Tailoring bespoke medicine using big genetics and genomics data in the NHS

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NIHR Biomedical Research Centre: Building a Learning Health System

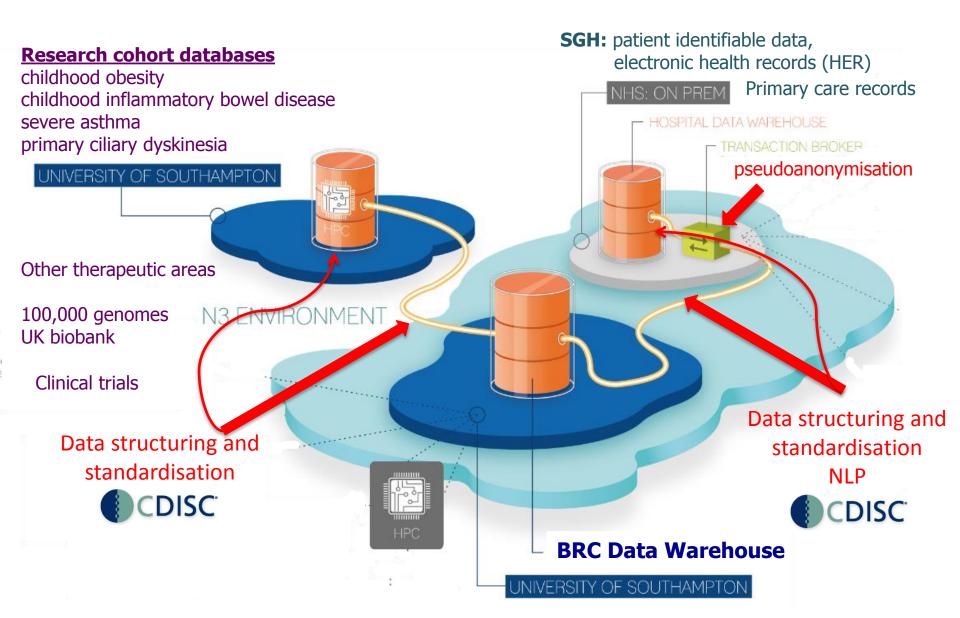
Aims:

- 1. Integrate high-dimensional e-health data such as imaging, electrograms, activity trackers and molecular profiles, so as to implement the advances of stratified medicine.
- 2. Use patient-relevant data from across the hospital and university to improve clinical care by datadriven decision making
- 3. Refine phenomarker profiles by iterative interrogation of patient data and enable replication and validation studies

To do this, we must

- 1. Get all the data together in a secure environment (BRC Datawarehouse) with structure, standardisation and normalisation
- 2. Interrogate it in the context of known and putative phenomarkers for stratified medicine.
- 3. Feed the results of these analyses back into the data and clinical care.

Where the data comes from



Hurdles to integrated data

Data safety and security

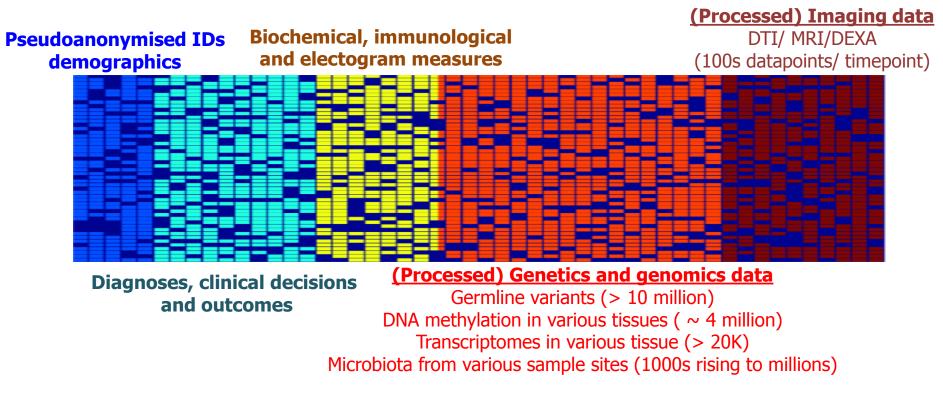
- N3 compliant but need HPC facility too.
- Data warehouse will be pseudoanonymised but there will be the ability to map back to patient identifiers.

• Some is unstructured and all must be standardised

- Apply Naturel language processing and standardise to CDISC
- Feedback structure and standards to individual databases
- Data cleaning may be necessary, have to evaluate and apply *ad hoc*
- Patient consent for use of clinical data in research and new contacts with subject for further investigations
 - Comprehensive ethical framework with pro-active consenting of SGH patients.

What the data will look like

Virtual data matrix: rows are patients, columns are variables



Nextgen: Activity trackers and other mobile connected apps

- Short and wide -> multiple testing problems, over-fitting problems
- Some data types will have high degree of missingness and/or associated confidence metrics
- Complex structure of longitudinal and/or hierachical relations e.g. DNA->RNA

How do we want to interrogate it?

Replicate

Many phenomarker signatures have already been reported with varying degrees of confidence in translation to clinical decision making. We can make these readily available to clinicians and use more data to improve on them.

Discover new patterns

Define diseases, disease subclasses and prognosis by supplementing symptom data with high-dimensional data.

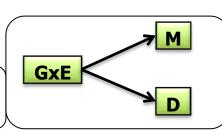
Make new predictions

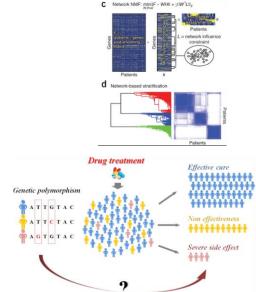
Predict response to treatments given full dataset for patient (includes pharmacogenomics)

Learn causal relationships

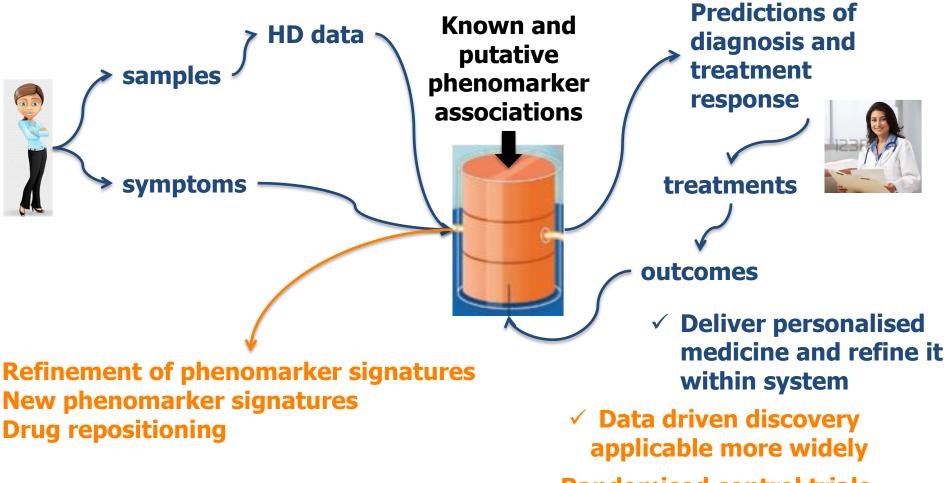
Longitudinal data analysis and causal inference (MR), to unpick casual relationships between phenomarkers and disease







Where we are going: What a true learning health system looks like



Replication in other health systems (Other Trusts and BRCs) Randomised control trials Select patients with particular profiles from data warehouse

We are not alone

Imperial College London





The Farr Institute of Health Informatics

NHS

NHS

National Institute for Health Research

National Institute for

Health Research

Data Science Institute









PRECISION MEDICINE INITIATIVE COHORT PROGRAM

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