

# National Institute of Health 'All of Us' Research Program

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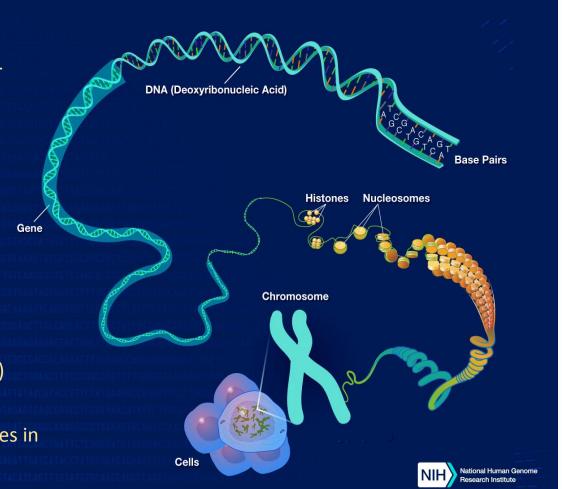
## Genomics is the study of the code of life

• DNA is composed of A-T and C-G base pairs similar to 0s and 1s in a computer

• Everyone has a unique DNA sequence

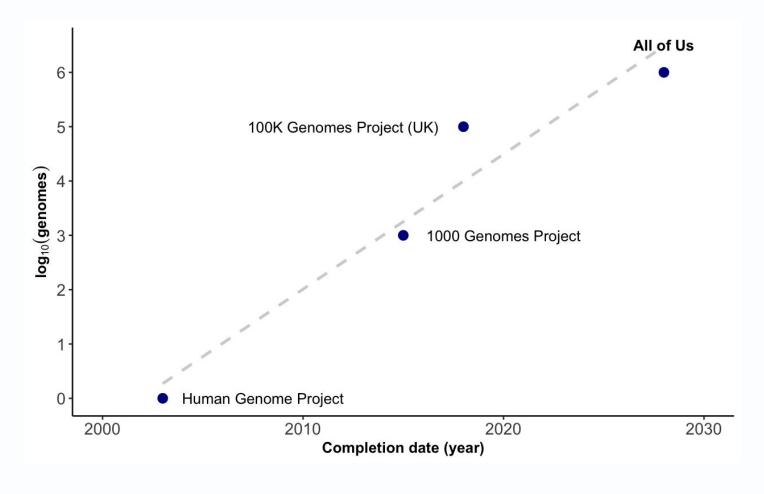


- Errors in the DNA sequence (ie, "bugs") can be fatal or cause disease
- 'All of Us' aims to understand differences in our genomes and improve medicine





## Increasing scale of genomics projects



Abbreviation: UK, United Kingdom.



## **Data curation**

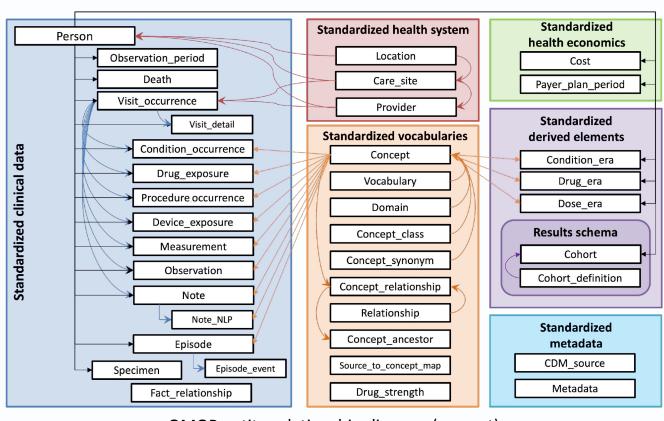


- Data sources include patient surveys, electronic health records, genomic sequences
- Data are harmonized using a common data model

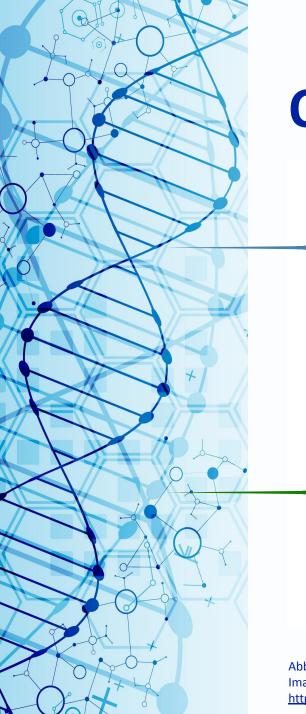


## Observational Medical Outcomes Partnership (OMOP) Common Data Model

- Open community standard for observational data from hospitals and insurance providers
  - Common format (data model)
  - Common representation (standardized vocabulary)
  - Library of standard analytic tools
- Implemented in SQL,
  R, and Python



OMOP entity-relationship diagram (excerpt)



## **OMOP Data Harmonization Process**



Data quality checks



Qualitative data quality analysis



#### **ETL-process**

Save OMOP data in target database



Convert source format to OMOP CDM



Map source vocabulary to OMOP CDM







Define scope of source data for use(s)





Assess source data structure, format

Vocabulary identification



## Coverage analysis of vocabularies



Assess extent that source vocabulary already exists in OMOP; identify missing vocabularies



### **Questions?**

For more information about the 'All of Us' research program, visit <a href="https://www.joinallofus.org/">https://www.joinallofus.org/</a>