NIHR Health Protection Research Unit in Blood Borne and Sexually Transmitted Infections

Epidemiology of sexually transmissible enteric infections in men who have sex with men (STEIM): protocol for a longitudinal pilot study in Brighton and Sussex

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1. BACKGROUND

- Increasing number of enteric pathogen outbreaks reported among men-who-have-sexwith-men (MSM), often associated with high levels of antimicrobial resistance (AMR)
- Evidence suggests that asymptomatic carriage might play an important role in sustaining transmission in specific sexual networks of MSM
- There is a need to better understand the drivers of sustained transmission of, and development of AMR in, enteric infections among MSM to inform the design and delivery of interventions.

2. PILOT STUDY OBJECTIVES

- To assess the feasibility and acceptability of longitudinal sample and epidemiological data collection among MSM attending NHS SHCs
- To compare rectal swabs with faecal samples for enteric pathogen and AMR detection, and molecular typing
- To provide preliminary data on:

of interventions

- Such a study will require the collection of samples and epidemiological data over a period of several months
- Rectal swabs are routinely collected from MSM attending sexual health clinics (SHCs) for routine care and may provide a practical alternative to stool samples
- Overall prevalence of enteric pathogens
- Duration of carriage
- Within-host evolution of pathogens, including AMR



3. STUDY POPULATION

Location and sample size

- Individuals attending SHCs in Brighton and Sussex for routine testing and care, regardless of symptoms
- 200 individuals

Eligibility criteria

- MSM, including men (cis/transgender), transwomen or gender-diverse people reporting sex with a man (cis/transgender) or non-binary person assigned male at birth
- Report sex with another man in the past 3 months

4. STUDY DESIGN



5. ANALYSES

- Testing and analysis of stool samples and rectal swabs
 - PCR testing to include Shigella spp., Campylobacter spp., Salmonella spp., E. coli, E. histolytica, Giardia lamblia, hepatitis A virus

1 stool sample (optional)

1 rectal swab

Baseline questionnaire (sexual behaviour, travel history, antibiotic use, symptoms)



Qualitative interview (optional)

1:1 interview exploring the barriers and motivators to study participation, and the collection of samples over time

As part of the Patient and Public Participation and Engagement strategy, cognitive interviews were used to explore general interest and acceptability of the study design and comprehension of participant information; suggestions for improvement were incorporated

- Culture and whole genome sequencing
- Metagenomics to compare complete diversity of pathogens and genetic determinants of AMR
- Linkage of test results and questionnaire responses to national STI surveillance data
 - Information on STI and HIV testing and diagnoses
 - Participants provide consent for data linkage
- Thematic analysis of 1:1 interviews
 - Identification of emerging themes and exploration of
- The study has been reviewed and approved by London South East NHS Research Ethics Committee (21/LO/0891). Recruitment will begin in March 2022.

any patterns observed

6. EXPECTED OUTCOMES

- Key analyses will address the following:
 - > Willingness to take part in the study and to return samples over a period of several months
 - > The characteristics of those who take part in the study compared to those who do not
 - > Proportion of participants who agree to the linkage of biological samples and questionnaire responses to STI surveillance data
 - > Sensitivity of rectal swabs compared to stool samples to determine whether future studies involving longitudinal sampling can use rectal swabs only
 - > Preliminary data on the prevalence of enteric pathogens and duration of carriage, and how these vary according to patient characteristics
 - Genomic comparisons to infer within-host evolutionary dynamics and processes

• The findings will be used to inform the design of a larger study at multiple SHCs involving the repeated collection of biological samples and epidemiological data

The research was funded by the National Institute for Health Research Health Protection Research Unit (NIHR HPRU) in Blood Borne and Sexually Transmitted Infections at University College London in partnership with UK Health Security Agency (UK HSA). The views expressed are those of the authors and not necessarily those of the NIHR, the Department of Health and Social Care or UK HSA. We acknowledge members of the NIHR HPRU in BBSTI Steering Committee: Professor Caroline Sabin (HPRU Director), Dr John Saunders (UK HSA Lead), Professor Catherine Mercer, Professor Gwenda Hughes, Professor Greta Rait, Dr Ruth Simmons, Professor William Rosenberg, Dr Tamyo Mbisa, Professor Rosalind Raine, Dr Sema Mandal, Dr Rosamund Yu, Dr Samreen Ijaz, Dr Fabiana Lorencatto, Dr Rachel Hunter, Dr Kirsty Foster and Dr Mamooma Tahir.









UK Health Security Agency