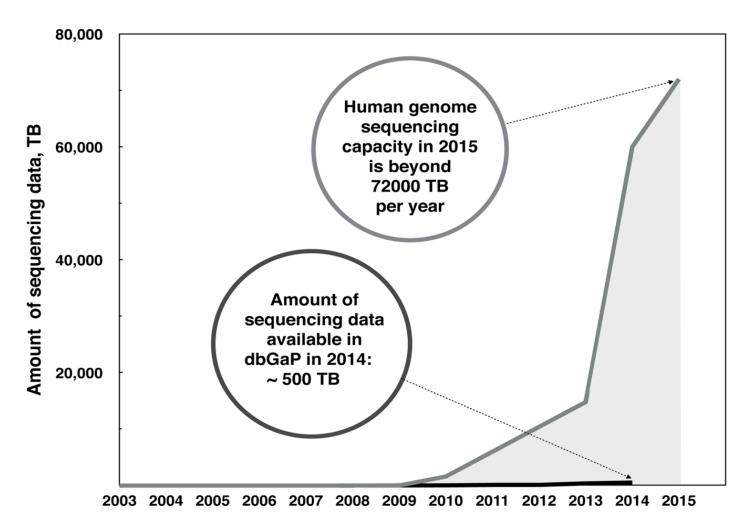
人類基因資料的開放與管理



何之行 中研院歐美所

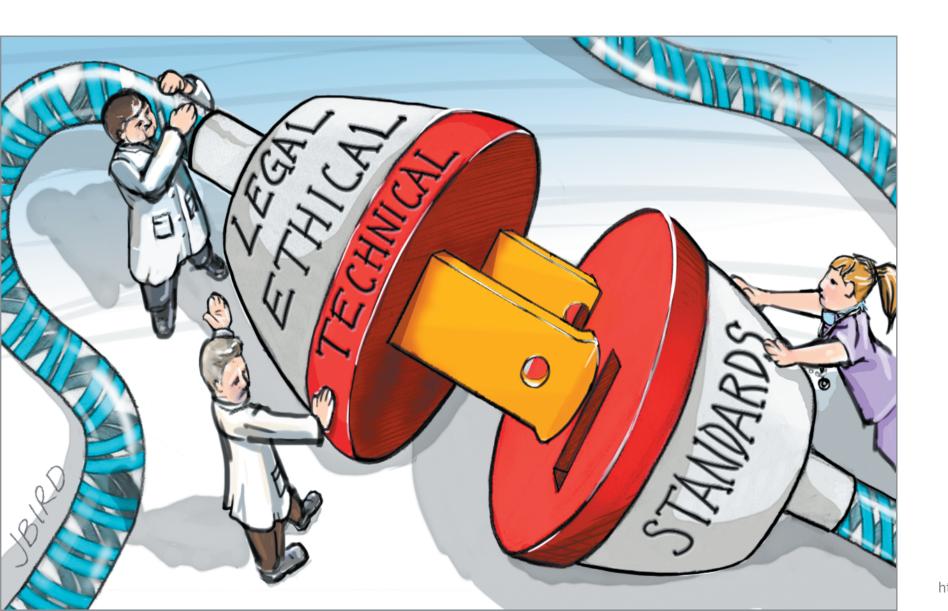
Email: <u>chihho@sinica.edu.tw</u> 2018 研究資料管理工作坊

Vast amount of genomic data is produced



The Database of Genotypes and Phenotypes (dbGaP) started to operate from 2007

Why is data not shared?



Barriers

- Legal
- Ethical
- Technical

Difficulties

FRAGMENTED

Poor visibility of available genomic data

ADMIN BURDEN

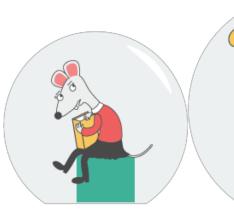
Huge overhead to manage data access

CULTURE OF SHARING

Lack of data sharing habits in research culture









Fiona Nielsen: Human Genomic Data Discoverability, Data Dialogue, Cambridge – July 28th 2016

National Institutes of Health (NIH) calls for a new Genomic Data Sharing Policy

US Federal Government Initiatives

- ➤ Proposed Common Rule revisions (July 2011) supports broad consent to maximize utility of bio-specimens and data
- February 2013 White House initiative to increase access to publications and the results of federally funded scientific research
- ➤ NIH Big Data to Knowledge Initiative (BD2K) supports the broad use of digital assets and resources to enhance the utility of biomedical big data and accelerate discovery

Global Alliance for Genomics and Health (GA4GH)

a policy-framing and technical standards-setting organization, seeking to enable responsible genomic data sharing

international, nonprofit alliance formed in 2013





The Comparison of Bermuda Principle (1996), and the GA4GH Genomic Data Sharing Policy 1.

International Human Genome Sequencing Consortium (Bermuda 1996)	Global Alliance for Genomics and Health (2017 and beyond)
Number of countries	
5	>70
Sources of data	
Major sequencing laboratories	Research laboratories of many types, clinical laboratories

The Comparison of Bermuda Principle (1996), and the GA4GH Genomic Data Sharing Policy II.

Type of data	
International Human Genome Sequencing Consortium (Bermuda 1996)	Global Alliance for Genomics and Health (2017 and beyond)
Assembled fragments of DNA sequence data contributing to an ultimate reference sequence	Data on genomic variants, clinical laboratory data, medical outcomes data, electronic medical records, exposure to environmental factors and toxins
Key diff	erences
broader array of data types; more databases with diverse formats/organisations	

The Comparison of Bermuda Principle (1996), and the GA4GH Genomic Data Sharing Policy III.

International Human Genome Sequencing Consortium (Bermuda 1996) Global Alliance for Genomics and Health (2017 and beyond)

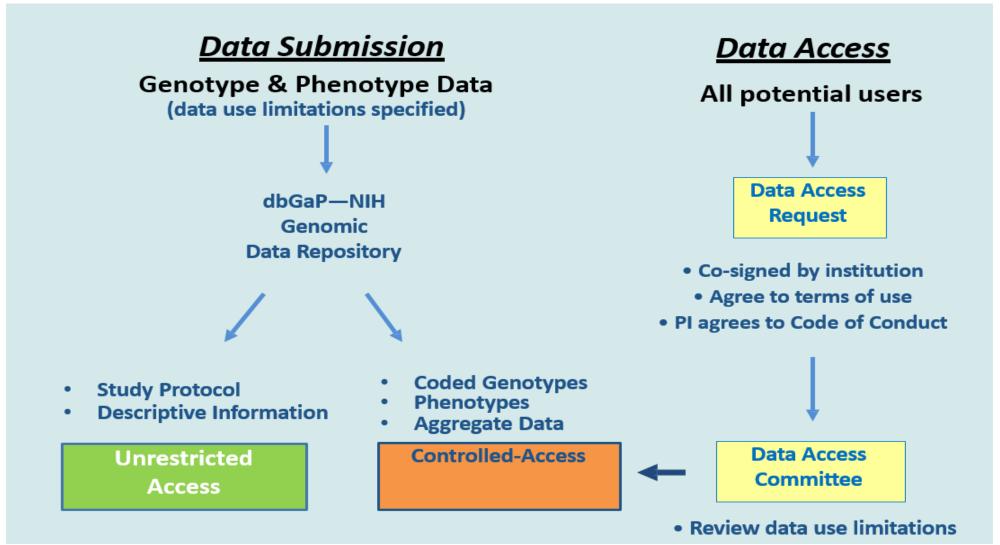
Regulatory Framework: Consent, Privacy, confidentiality

Reference samples from a small number of donors who gave informed consent

- Sequencing technology can enable re-identification,
- -Standards for informed consent are changing,
- more **complex legal framework** for sharing personal data

Robert Cook-Deegan and Amy L. McGuire: Moving beyond Bermuda: Sharing data to build

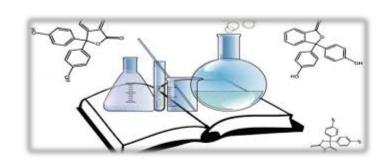
NIH Data Submission and Access



NIH's Genomic Data Sharing Policy Overview https://osp.od.nih.gov/scientific-sharing/genomic-data-sharing/

Repositories for Human Genomic Data

- Unrestricted-access: data are accessible to anyone via public website (previously referred to as "open access")
 - e.g., 1000 Genomes project
- Controlled-access: data are available to an investigator for a specific project only if certain stipulations are met
 - e.g., dbGaP, Cancer Genomics Hub



NIH Data Sharing Policy Informed Consent Expectations I.

PI should ask **participant's consent** for genomic and phenotypic data to be used for future research purposes and to be shared **broadly**

- Explanation as to whether the data will be shared via unrestricted- or controlled-access repositories
- If participant does not consent to broad sharing of data, he or she may still be enrolled in the study, but the data may not be shared



NIH Data Sharing Policy Informed Consent Expectations II.

For studies using <u>cell lines or clinical specimens</u> created or collected after the effective date:

- Informed consent for future research use and broad data sharing should have been obtained, even if samples are de-identified (studies initiated after Jan 2015)
- Exception to consent expectation can be granted, if there are compelling scientific reasons

`Compelling scientific reasons`

No guidance on this exemption yet, but the NIH promised it

`Further guidance on what constitutes compelling scientific reasons will be made available on the GDS website and will likely evolve over time`

Terms and Conditions for Research Use of Controlled-Access Data

- The data can be used only for approved research

it is prohibited

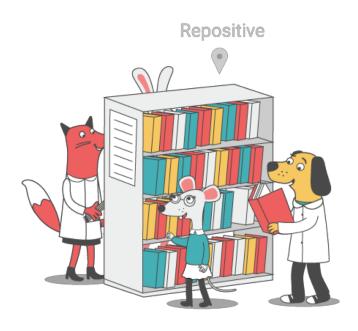
- to re-identify individual participants from whom the data were obtained
- to sell any of the data obtained from NIH-designated data repositories
- sharing any of the data obtained from controlled-access NIH-designated data

FAIR data: Findable, Accessible, Interoperable, Reusable

MAKE DATA
DISCOVERABLE

SIMPLIFY WORKFLOWS

CONTRIBUTE TO COMMUNITY







Fiona Nielsen: Human Genomic Data Discoverability, Data Dialogue, Cambridge – July 28th 2016

Thank you for your attention!

