Balancing the Local and the Universal in Maintaining Ethical Access to a Genomics Biobank

Catherine Heeney Science Technology and Innovation Studies & Shona Kerr MRC Institute of Genetics and Molecular Medicine Edinburgh University

## Oppositions in data sharing

- Open access v bureaucracy/ownership
- Standardisation v local arrangements
- Public good v protection of data subject autonomy
- Broad consent v restrictive access agreements
- Platforms/consortia v repositories/studies

## The em

 Huma •Fort L (Well •UK Bid •The W Conso

Biostatistics (2010), 11, 4, pp. 661–673 doi:10.1093/biostatistics/kxq035 Advance Access publication on June 3, 2010

### On inferring presence of an individual in a mixture: a Bayesian approach

### DAVID CLAYTON

Wellcome Trust/Juvenile Diabetes Research Foundation, Diabetes and Inflammation Laboratory and Department of Medical Genetics, Cambridge Institute for Medical Research, Cambridge University, Wellcome Trust/MRC Building, Addenbrooke's Hospital, Hills Road, Cambridge CB2 0XY, UK david.clayton@cimr.cam.ac.uk

### SUMMARY

Homer and others (2008. Resolving individuals contributing trace amounts of DNA to highly complex mixtures using high-density SNP genotyping microarrays. *PLoS Genetics* 4, e1000167) recently showed that, given allele frequency data for a large number of single nucleotide polymorphisms in a sample together with corresponding population "reference" frequencies, by typing an individual's DNA sample at the same set of loci it can be inferred whether or not the individual was a member of the sample. This observation has been responsible for precautionary removal of large amounts of summary data from public access. This and further work on the problem has followed a frequentist approach. This paper sets out a Bayesian analysis of this problem which clarifies the role of the reference frequencies and allows incorporation of prior probabilities of the individual's membership in the sample.

Downloaded from http://biostatistics.ord/ordjournals.org/atEdinbutg

## Openess continued...

- Global Alliance for Genomics and Health
  - Harmonisation, setting universal standards for governance (Knoppers et al 2014) (Birney et al 2017)
- RCUK Concordat on Open Research Data (2016)
  - All data
- The FAIR Guiding Principles for scientific data management and stewardship (2016)
  - Findable, Accessible, Interoperable and Reusable (Wilkinson et al 2016)

## Broad Consent v Managed access

- Toronto Statement acknowledges 'access may be restricted' in circumstances where detailed genomic or clinical data pose a risk of deidentification of individuals research subjects (Toronto Statement 2009)
- Consent 'is not a panacea...robust governance is essential for the ethical conduct of research' (WT Expert Advisory Group on Data Access 2015).
- RCUK Principle 5 reasons for not sharing: commercial interests and again the privacy and confidentiality of research subjects - (RCUK 2016)

# The Scottish Place

- Expert Working Groups and the GS Executive Committee
- Access Committee
  - Scientific merit
  - Governance
  - Data/Material
  - Sustainability

 Co-authorship, which is stated in the data and materials transfer agreement and the GS Authorship & Acknowledgement Policy

## Sustainability and governance

- Adequate acknowledgement of those involved in maintaining the study - authorship
- Cost recovery through access fees helping to maintain the governance and curation of the GS resource
- Original researchers ability to fulfil commitments given 'moral distance' between new and original contexts for data use and sharing (Bull, Roberts and Parker 2015)
- Implications of requests for data to be housed on platforms such as MRC Dementias Platform UK (<u>http://www.dementiasplatform.uk/</u>)?
  - Creating 'tremendous consent challenges' (Caulfield et al 2008)





CSH) Laboratory bioRxiv	Search		
THE PREPRINT SERVER FOR BIOLOGY	Ad	Ivanced Sea	
New Results	O Previous		
Balancing the Local and the Universal in Maintaining Ethical Ac	ess to a Posted August 16, 2017.		
Genomics Biobank			
	Download PDF	Share	
Catherine Heeney, Shona M. Kerr	🖬 Email	Citation Too	
Ioi: https://doi.org/10.1101/157024 This article is a preprint and has not been peer-reviewed [what does this mean?].			
nis arocie is a preprint and has not been peer-reviewed [what does this mean;].	Tweet Like 0	-	
Abstract Info/History Metrics	Preview PDF Subject Area		
	Genomics		
Abstract			
Issues of balancing data accessibility with ethical considerations and genomics research biobank, Generation Scotland, are explored within	Subject Areas		
policy landscape of the past ten years. During this time data sharing			
access have become increasingly important topics in biomedical rese	Animal Behavior and Cognition	6	
around data access are influenced by local arrangements for govern	Biochemistry		
such as linkage to health records, and the global through policies fo	Bioenzineering	Bioengineering	
the sharing of data with large-scale biomedical research data resource	Bioinformatics	Bioinformatics	
Methods: We use a literature review of policy relevant documents wh	Biophysics		
conduct of biobanks in two areas: support for open access and the p	Cancer Biology		
subjects and researchers managing a bioresource. We present exam	Cell Biology	Cell Biology	
making within a biobank based upon observations of the Generation	Clinical Trials	Clinical Trials	
Committee. We reflect upon how the drive towards open access raise	ethical		

A Real and A Real Property of the local states of the local states

#### Abstract