A study of multidrug resistant organisms in the adult intensive care unit at Addenbrooke's Hospital

Submission date 13/02/2020	Recruitment status No longer recruiting	Prospectively registered		
		[X] Protocol		
Registration date 20/02/2020	Overall study status Completed	Statistical analysis plan		
		Results		
Last Edited 06/03/2020	Condition category Infections and Infestations	Individual participant data		
		Record updated in last year		

Plain English summary of protocol

Background and study aims

Healthcare-associated infections (HCAI) affect up to 10% of hospital patients, and are associated with an increased risk of death. Multidrug-resistant organisms (MDRO) are more common in hospitals. Patients in intensive care units (ICU) are particularly vulnerable to HCAI, and several outbreaks of infection with MDRO have been reported.

Current practices for screening for MDRO vary between countries, hospitals and units, reflecting a lack of information, and uncertainty about best practice. One strategy to reduce HCAI in ICUs would be to perform screening to look for MDRO in patients admitted to ICUs. This would enable earlier identification and treatment of MDRO and implementation of appropriate infection control measures to prevent their spread.

Whole-genome sequencing (WGS) is novel technology, which is more discriminatory than currently available typing methods. We are conducting a study to determine the rates of carriage, infection, and transmission of MDRO in the adult ICU at Addenbrooke's Hospital, using WGS. This study will facilitate translation of this technology from a research tool into day-to-day clinical practice. Information from this study will be used to inform infection control and public health policies and procedures.

Who can participate?

Any adult admitted to the John Farman Intensive Care Unit during the study period.

What does the study involve?

All patients admitted to ICU during the study period will be screened for MDRO and clinical data on infections and antimicrobial use will be collected.

What are the possible benefits and risks of participating?

No direct benefits for study participation. In terms of the risks of participation in the study,

these are negligible as the study is observational in nature and there are no study-specific interventions or treatments. All specimens will be collected by experienced nursing staff, in accordance with routine clinical practice.

Where is the study run from? Cambridge Biomedical Campus (UK)

When is the study starting and how long is it expected to run for? June 2016 to December 2016

Who is funding the study?

- 1. Academy of Medical Sciences (UK)
- 2. The Health Foundation (UK)
- 3. National Institutes of Health Research Cambridge Biomedical Research Centre (UK)

Who is the main contact? Dr Estee Torok et317@cam.ac.uk

Contact information

Type(s)

Scientific

Contact name

Dr Estee Torok

ORCID ID

http://orcid.org/0000-0001-9098-8590

Contact details

University of Cambridge, Department of Medicine, Box 157, Addenbrooke's Hospital, Hills Road, Cambridge
United Kingdom
CB1 7TP
01223 336845
et317@cam.ac.uk

Additional identifiers

EudraCT/CTIS number

Nil known

IRAS number

180415

ClinicalTrials.gov number

Nil known

Secondary identifying numbers

Study information

Scientific Title

Whole-genome sequencing to investigate colonisation and transmission of multidrug-resistant organisms in the adult intensive care unit at Addenbrooke's Hospital

Acronym

ICU001

Study objectives

What is the prevalence of the colonisation and is there evidence of transmission of multidrugresistant organisms in the adult intensive care unit?

Ethics approval required

Old ethics approval format

Ethics approval(s)

Approved 21/04/2016, NHS Health Research Authority, East of England - Cambridge Central Research Ethics Committee (Royal Standard Place, Nottingham NG1 6FS, UK; +44 (0)207 104 8388; NRESCommittee.EastofEngland-CambridgeCentral@nhs.net), ref: 15/EE/0318

Study design

Observational cohort study

Primary study design

Observational

Secondary study design

Cohort study

Study setting(s)

Hospital

Study type(s)

Screening

Participant information sheet

Not available in web format, please use the contact details below to request a patient information sheet

Health condition(s) or problem(s) studied

Surveillance for multidrug-resistant organisms

Interventions

All patients admitted to ICU during the study period will be screened for MDRO and clinical data on infections and antimicrobial use will be collected.

Participants will be screened for MDRO on admission to the John Farman Intensive Care Unit. All specimens will be assigned a unique anonymised identification number prior to transfer to the research laboratory the Department of Medicine for processing.

Participants will be screened for MDRO on discharge from ICU, and weekly during their ICU admission if the duration of admission is 7 days or longer.

Intervention Type

Other

Primary outcome measure

Measured using patient records during the study period:

- 1. Number of patients colonised with multidrug-resistant organisms
- 2. Number of patients with clinical evidence of infection with multidrug-resistant organisms
- 3. Number of transmission events of multidrug-resistant organisms

Secondary outcome measures

Measured using patient records during the study period:

- 1. Risk factors for colonisation / infection with multidrug-resistant organisms
- 2. Outcome of patients colonised / infected with multidrug-resistant organisms
- 3. Cost-consequences of whole-genome sequencing versus standard epidemiological investigation / typing for surveillance and investigation of suspected outbreaks and the management of confirmed outbreaks

Overall study start date

01/01/2016

Completion date

30/06/2017

Eligibility

Key inclusion criteria

- 1. Aged 18 years old or older
- 2. Male or female
- 3. Admitted to the John Farman Intensive Care Unit during the study period

Participant type(s)

Patient

Age group

Adult

Lower age limit

18 Years

Sex

Both

Target number of participants

All patients admitted to ICU in a 6 month period (N=400)

Key exclusion criteria

Does not fulfil study inclusion criteria

Date of first enrolment

20/06/2016

Date of final enrolment

20/12/2016

Locations

Countries of recruitment

England

United Kingdom

Study participating centre

Cambridge Biomedical Campus

Cambridge University Hospitals NHS Foundation Trust Hills Road Cambridge United Kingdom CB2 0QQ

Sponsor information

Organisation

Cambridge University Hospitals NHS Foundation Trust

Sponsor details

R&D Department
Cambridge University Hospitals NHS Foundation Trust
Hills Road
Cambridge
England
United Kingdom
CB2 0QQ
+44 (0)1223 348490
research@addenbrookes.nhs.uk

Sponsor type

Hospital/treatment centre

Website

http://www.cuh.org.uk/

ROR

https://ror.org/04v54gj93

Organisation

University of Cambridge

Sponsor details

School of Clinical Medicine
Box 111 Addenbrooke's Hospital
Hills Road
Cambridge
England
United Kingdom
CB2 0QQ
+44 (0)1223 766362
jo.deckers@admin.cam.ac.uk

Sponsor type

University/education

Website

http://www.cam.ac.uk/

ROR

https://ror.org/013meh722

Funder(s)

Funder type

Research organisation

Funder Name

Academy of Medical Sciences

Alternative Name(s)

The Academy of Medical Sciences

Funding Body Type

Private sector organisation

Funding Body Subtype

Other non-profit organizations

Location

United Kingdom

Funder Name

The Health Foundation

Funder Name

National Institutes of Health Research Cambridge Biomedical Research Centre

Results and Publications

Publication and dissemination plan

Planned publication in a high-impact peer-reviewed journal.

Intention to publish date

31/12/2020

Individual participant data (IPD) sharing plan

All data generated or analysed during this study will be included in the subsequent results publication.

IPD sharing plan summary

Other

Study outputs

Output type	Details version v2.0	Date created	Date added	Peer reviewed?	Patient-facing?
Protocol file			06/03/2020	No	No
HRA research summary			28/06/2023	No	No