strength of the detected relationships. The advantage of SaTScan analysis is that it
32 does not require an *a priori* hypothesis about cluster location, size, or duration. The software adjusts
33 for inhomogeneous population density, addresses multiple testing, preserves patient confidentiality
34 at small area level and adjusts for inhomogeneous population density within the base population. In
35 addition, since SaTScan analysis and kernel density maps are not dependent on the denominators
36 created by administrative units, the analysis was not influenced by the modifiable areal unit problem
37 [13,14].
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41 During the period covered by this study a number of outbreak investigations amongst MSM and
42 young heterosexuals were undertaken by local outbreak control teams across England. Some
43 investigations were published but many could not be put into the public domain because of the
44 need to protect the confidentiality of the patients who were involved in small scale investigations
45 [15,16]. Of the space-time clusters detected all had been the subject of investigations but the the
46 analysis did not identify two investigations based on MSM living in East Anglia [17]. The clusters may
47 have been pushed below the limit of detection because of the small number of diagnoses involved in
48 the outbreak. The size of the epidemic may also have been underestimated for example due to the
49 limited success of partner notification and variations in patient clinical management between
50 locations. Although all diagnoses conformed to British Association for Sexual Health and HIV
51 standards, testing frequency varies across England [18]. Genitourinary medicine services within
52 London generally undertake more intensive patient management, a process which could increase
53 the syphilis diagnostic rate compared to HIV negative patients. This could account for the increased
54 detection of early latent syphilis within London.
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1 The epidemic explored here has the characteristics of a metapopulation. Essentially, populations in
2 which syphilis is endemic sustain the epidemic and seed sexual networks in smaller satellite
3 populations through bridging populations creating local clusters. Infection cycles relatively
4 independently within these populations [19]. Smaller populations are more likely to become extinct,
5 elimination occurring due to density dependent or independent factors or stochastic events. The
6 sporadic diagnoses that could not be classified as endemic or cluster could have been recent
7 infections that had the potential to start localised epidemics. Alternatively they may have been more
8 long standing infections from which local sustained transmission had not emerged.

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11 Since 2000 the epidemiology of syphilis, *Neisseria gonorrhoeae*, lymphogranuloma venereum (LGV),
12 hepatitis C, and sexually transmissible infections such as *Shigella flexneri*, *S. sonnei* and VTEC0117
13 has changed significantly, influenced by population flow, sero-adaptive behaviour amongst HIV
14 positive MSM, antimicrobial resistance together with advances in diagnostic techniques and
15 therapeutic agents [20–22]. Chemsex drugs have become widely used and, since their first launch in
16 early 2009, location based sexual networking applications have become an increasingly popular
17 method of meeting sex partners and organising sex parties [23]. In effect sexual network structure is
18 changing from a density dependent factor into a density independent factor thereby increasing the
19 potential for the infection transmission. Such developments make infection control increasingly
20 challenging particularly in locations where infection is endemic. Clusters can vary in size and usually
21 take months to control. Outbreaks may be highlighted by local sexual health professionals or
22 through exceedance reporting at Public Health England (PHE) Colindale, PHE Field Epidemiology
23 Services and laboratories. PHE has introduced automated spatio-temporal detection tools based on
24 SaTScan and *R* to monitor the epidemiology of gonorrhoea in England [24,25]. This tool, which is
25 available to public health professionals, is used by local services as part of their outbreak detection
26 strategy to increase awareness to existing and emerging sexual health challenges and inform service
27 planning. The method is to be extended to include infectious syphilis. Outbreak detection methods
28 also need to be fine-tuned to enable the detection and characterisation of infection clusters
29 amongst young heterosexuals and women in low population density areas so we can better
30 understand why outbreaks occur and assess the potential for vertical transmission.

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38 Heterogeneity across the syphilis epidemic emphasises the importance of planning local control and
39 prevention strategies. Here we have demonstrated how spatio-temporal analysis can be used to
40 explore the characteristics that make locations inherently similar or unique in space and time. Such
41 information could be used to consider the structure of future outbreaks, plan, and evaluate sexual
42 health services, and provide a starting point for public health investigation, hypothesis generation
43 and research. The future challenge is adapting such analytical and visualisation techniques for use
44 with real-time GUMCADv2 reporting to provide the evidence base that enables health care
45 professionals to respond to changes within the developing epidemic.

46 47 48 49 **Key points**

- 50 • Spatial-temporal variation is a critical feature of the infectious syphilis epidemic and
51 emphasises the importance of planning local control and prevention strategies.
- 52 • Most diagnoses were seen within four endemic areas, London, Manchester, Brighton and
53 Blackpool. The characteristics of each of these epidemics were different.
- 54 • The heaviest burden of diagnoses in terms of incidence, density and absolute numbers was
55 seen in London.
- 56 • Space-time clusters, which included 10% of all diagnoses, were found in urban and rural
57 areas.
- 58
- 59
- 60

- Space-time clusters developed over several months and varied in terms of stage of infection, HIV-serostatus, sexual orientation, age, and place of birth.

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Conflict of interest

None.

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Contributorship statement

JP, MG and IS were closely involved with the design, conduct, analysis, presentation and interpretation of the study findings. BS extracted and quality assured the GUMCAD dataset.

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Confidential: For Review Only

Web-resources for Methods used in:

Identifying and interpreting spatial temporal variation in diagnoses of infectious syphilis amongst men, England: 2009 – 2013.

Data source

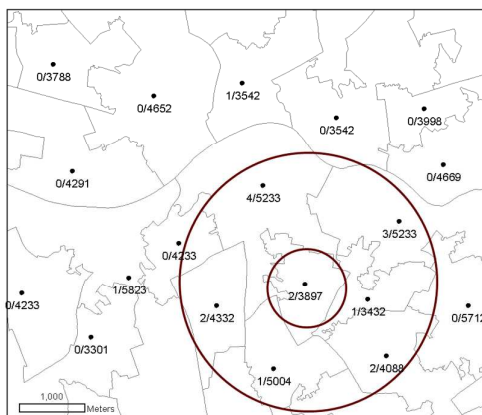
The NHS Genitourinary Medicine Clinic Activity Dataset (GUMCAD) was set up in 2008 to create a single, national database to monitor trends in STI diagnoses, to determine specific risk groups and to assess local service needs [1].

Statistical methods

A variety of space-time cluster detection techniques are available, e.g. SaTScan, ClusterSeer, Geosurveillance, and Surveillance for R; of which SaTScan is the most popular technique [2]. SaTScan has been used in hundreds of epidemiological and simulated health service datasets [3] and Kulldorff's prospective scan statistics method with discrete Poisson model (SaTScan 9.3) was used here to explore high risk clusters [4,5]. SaTScan has a number of strengths: no *a priori* hypotheses about cluster location, size or duration is needed; the software adjusts for inhomogeneous population density; takes multiple testing into account and provides a single test statistics for the most likely cluster. Potential secondary clusters are also ranked.

At the start of the analytical process SaTScan creates concentric circles around a centroid. In the example, the centroid of each Middle Super Output Area is located (Figure). SaTScan sums the number of cases and population-at-risk inside and outside of the circle. Candidate circles with a higher ratio of observed-to-expected are selected. To assess how likely the results could have been generated by chance alone the clusters are tested using a large number of Monte-Carlo replications.

Figure Likelihood ratio test: identifying cases and at-risk-population. Two of the many concentric circles of varying radii evaluated are shown (fictive data).



● Middle Layer Super Output Area centroid

Space-time cluster analysis - SaTScan

For space-time analysis cluster cylinders were constructed in which height represented the time dimension. The analyses can be computationally intensive especially if a large number of geographical data points are included. Here, an exploratory analysis indicated that it was possible to

1 detect the same clusters at MSOA01 as at LSOA01 level and so, to reduce computation time, the
2 analysis was performed at MSOA01 level. Large, relatively low risk clusters tend to have higher
3 significance levels than the smaller high-risk clusters that they may contain [6]. Data from the
4 smaller clusters with higher risk are therefore suppressed by default from the output if they fall
5 inside a larger significant cluster. Initially, such high-risk clusters were explored by progressively
6 restricting the maximum base population from the default to $\leq 0.125\%$ (equivalent to a base
7 population of around 30,000 men per year) by changing the output criteria of the analyses with the
8 $\leq 50\%$ base population restriction. No additional tests were carried out. The changes to clusters
9 outside London were minimal below the 20% population threshold.

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14 A purely spatial cluster analysis was carried out for each calendar year to identify persistent clusters
15 over time. Persistent clusters were found each year in London, Manchester, Blackpool and Brighton.
16 Spatial clusters were found in Bournemouth for three out of the four years studied. These clusters
17 were interpreted as the major core areas.

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20 Prospective space-time cluster analysis was carried out monthly between January 2010 and
21 December 2013 to identify outbreak areas outside the core areas. The total number of consecutive
22 tests was 48 and only clusters with recurrence intervals at or above 50 have been reported [5].
23 Population thresholds were evaluated and set to 1% of person-time. For these analyses core areas
24 were removed from both the numerator and denominator side. The temporal window was selected
25 as 1-6 months as syphilis outbreaks are known to build up over order several months. This two-tier
26 strategy was used to avoid smaller and temporal outbreaks to be swamped by the larger and
27 endemic core areas. Localities were identified by overlaying a map of the cluster cylinder bases with
28 the boundary sets from ONS's 2011 built-up area and built-up area subdivisions.

31 Web resource references

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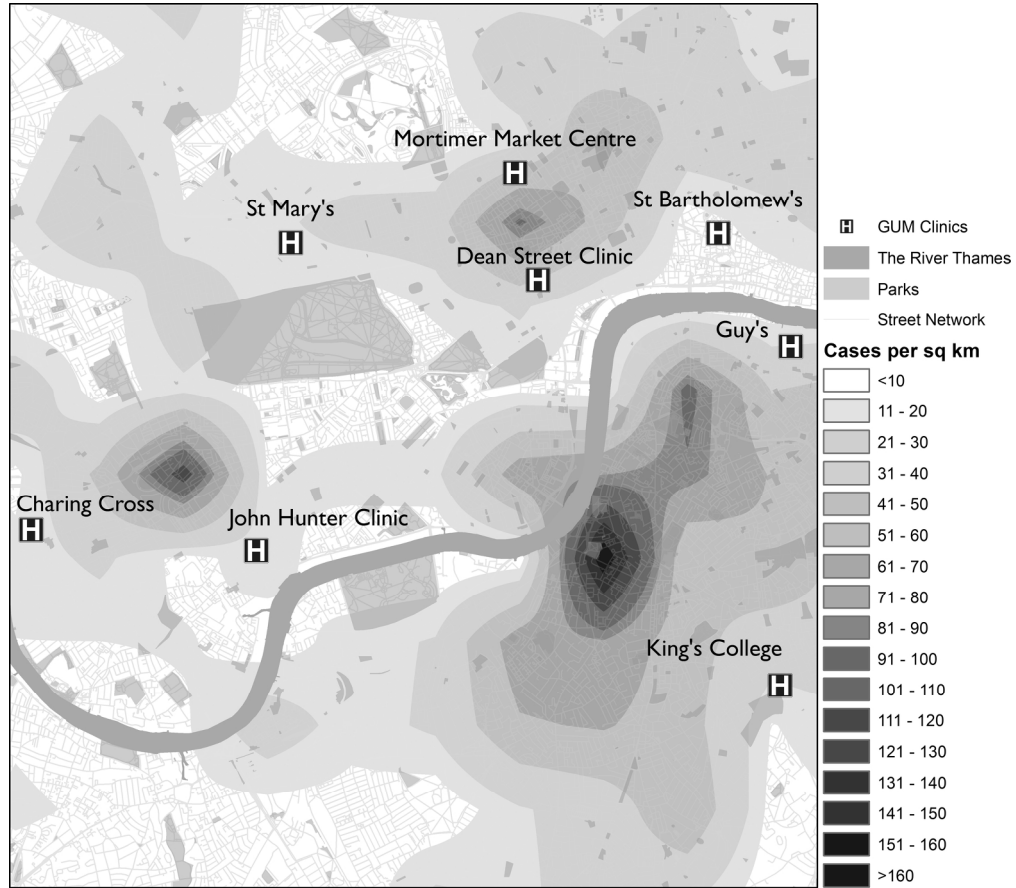
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