

Pharmacogenetics of human immunodeficiency virus therapy

Submission date 28/05/2010	Recruitment status No longer recruiting	<input type="checkbox"/> Prospectively registered
		<input type="checkbox"/> Protocol
Registration date 28/05/2010	Overall study status Completed	<input type="checkbox"/> Statistical analysis plan
		<input checked="" type="checkbox"/> Results
Last Edited 21/01/2019	Condition category Infections and Infestations	<input type="checkbox"/> Individual participant data

Plain English summary of protocol
Not provided at time of registration

Contact information

Type(s)
Scientific

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Additional identifiers

EudraCT/CTIS number

IRAS number

ClinicalTrials.gov number

Secondary identifying numbers
7342

Study information

Scientific Title

Host genetic factors influencing drug disposition and response to human immunodeficiency virus treatment

Study objectives

This is a study to investigate the association between genetic polymorphisms and:

1. Treatment response (viral load and CD4 count), or
2. Drug exposure in human immunodeficiency virus (HIV) positive patients

The cohort study examines treatment response after starting or switching antiretroviral therapy (ART) regimen according to genotype. There is also a cross-sectional study where the primary endpoint is the measured concentration of antiviral drug. The relationship between drug exposure and genetic polymorphism will also be examined.

More details can be found here: <http://public.ukcrn.org.uk/Search/StudyDetail.aspx?StudyID=7342>

Ethics approval required

Old ethics approval format

Ethics approval(s)

North West MREC (now changed to North West 5 Research Ethics Committee), 07/11/2003, ref: 02/8/87

Study design

Multicentre non-randomised observational treatment cohort study

Primary study design

Observational

Secondary study design

Cohort study

Study setting(s)

GP practice

Study type(s)

Treatment

Participant information sheet

Health condition(s) or problem(s) studied

Topic: Infection; Subtopic: Infection (all Subtopics); Disease: Infectious diseases and microbiology

Interventions

Study A:

This study is examining treatment response after starting or switching antiretroviral therapy according to genotype with the primary endpoint of a change in CD4 count and viral load at 24 weeks.

Study B:

This study involves obtaining a single blood sample in which drug concentrations will be measured. The primary endpoint is the measurement of the antiretroviral drug.

Genomic DNA will be purified and quantified from both studies. Genetic polymorphisms will be defined by PCR-RFLP, sequence-specific PCR or SNaPshot as optimised for each allele to be examined.

Intervention Type

Other

Phase

Phase IV

Primary outcome measure

Change in CD4 count and viral load, measured at 24 weeks, with secondary endpoints of viral load at 12 weeks and time to/proportion achieving undetectable viral load

Secondary outcome measures

1. Change in viral load at 12 weeks
2. Time to/proportion achieving undetectable viral load

Overall study start date

24/04/2007

Completion date

31/12/2012

Eligibility

Key inclusion criteria

Study A:

Recruitment from existing cohort studies

Study B:

1. Aged greater than 18 years
2. Know HIV-seropositive
3. Receiving antiretroviral therapy
4. Having drug concentration measured

Participant type(s)

Patient

Age group

Adult

Lower age limit

18 Years

Sex

Both

Target number of participants

Planned sample size: 900

Key exclusion criteria

Study A:

Recruitment from existing cohort studies

Study B:

1. Aged less than 18 years

2. Not on antiretroviral therapy

Date of first enrolment

24/04/2007

Date of final enrolment

31/12/2012

Locations**Countries of recruitment**

England

United Kingdom

Study participating centre

Royal Liverpool Hospital

Liverpool

United Kingdom

L69 3GA

Sponsor information**Organisation**

University of Liverpool (UK)

Sponsor details

Thompson Yates Building

Quadrangle Brownlow Hill

Liverpool

England
United Kingdom
L69 3GB
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research@rlbuht.nhs.uk

Sponsor type
University/education

Website
<http://www.liv.ac.uk/>

Organisation
Royal Liverpool and Broadgreen University Hospitals NHS Trust (UK)

Sponsor details
Prescot Street
Liverpool
England
United Kingdom
L7 8XP

Sponsor type
Hospital/treatment centre

Organisation
University of Liverpool

Sponsor details

Sponsor type
Not defined

Website
<http://www.liv.ac.uk/>

ROR
<https://ror.org/04xs57h96>

Funder(s)

Funder type
Research council

Funder Name

Medical Research Council (MRC) (UK)

Alternative Name(s)

Medical Research Council (United Kingdom), UK Medical Research Council, MRC

Funding Body Type

Government organisation

Funding Body Subtype

National government

Location

United Kingdom

Results and Publications

Publication and dissemination plan

Not provided at time of registration

Intention to publish date**Individual participant data (IPD) sharing plan****IPD sharing plan summary**

Not provided at time of registration

Study outputs

Output type	Details	Date created	Date added	Peer reviewed?	Patient-facing?
Results article	results	01/11/2008	21/01/2019	Yes	No
Results article	results	01/02/2014	21/01/2019	Yes	No
Results article	results	01/05/2009	21/01/2019	Yes	No
Results article	results of population pharmacokinetic modeling of the association between 63396C->T pregnane X receptor polymorphism and unboosted atazanavir clearance	01/12/2010	21/01/2019	Yes	No
Results article	results of the association of ABCC10 polymorphisms with nevirapine plasma concentrations	01/01/2012	21/01/2019	Yes	No
Results article	results of the effect of SLCO1B1 polymorphisms on lopinavir plasma concentration in HIV-infected adults	01/02/2012	21/01/2019	Yes	No
Results article	results of the effects of SNPs within OATP1A2, OATP1B1 and OATP1B3 on the pharmacokinetics of lopinavir	01/02/2010	21/01/2019	Yes	No