

Inclusion/Exclusion

GenOMICC participant IDs are found inside each specimen kit.

GenOMICC ID number _____

Please ensure that the ID Number is correct and is in the format 'GCC' followed by exactly 5 digits before saving this page

Inclusion Criteria

Is the patient critically ill (requires continuous monitoring, mechanical ventilation or organ support)? Yes No

Does the patient have a primary diagnosis that meets the entry criteria? (Any suspected or confirmed infection, any non-infectious syndromes such as pancreatitis or burns, or other rarer conditions) See <https://genomicc.org/countries/uk/entry/> Yes No

Patients can be recruited once they become well again - please update this section as at the date patient first met eligibility.

Exclusion criteria

Has the patient ever received a bone marrow transplant? Yes No

Co-enrolment

Is the pt co-enrolled in any other clinical trials/studies? Yes No

Please complete the details for all studies that this participant is currently involved with.

Name of the study _____

Participant ID for the study _____

Name of the study _____

Participant ID for the study _____

Name of the study _____

Participant ID for the study _____

THIS PATIENT IS NOT ELIGIBLE FOR THE STUDY

This warning will appear if 'yes' has been selected to the exclusion criteria or if 'no' has been selected to 'Is the patient critically ill'.

This warning will appear if the patient has not received continuous monitoring (as selected above). Disregard for outbreaks or exposures of public health interest cohort.

THIS PATIENT MAY NOT BE ELIGIBLE FOR THE STUDY

This patient will only be eligible for the study if they are being recruited within the Outbreaks or exposures of public health interest cohort (see <https://genomicc.org/countries/uk/entry/>).

Patients from the groups specified in this cohort may be recruited even if they are not admitted to critical care.

Consent

Consent version used
(not all versions are available/valid at all sites)

5

We are now using version 5 consent documents.

Consent/assent was provided by

- Patient
- Next-of-kin
- Personal consultee [E,W,NI]
- Nominated (professional) consultee [E,W,NI]
- Welfare guardian/attorney [S]

Date of consent

Initials of person recording consent

Note - if consent was obtained from a next of kin or consultee in the first instance, please follow up consent directly with the patient if they regain capacity.

If a patient will never regain capacity or be able to consent directly, please email the GenOMICC study team to inform them.

If direct patient consent is obtained, please add a new consent page to reflect this and record the new date of consent.

If a patient does not regain capacity, they will remain in the study with the NOK/consultee consent.

Sampling Details

How was this participant recruited?

The following fields are dependent on the choices made in this section - if 'prospective recruitment' is selected, the below fields will appear.

- Prospective Recruitment (Patient is currently in ICU or still in hospital)
 Patient attended for follow-up appointment after discharge and was recruited by face-to-face discussion
 Retrospective Recruitment (Participant was contacted at home retrospectively after discharge)

Is this participant part of the ACUTE sub-study?

ACUTE is a sub-study initially recruiting at a limited number of sites. Contact GenOMICC study team if you would like more details.

Yes No

The ACUTE sub-study is a pilot running at a limited number of sites. Please only select 'yes' if your site is already taking part in the ACUTE sub-study.

Confirm sample for DNA extraction (and additional samples if this is an ACUTE patient) has/have been taken using GenOMICC sample kit, boxed and sent for posting

If 'retrospective recruitment' is selected the below fields will appear.

Which option has the participant chosen for providing a sample?

- By Third-party Research Nurse visit (Blood)
 At Hospital Out-patient appointment (Blood)
 By Self-administered saliva kit (Saliva)

Has the participant been informed that contact details (name, address, phone number, e-mail) will be passed to the Third Party blood sampling service?

Yes No

If 'By Third-party Research Nurse visit' is selected as the sampling option, the below fields will appear.

Has the participant consented to receive SMS messages from the Third Party blood sampling service as part of the process of arranging and carrying out the appointment?

Yes No

Participant Preferred Title

- Miss
- Mrs
- Ms
- Mx
- Mr
- Dr
- Professor

Participant First Name

Participant Last Name

Address Line 1

Address Line 2

Address Line 3

City

County

Country

Postcode

Contact Phone number(s). If entering multiple numbers, please separate them with commas.

Contact email

Please confirm that a sample collection kit box containing a correctly labelled EDTA tube, along with a Patient Information Sheet and copy of consent form has been packaged for posting to the participant

Please confirm that a sample collection box containing a correctly labelled Saliva collection kit, along with a Participant Information Sheet and copy of consent form has been packaged for posting to the participant

If 'By Self-administered saliva kit' is selected as the sampling option, this field will appear.

Data At Recruitment

National Audit Database ID number i.e. ICNARC number,
SICSAG number or equivalent _____

NHS or CHI number _____

If no NHS/CHI number is available,
please enter 10 zeros '0000000000'

Patient name _____

Date of birth (DD-MM-YYYY using hyphens to separate) _____

Date the patient first met the inclusion criteria
(DD-MM-YYYY using hyphens to separate)

(Date of admission to ICU (Day 0))

Date of admission to ICU (Day 0)

Primary Diagnosis

Primary diagnosis for admission to critical care
(confirmed or suspected)

- Acute hepatitis (unexplained) in children
- Acute hepatitis associated with gene therapy
- Acute pneumonia complicating confirmed infection with influenza virus
- Appendicitis
- CAR T-cell reactions
- Cholecystitis
- COVID-19
- COVID-19 MISC (Multisystem inflammatory syndrome temporally associated with COVID-19)
- Cytokine storm secondary to therapy
- ECLS
- Encephalitis
- Endocarditis
- Full thickness burns covering > 20% of body surface area
- Haemophagocytic syndrome
- Heat stroke
- Influenza
- Meningitis
- Pancreatitis of any aetiology
- Peritonitis
- Pneumonia with radiographic changes at presentation to critical care
- Pyelonephritis
- Radiation poisoning
- Reaction to vaccination
- RSV (respiratory syncytial virus) infection
- Soft tissue infections causing systemic sepsis
- Stevens-Johnson Syndrome/Toxic Epidermal Necrolysis
- Still disease
- Urinary tract infection
- Any emerging critical care syndrome not listed above
- Any primary infection not listed above
- Any reaction to therapy not listed above

Confirmed suspected or emerging infection

If 'Any primary infection not listed above' is selected, these fields will appear.

- HPAI (Highly pathogenic avian influenza)
- MERS (Middle East Respiratory Syndrome)
- SARS (Severe Acute Respiratory Syndrome)
- Ebola
- Zika virus
- Dengue
- Mpox virus
- Marburg virus
- Lassa fever virus
- Crimean-Congo Haemorrhagic Fever orthonairovirus
- Hendra henipavirus
- Nipah henipavirus
- Variola virus (minor or major)
- Bacillus anthracis
- Coxiella burnetti
- Rickettsia spp
- Yersinia pestis
- West Nile fever virus
- Yellow fever virus
- Rift Valley fever virus
- other

HAZARD - Group 3 pathogens

Please follow all local Health & Safety protocols and contact the GenOMICC Team for further details before shipping this sample

If 'Any primary infection not listed above' is selected, this warning will appear.

Percentage of body area with full thickness burns

If 'Full thickness burns' is selected, this field will appear.

Highest known transaminase (AST or ALT) assay result reported for this patient on or prior to the date of recruitment.

If 'Acute hepatitis (unexplained) in children' is selected, this field will appear.

Vaccine reaction

If 'Reaction to vaccination' is selected, these fields will appear.

- Cerebral venous sinus thrombosis
- Deep vein thrombosis/pulmonary embolism
- Other thrombotic events, with or without thrombocytopenia
- Neuroinflammatory disorders including Guillian-Barre syndrome
- Anaphylaxis
- Vasculitis
- Other potentially life-threatening suspected complications of vaccine

Additional details relating to primary diagnosis (e.g. suspected syndrome, vaccine type, therapy type, symptoms etc.)

If 'Any emerging critical care syndrome not listed above' or 'Any reaction to therapy not listed above' is selected, this field will appear.

Has the patient received invasive ventilation for their current eligible condition?

Yes No

Answering no here does not exclude a patient.

Did the patient have functionally-limiting comorbidity before this illness?
(such as heart failure, chronic obstructive pulmonary disease (COPD), or reduced exercise tolerance of any cause)?

Yes No

Answering yes here does not exclude a patient.

Did the patient have significant immunosuppression before this illness?
(such as cancer chemotherapy or acquired immune deficiency syndrome)?

Yes No

Answering yes here does not exclude a patient.

Do you think this patient's illness is unusual*?

Yes No

* answer yes if you think, in your own opinion, that this patient's illness is unusually severe, or has unusual or unexplained clinical features. Imagine if you had 2 or 3 similar patients at the same time - would you consider the possibility of a new illness or outbreak?

Demography

Age (Years) at the time when the patient first met the inclusion criteria

(If including months use decimal point and number of months, e.g. 0.3 for 3 months, 1.11 for 1 yr 11 months)

Please record the patients age at the time they met the inclusion criteria here - not their age at the time of recruitment.

Sex at birth

Male Female

Postcode

(Please provide the first half of the postcode)

Presumed Primary infection

Presumed primary infection causing critical illness
(if applicable)

- Not applicable
- SARS-CoV-2 (COVID-19)
- Acinetobacter Baumannii.
- Chlamydia pneumoniae.
- Clostridium (other)
- Dengue
- Escherichia Coli
- Enterococci
- Influenza A.
- Influenza B.
- Klebsiella pneumoniae.
- Legionella pneumophila.
- Mycoplasma pneumoniae.
- Parainfluenza species.
- Respiratory syncytial virus.
- Staphylococcus aureus
- Staphylococci (other)
- Streptococcus pneumoniae
- Group A Streptococcus / Streptococcus pyogenes
- Streptococci (other)
- Other

Presumed Primary Infection: Other - please specify _____

Note - the following pages cover day 3 and day 7 microbiology results.

All day 3 and day 7 microbiology results are required for all conditions where available (including for patients recruited with Covid).

Simply leave microbiology pages blank if there is nothing to report and mark the page as complete.

Blood culture

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Blood culture sample: Organism detected

- Staph. Aureus
- Strep. Pneumoniae
- Chlamydia pneumoniae
- Mycoplasma pneumoniae
- Klebsiella pneumoniae
- Acinetobacter Baumannii
- Other

Other - please specify _____

Date sample taken _____

Serology

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Serology sample: Organism detected

- SARS-CoV-2(COVID-19)
 Dengue
 Other

Other - please specify

Date sample taken

Urinary antigen test

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Urinary antigen sample: Organism detected

- Legionella pneumophila.
 Streptococcus pneumoniae.
 Other

Other - please specify

Date sample taken

Throat/nose swab

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Throat/nose swab sample: Organism detected

- Influenza A
- Influenza B
- Respiratory syncytial virus
- Parainfluenza species
- SARS-CoV-2(COVID-19)
- Mycoplasma pneumoniae
- Chlamydia pneumoniae
- Other

Other - please specify

Date sample taken

Tracheal aspirate

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Tracheal aspirate sample: Organism detected

- SARS-CoV-2(COVID-19)
- Staph. Aureus
- Strep. Pneumoniae
- Klebsiella pneumoniae
- Other

Other - please specify

Date sample taken

Urine Culture

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Urine culture sample: Organism detected

- E. Coli
 Enterococci
 Staph. Aureus
 Klebsiella pneumoniae
 Strep. Pneumoniae
 Other

Other - please specify

Date sample taken

Fluid from infected collection

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Fluid from infected collection sample: Organism detected

- Staph. Aureus
 Other

Other - please specify

Date sample taken

Wound swab

Only record samples taken within the first 3 calendar days from admission to ICU (day 0)

Wound swab sample: Organism detected

- Staph. Aureus
 Staphylococci (other)
 Streptococci (Group A)
 Streptococci (other)
 Clostridium (other)
 Other

Other - please specify

Date sample taken

mini-BAL

Only record samples taken within the first 7 calendar days from admission to ICU (day 0)

mini-BAL sample 1: Organism detected

- SARS-CoV-2(COVID-19)
- Staph. Aureus
- Strep. Pneumoniae
- Klebsiella pneumoniae
- Other

Other - please specify

Date sample taken

BAL

Only record samples taken within the first 7 calendar days from admission to ICU (day 0)

BAL sample: Organism detected

- SARS-CoV-2(COVID-19)
- Staph. Aureus
- Strep. Pneumoniae
- Klebsiella pneumoniae
- Other

Other - please specify

Date sample taken

Cerebrospinal Fluid

Only record samples taken within the first 7 calendar days from admission to ICU (day 0)

CSF: Organism detected

- Neisseria meningitidis
- Strep. Pneumoniae
- Haemophilus influenzae
- Klebsiella pneumoniae
- Streptococci (other)
- Listeria monocytogenes
- Other

Other - please specify

Date sample taken

CSF non-culture diagnostics

Only record samples taken within the first 7 calendar days from admission to ICU (day 0)

CSF non-culture diagnostics: Organisms detected

- Herpes simplex virus
 Herpes virus (other)
 Enterovirus
 Cryptococcus
 Mycobacterium tuberculosis
 SARS-CoV-2(COVID-19)
 Other

Other - please specify

Date sample taken

Follow Up 60 Days

Date of 60 day follow up (DD-MM-YYYY using hyphens to separate)

60 days from date patient first met criteria

60 day checker

Alive at 60 days

Date of death (DD-MM-YYYY using hyphens to separate)

Participant Change of Status

Date of Change of Status _____

Who is withdrawing the participant from the trial?

- Participant
 PI or clinical delegate
 Carer/Guardian
-

Reason for withdrawal

- Participant declined to give reason
 Other
-

Withdrawal status

- Partial Withdrawal
 Full Withdrawal

Partial Withdrawal

Data WILL continue to be updated and used for research, but no further contact will be made with the participant

Total Withdrawal

- no further contact will be made with the participant;
- data will not be updated from health records;
- data will not be removed from research that is underway or has already been done, and an audit record will be maintained to confirm participation.

Record Active Status

Confirm if this Record is inactive

Mark this record as inactive

Date of Inactivation

Reason for Inactivation

- Ineligible participant
 - Participant enrolled twice in error
 - Sample collection not completed
 - Sample not available - logistical issue
 - Sample not available - technical issue
-

This 'Record Active Status' page was introduced in May 2026.

Updating this page should only happen after discussion with the study team on a case-by-case basis.

Please don't hesitate to contact us with any questions you may have.

Email us on - genomicc@roslin.ed.ac.uk

Telephone - 0300 365 7660