BIODS215

Topics in Biomedical Data Science: Large-scale inference

Winter Quarter 2018

Course Instructors

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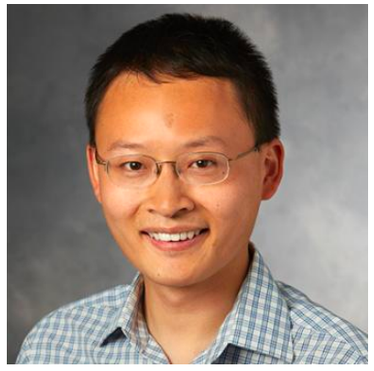

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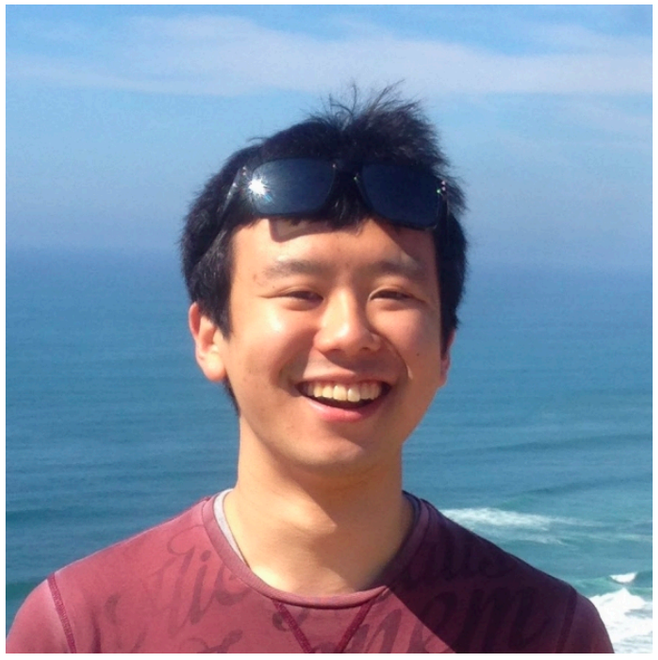
Teaching Assistant



Yosuke Tanigawa
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rivaslab.stanford.edu

Lecture structure

~20-45 minutes motivating biomedical example

~30-55 minutes statistical inference concept lecture

5-7 minute break in the middle

Course requirements and grading

Two homework assignments (40%)

Final project (50%)

Class participation (10%)

Announcements from TA

- Canvas access
  If you don't have access to Canvas, please see Yosuke

- Gradescope for assignments
  https://gradescope.com/
  Entry code: MWYKE4

- TA Office hour
  12 Jan. (Fri.) 10-11am at MSOB x399
  Tuesdays 1:30-2:30pm at MSOB x393

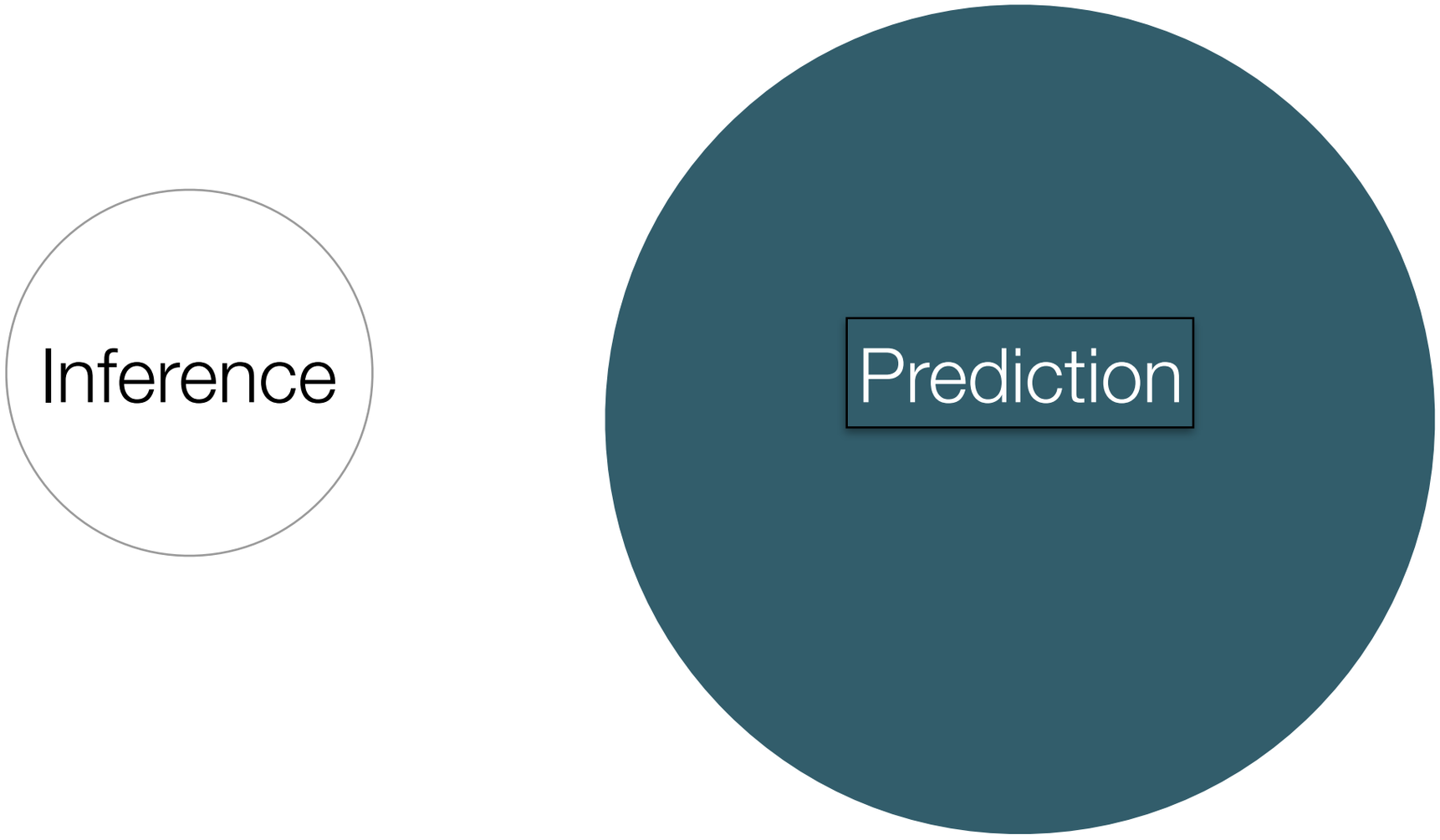
RIVASLAB

Data explosion and worldview across fields

Inference: To [infer] how nature is associating the response variables to the input variables

Statisticians, Biomedicine (therapeutics)

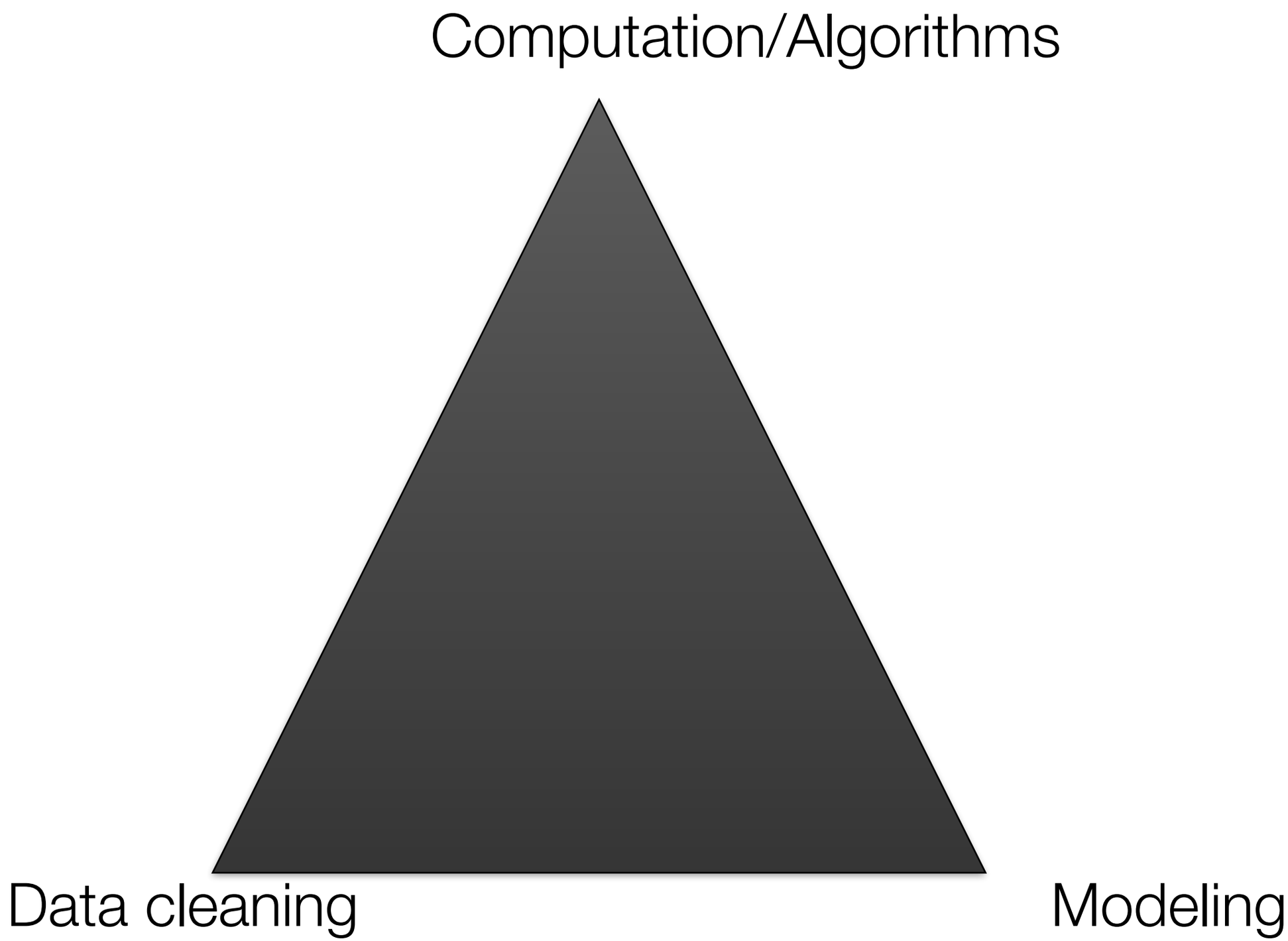
Data explosion and worldview across fields



Prediction: To be able to predict what the responses are going to be to future input variables

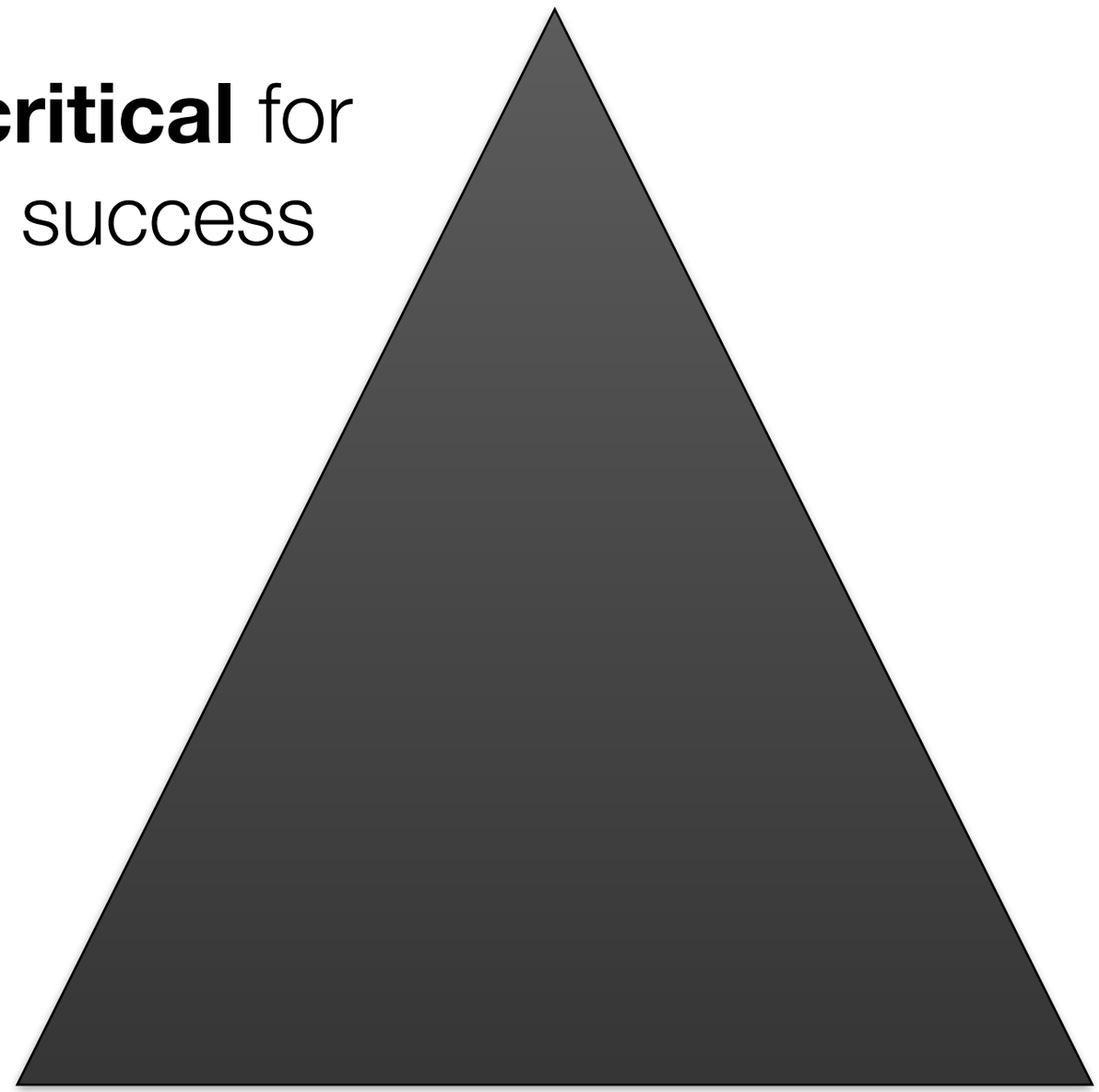
Machine Learning, Computer science

Learning objectives



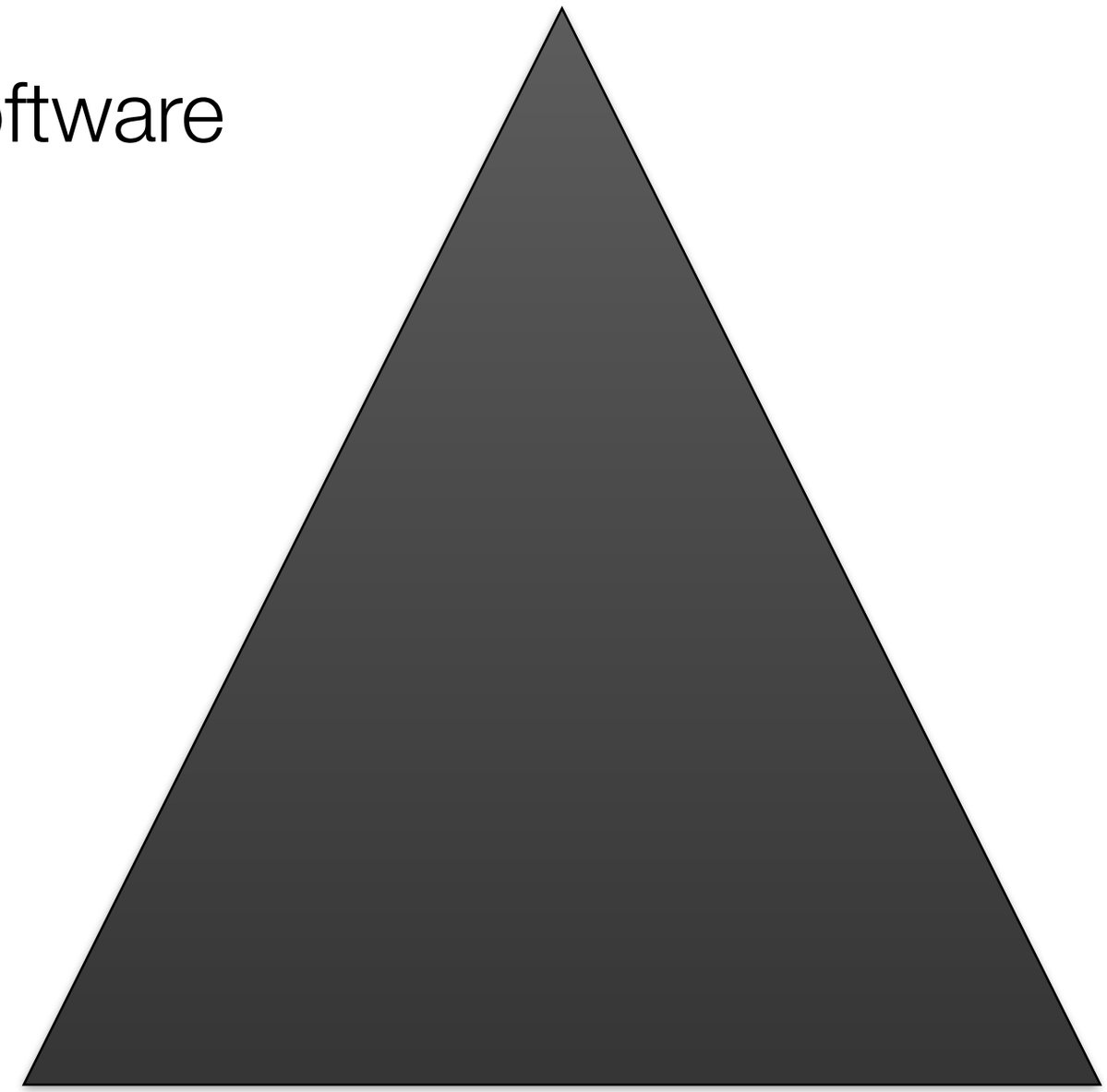
Learning objectives

Not fun, but **critical** for downstream success



Learning objectives

Algorithms: software and hardware



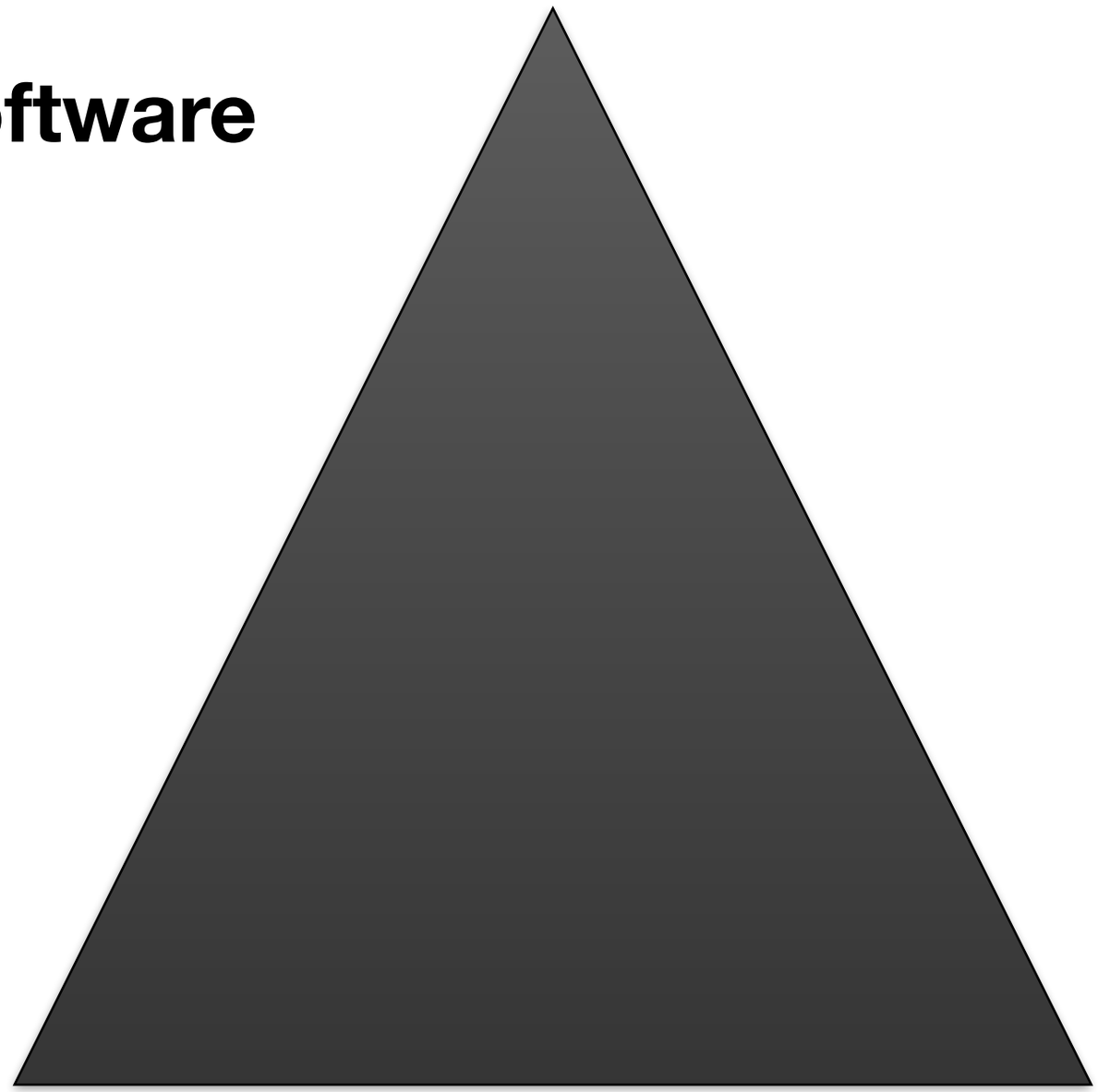
Data cleaning

Modeling

Learning objectives

Algorithms: **software** and hardware

Computation/Algorithms

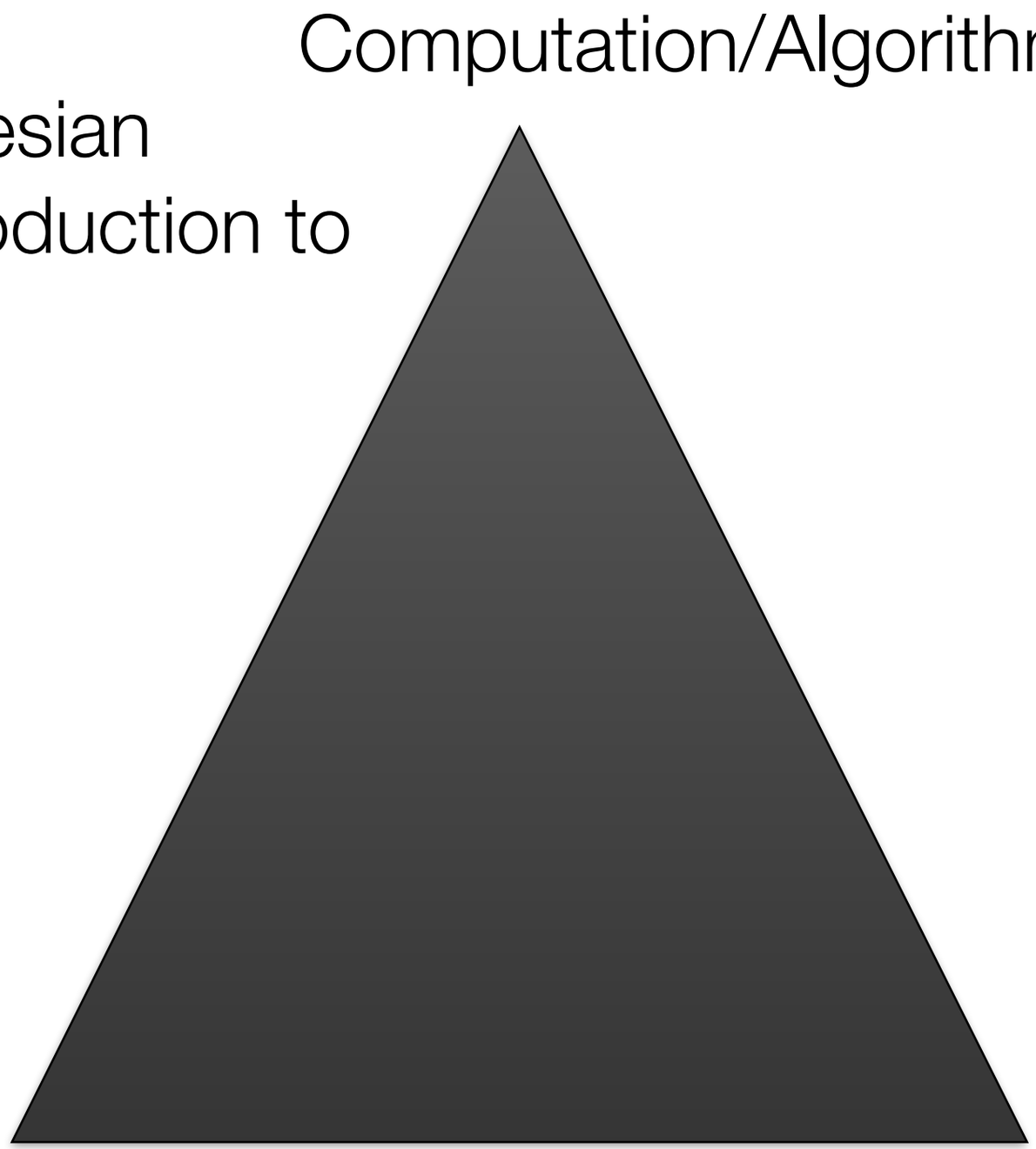


Data cleaning

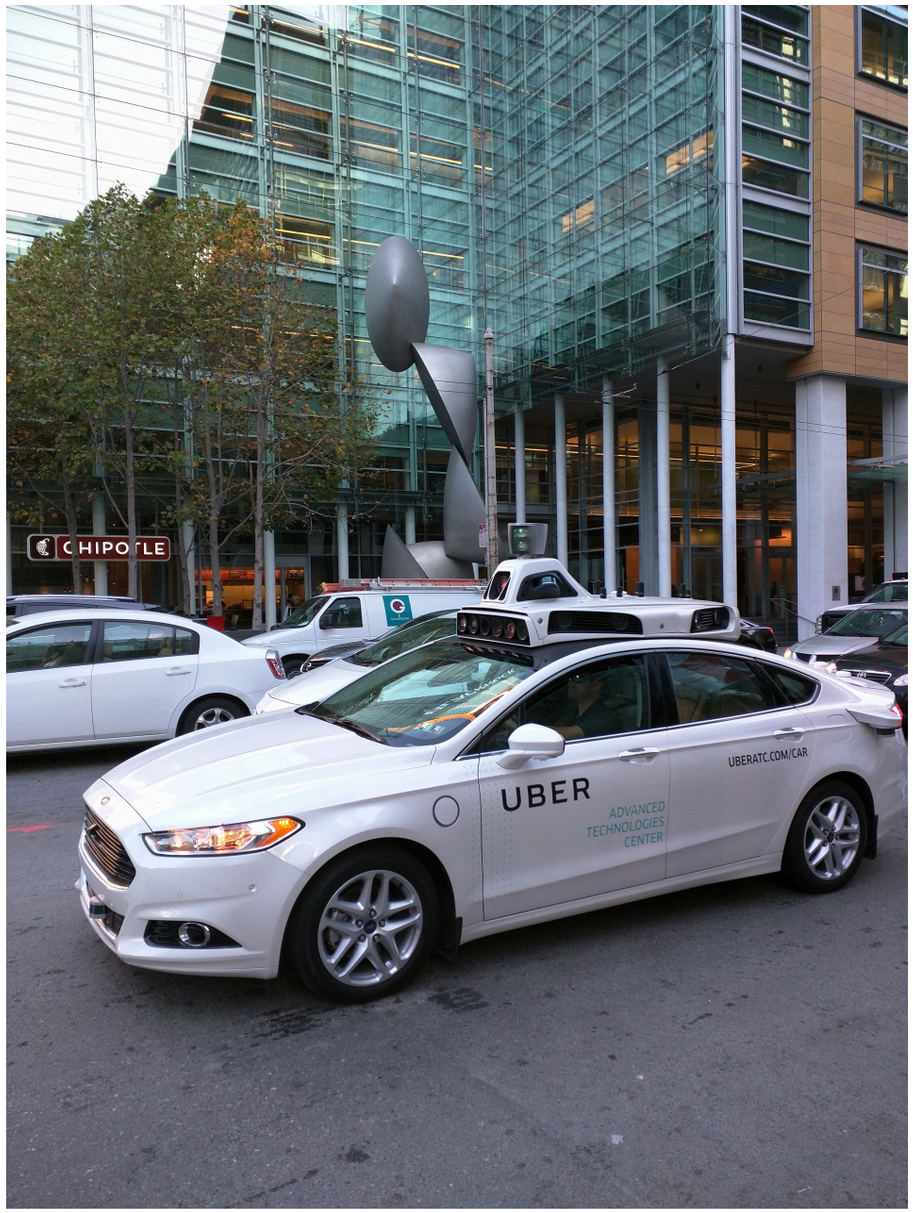
Modeling

Learning objectives

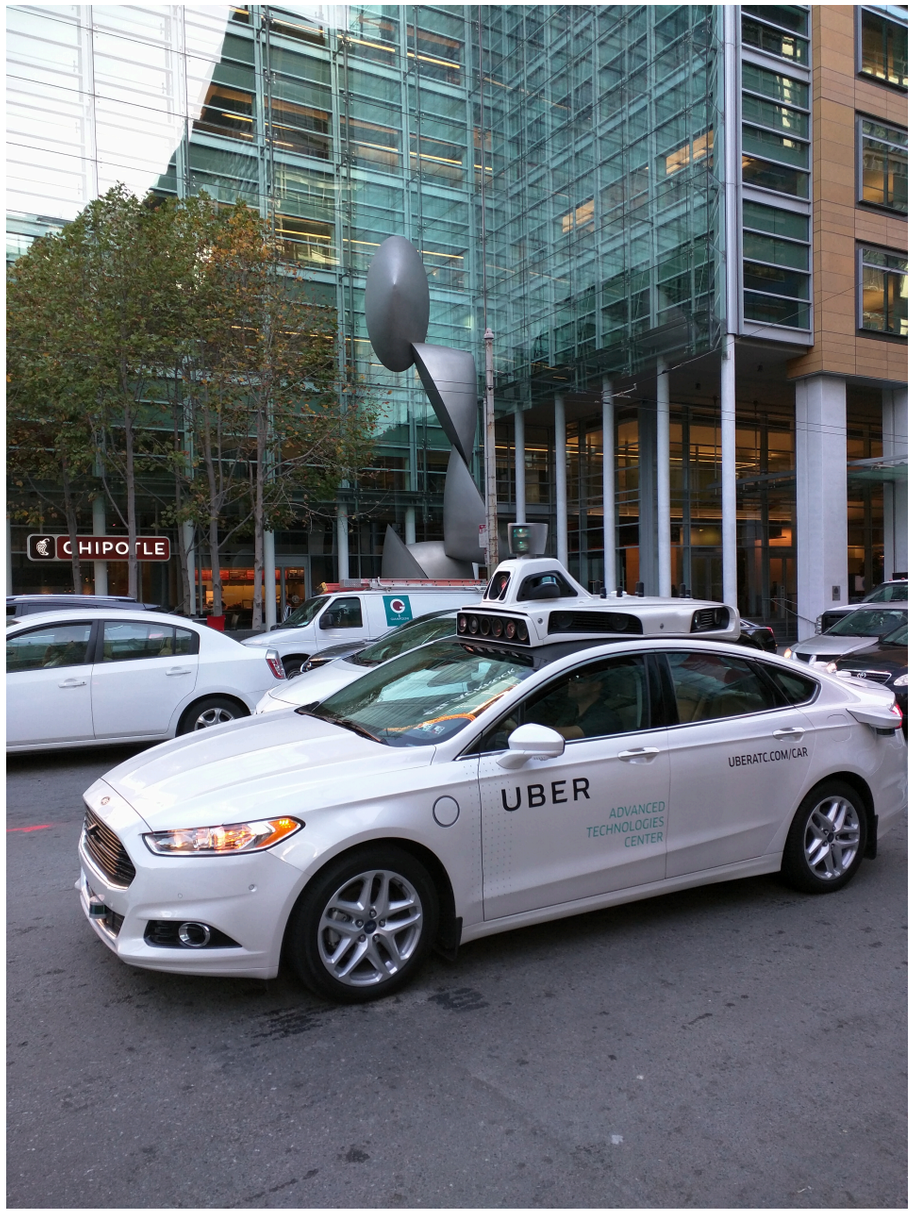
Focus on Bayesian modeling, introduction to deep learning



Transformation of many industries



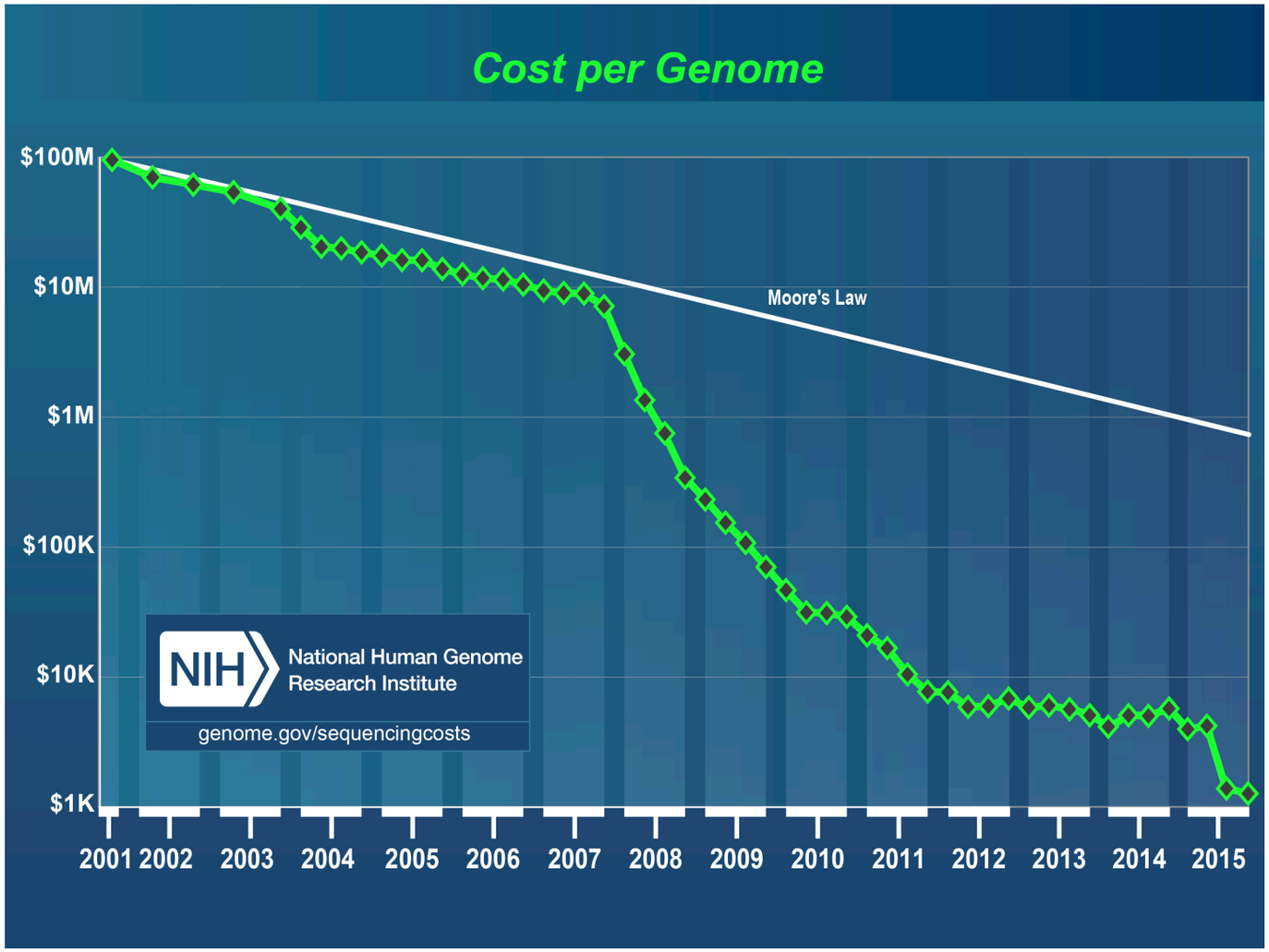
Transformation of many industries



What is missing?

Technologies transforming biomedicine

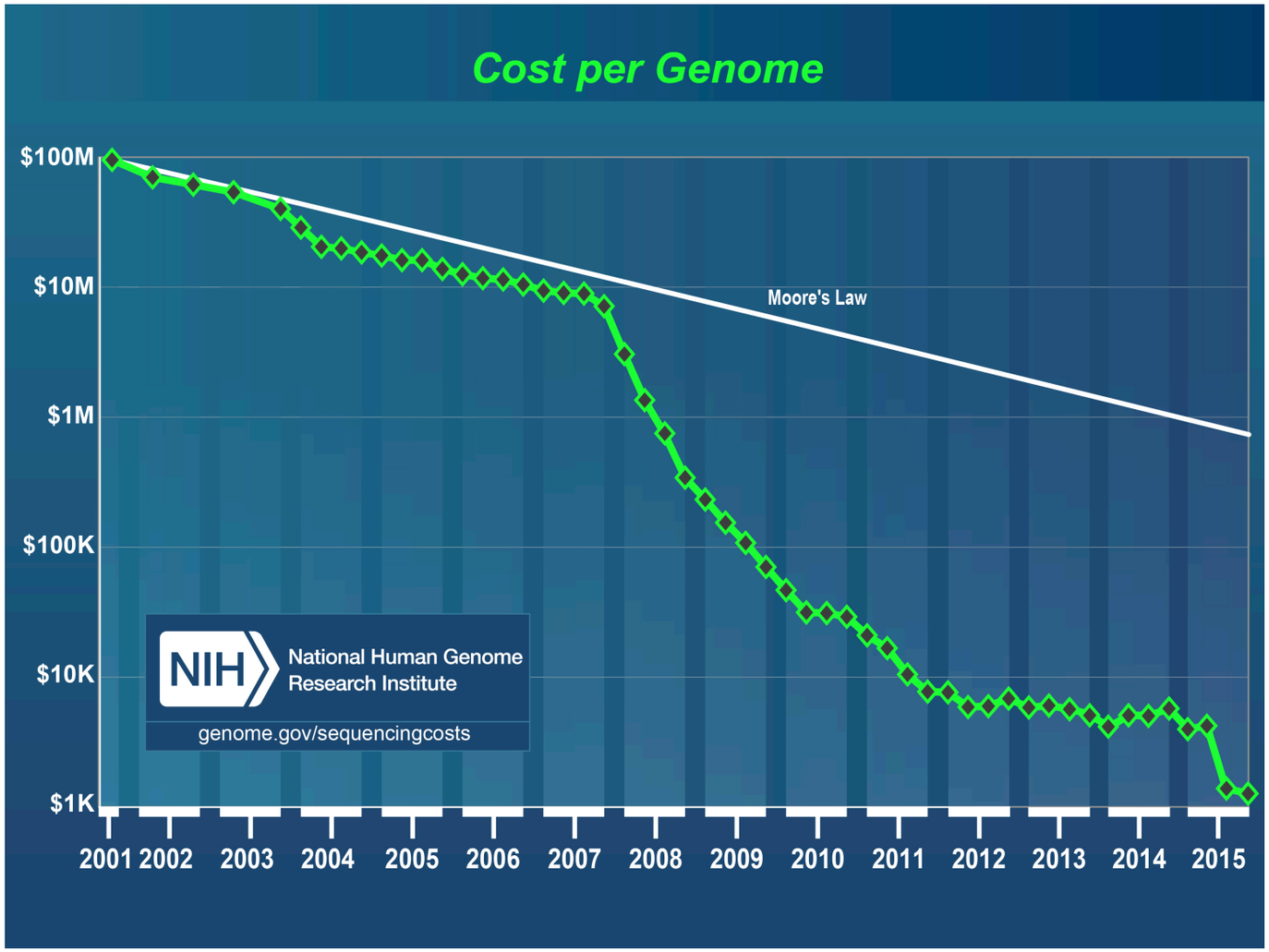
Cost of sequencing has plummeted over the past 15 years



Technologies transforming biomedicine

Cost of sequencing has plummeted over the past 15 years

~$1000 cost point projected for 18/19



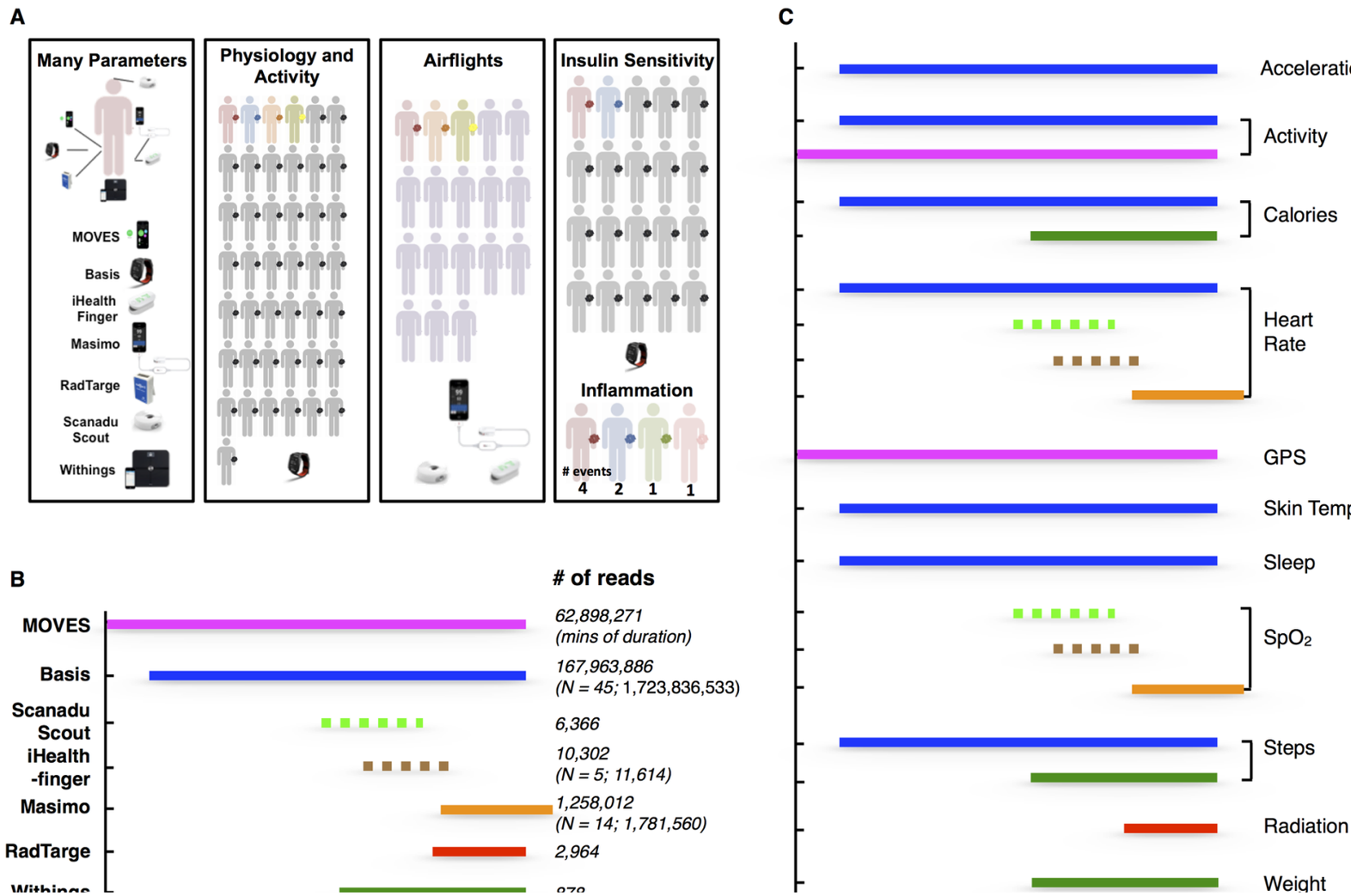
Technologies transforming biomedicine

Wearables and sensors

Ability to continuously monitor health measurements

Technologies transforming biomedicine

Li et al. 2017, PLoS Biology



Digital Health: Tracking Physiomes and Activity Using Wearable Biosensors Reveals Useful Health-Related Information

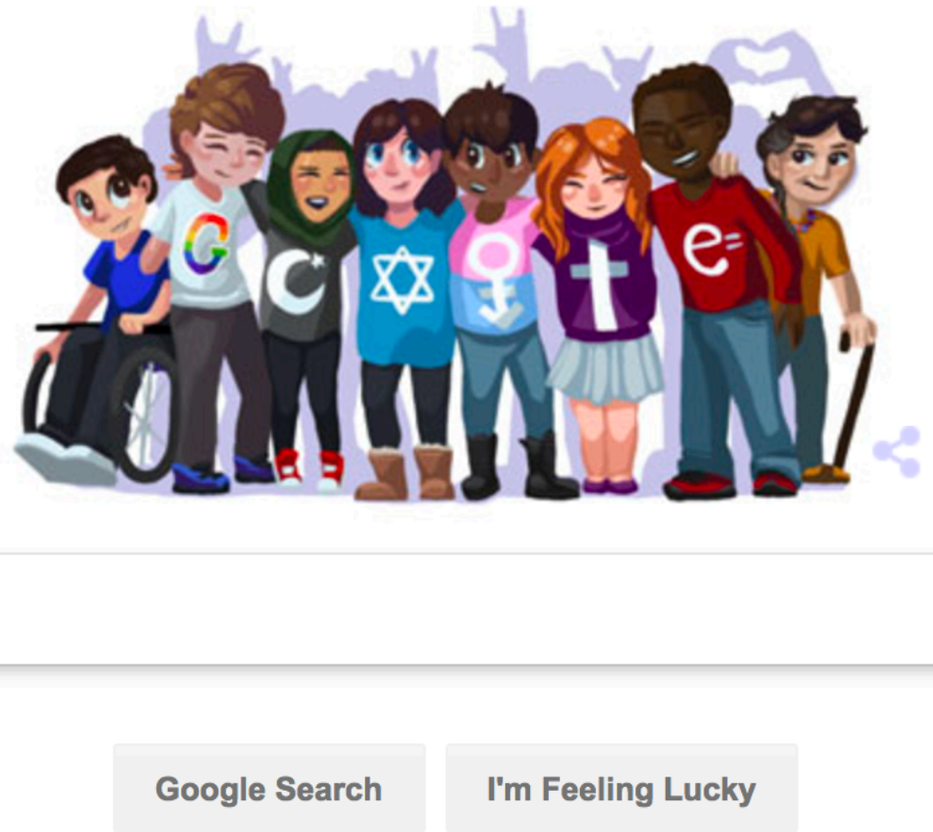
Technologies transforming biomedicine

Data streams from individuals participating in social networks

Social network data

Technologies transforming biomedicine

Data streams
from individual's
search activity



Search engine data

Technologies transforming biomedicine

Google

Detecting influenza epidemics using search engine query data

Jeremy Ginsberg[1], Matthew H. Mohebbi[1], Rajan S. Patel[1], Lynnette Brammer[2], Mark S. Smolinski[1] & Larry Brilliant[1]

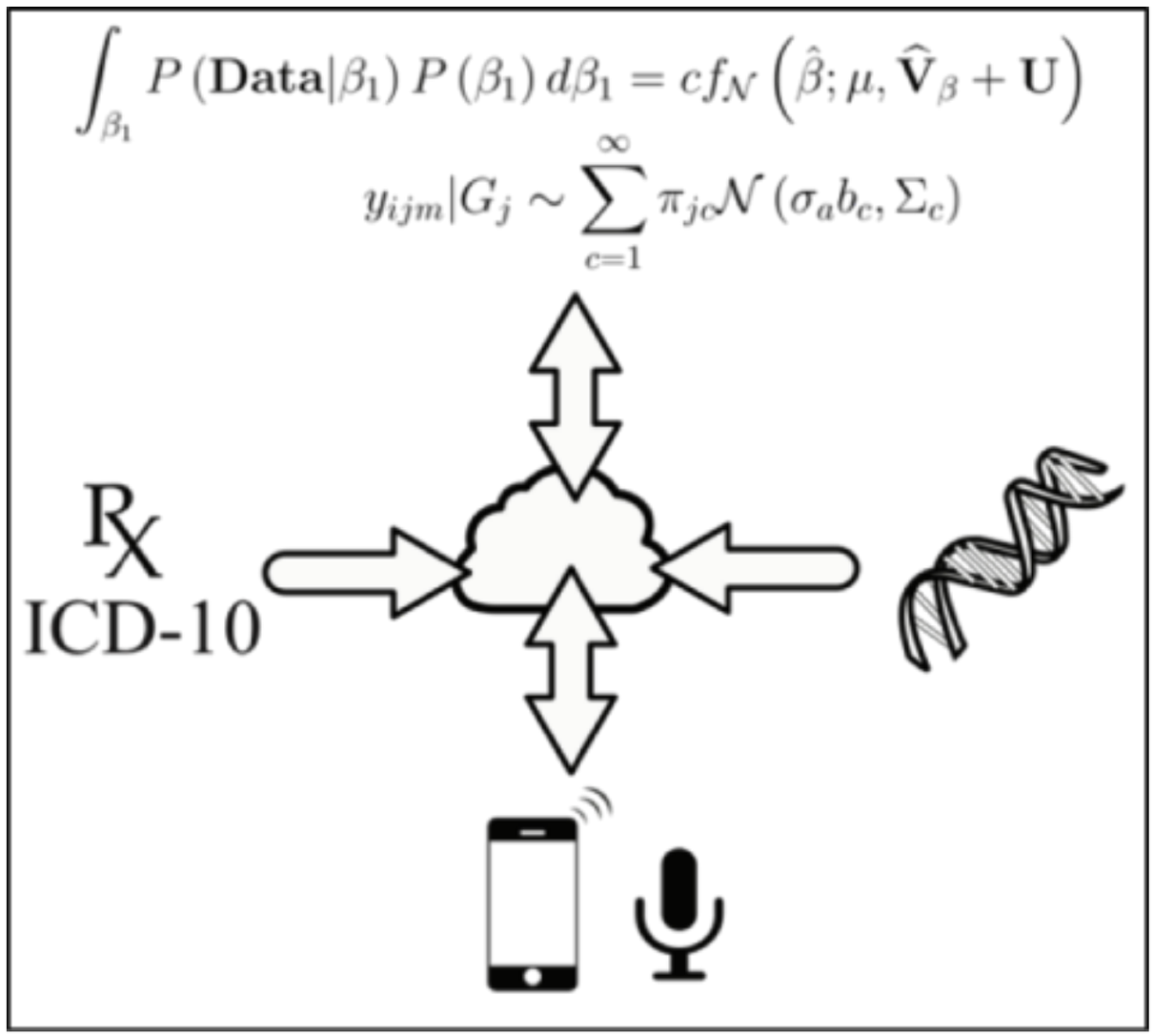
[1]Google Inc. [2]Centers for Disease Control and Prevention

Search engine data

State of health records in many places

Old technologies - health records across many regions of the world are annotated in pencil and paper

How to digitize and put data into action?



Challenge for this generation

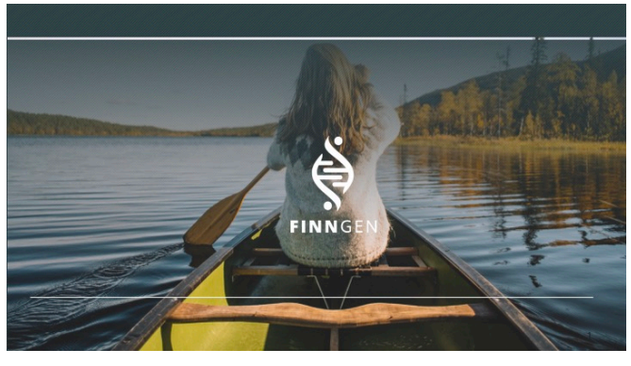
Precision health and biobank initiatives

Precision health and biobank initiatives

UK Biobank

China Kadoorie Biobank

Precision Medicine Initiative



FinnGen

Introduction to the UK Biobank project

Major source of data for this course

About the UK Biobank

National and international health resource

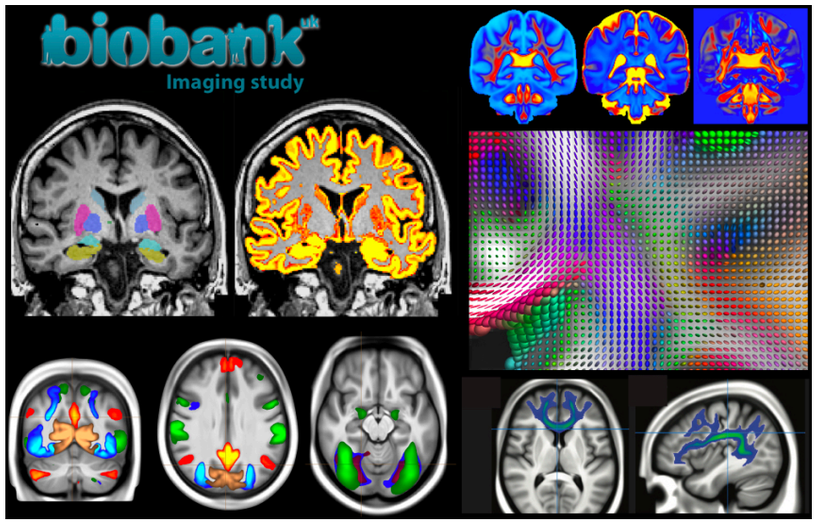
About the UK Biobank

National and international health resource

Hospital in-patient record

Primary care clinical notes

About the UK Biobank



National and international health resource

Hospital in-patient record

Primary care clinical notes

Imaging

~10,000 individuals -> 100,000

About the UK Biobank

National and international health resource

Hospital in-patient record

Primary care clinical notes

Imaging

Physical activity

About the UK Biobank

National and international health resource

Hospital in-patient record

Primary care clinical notes

Imaging

Physical activity

Biomarkers, etc

UK Biobank data showcase webpage

http://biobank.ctsu.ox.ac.uk/crystal/

Please visit

UK Biobank data showcase webpage

biobank uk

Index | Browse | Search | Catalogues | Downloads | Help

Welcome to the online showcase of UK Biobank resources. If you are new to using the showcase we recommend you begin by reading the short introductory **User Guide**. Please note that the showcase contains only anonymous summary information.

Essential Information

Information regarding timelines, updates, release schedules etc.

Browse

Find data items by navigating according to their category of origin.

Search

Find data items by searching on keywords and other characteristics.

Catalogues

Simple listings of database contents and additional resources.

Downloads

Download supporting utilities.

Login

Request data access and view cross-tabulations.

Improving the health of future generations

UK Biobank data showcase webpage

biobank uk

Index | Browse | Search | Catalogues | Downloads | Help

Browse by Primary Category of Origin

Category	Items
Population characteristics	8
UK Biobank Assessment Centre	2023
Biological samples	184
Genomics	12
Genotyping process	6
Genotyping intensities	27
Genotype confidences	25
Genotype calls & imputation	26
Online follow-up	466
Additional exposures	221
Health-related outcomes	149
Returned datasets	1

Top Level

Level 1

Level 2

Level 3

Summary generated 4 February 2017

Improving the health of future generations

UK Biobank data showcase webpage

biobank uk

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Browse by Primary Category of Origin

Category	Items
Population characteristics	8
UK Biobank Assessment Centre	0
Recruitment	13
Touchscreen	385
Verbal interview	31
Physical measures	396
Cognitive function	69
Imaging	1108
Biological sampling	10
Procedural metrics	11
Biological samples	184
Genomics	96
Online follow-up	466
Additional exposures	221
Health-related outcomes	149
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Top Level

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UK Biobank Assessment Centre	2023
Biological samples	184
Genomics	96
Online follow-up	466
Additional exposures	221
Health-related outcomes	0
Hospital in-patient	121
Death register	6
Cancer register	8
Algorithmically-defined outcomes	14
Returned datasets	1

Top Level

Level 1

Level 2

Level 3

Summary generated 4 February 2017

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Category	Items
Population characteristics	8
UK Biobank Assessment Centre	2023
Biological samples	184
Genomics	96
Online follow-up	0
Diet by 24-hour recall	317
Cognitive function follow-up	48
Work environment	101
Mental health	0
Additional exposures	221
Health-related outcomes	149
Returned datasets	1

Summary generated 4 February 2017

Top Level

Level 1

Level 2

Level 3

Improving the health of future generations

UK Biobank announcement

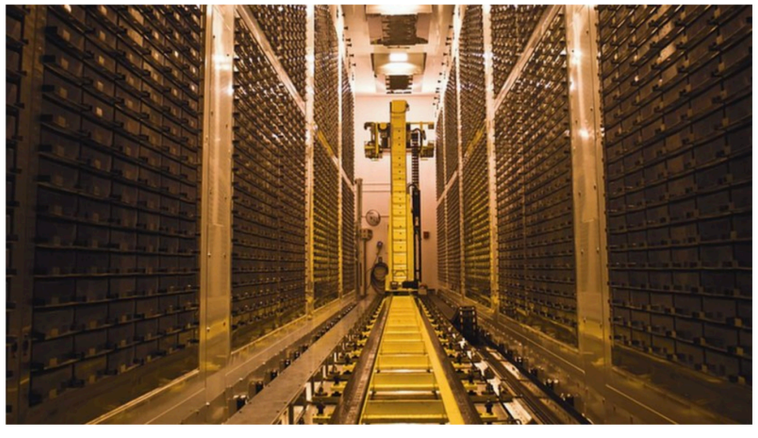
Drug Company Consortium To Sequence The Genes Of 500,000 Britons Over Next Two Years

Matthew Herper, FORBES STAFF
I cover science and medicine, and believe this is biology's century. FULL BIO

Adi Gaskell, Contributor
A London based innovation scout

Medical Consortium Aim To Find Treasure In UK Biobank Data

01/09/2018 02:55 am ET



The power of genetics is something that I've touched on a number of times. Technology

Rewriting Life

500,000 Britons' Genomes Will Be Public by 2020, Transforming Drug Research

Yancopoulos calls the slow start by the U.S. "a national embarrassment." The U.K. data trove is set to dominate "for the foreseeable future, the next five to 10 years," he says. "It's going to be the best resource. It's the first place people will go."

Rare diseases run in families

If you have **cystic fibrosis**, what is your risk for:

Your neighbor (unrelated)?

Your sibling?

Your twin?

Variation in your DNA influences your risk

Common diseases also run in families

If you have **cystic fibrosis**, what is your risk for:

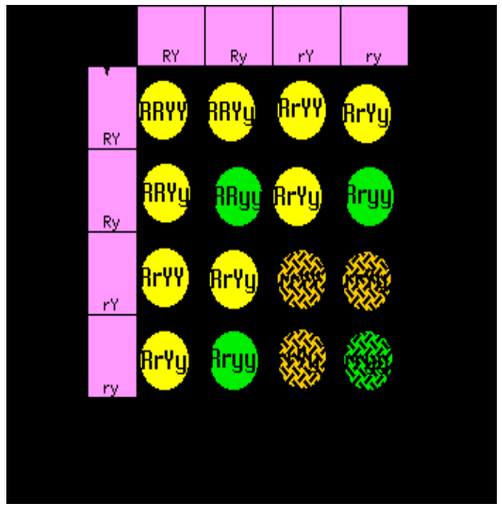
Your neighbor (unrelated)?

Your sibling?

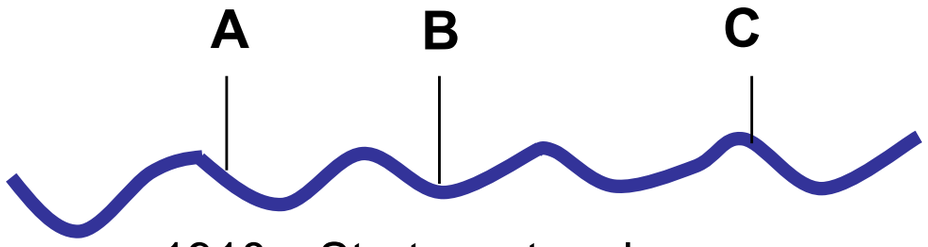
Your twin?

Variation in your DNA influences your risk

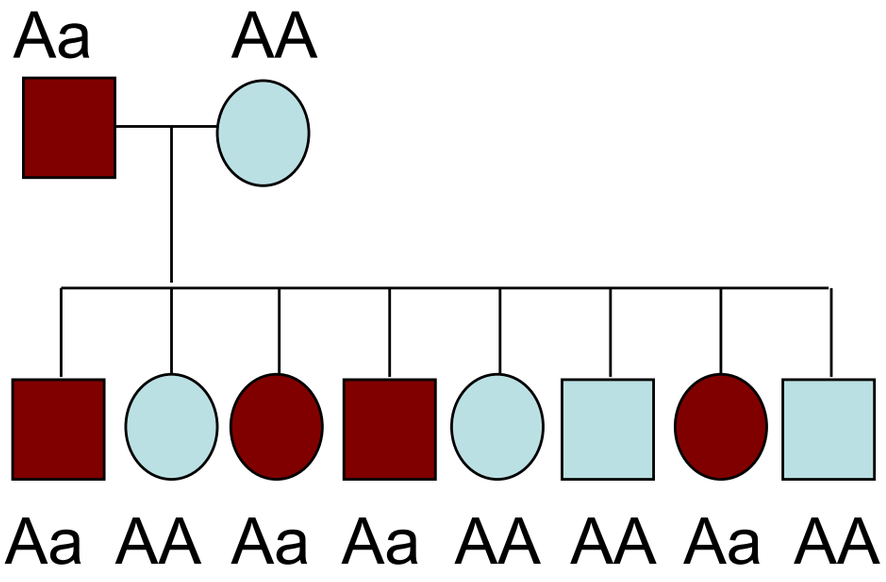
20th century genetics



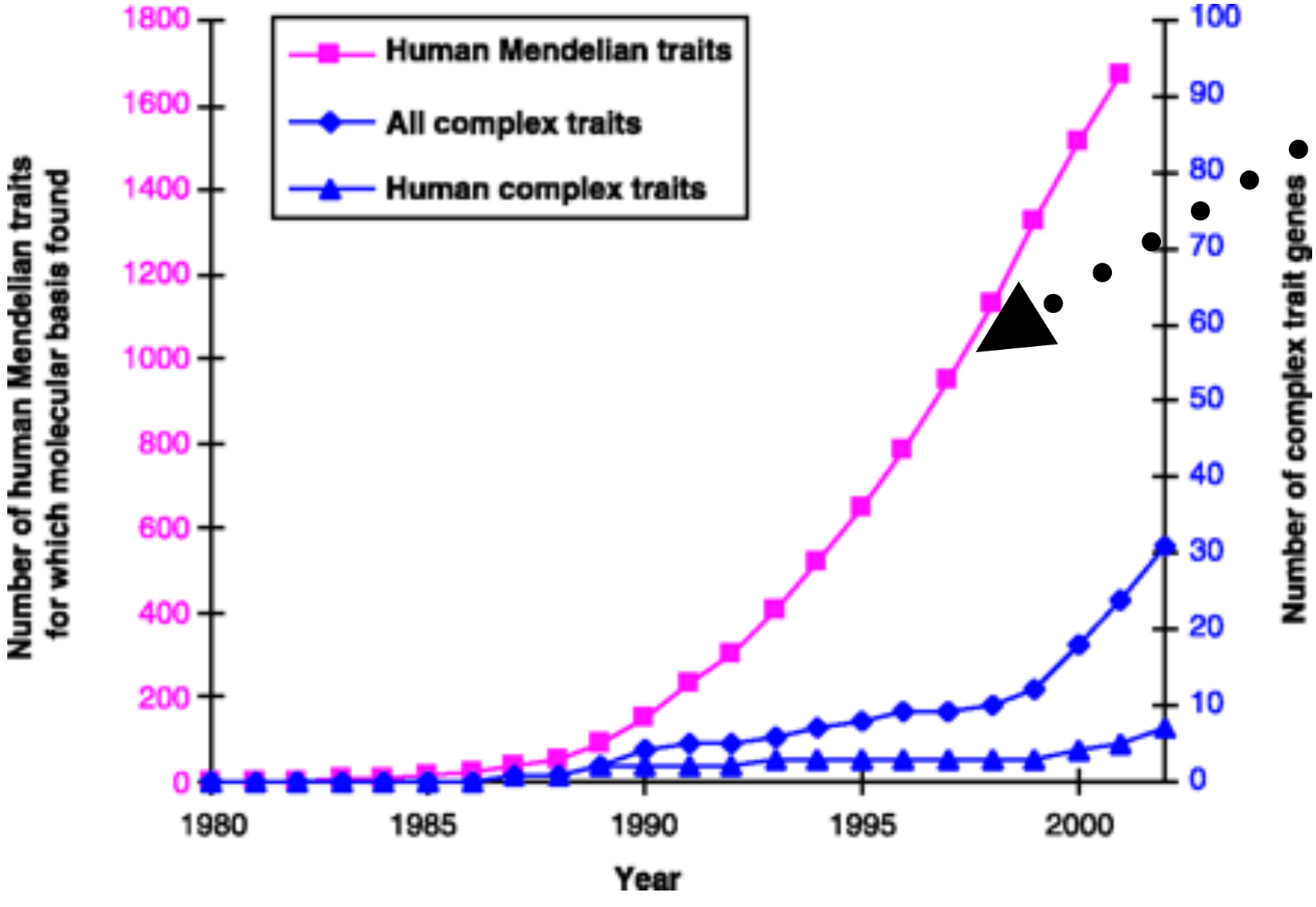
1860s: Mendel's laws of inheritance – discrete, transmissible units of inherited variation resulting in phenotypic differences



1910s: Sturtevant and Morgan create the first genetic map

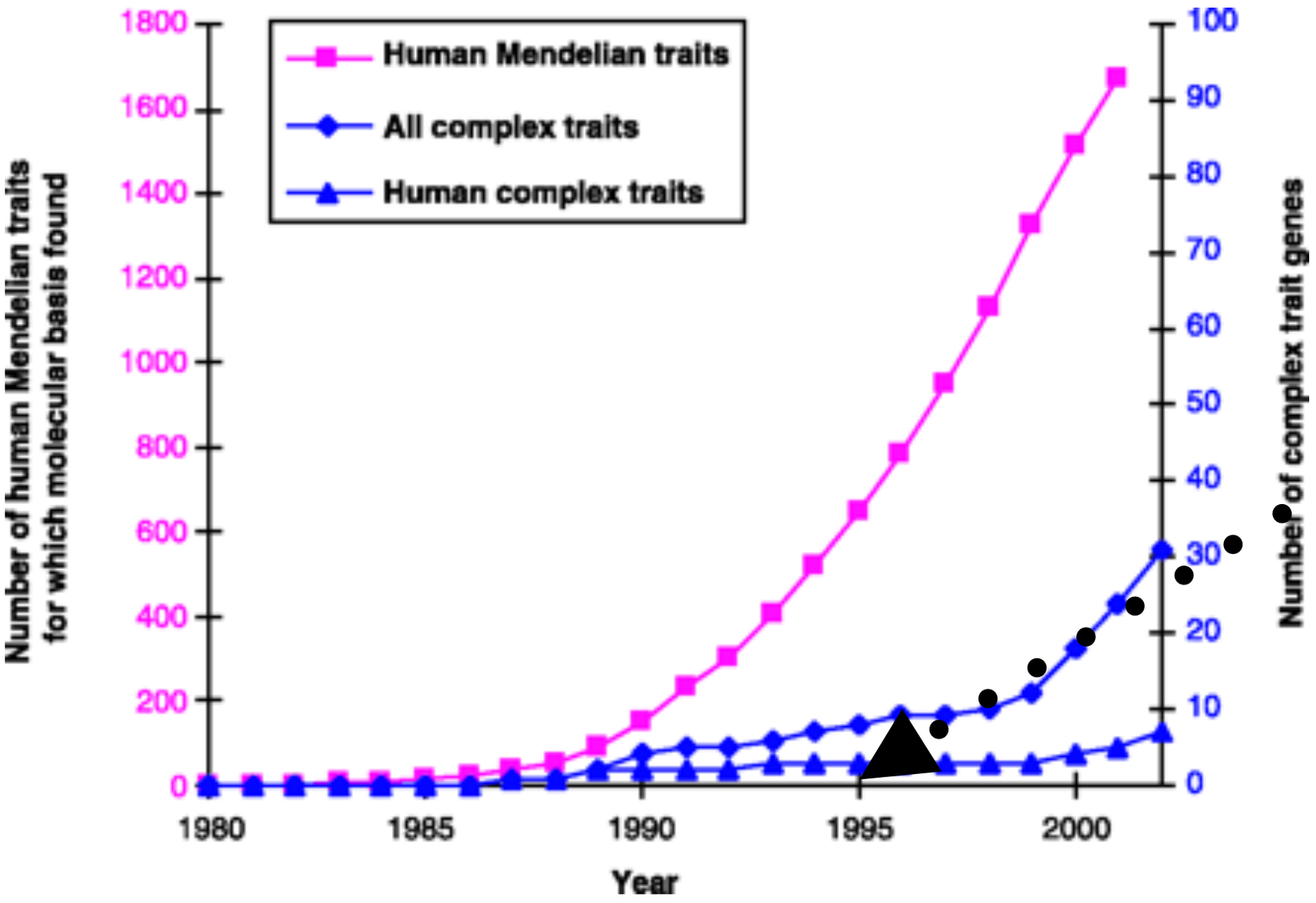


Dark ages of human genetics



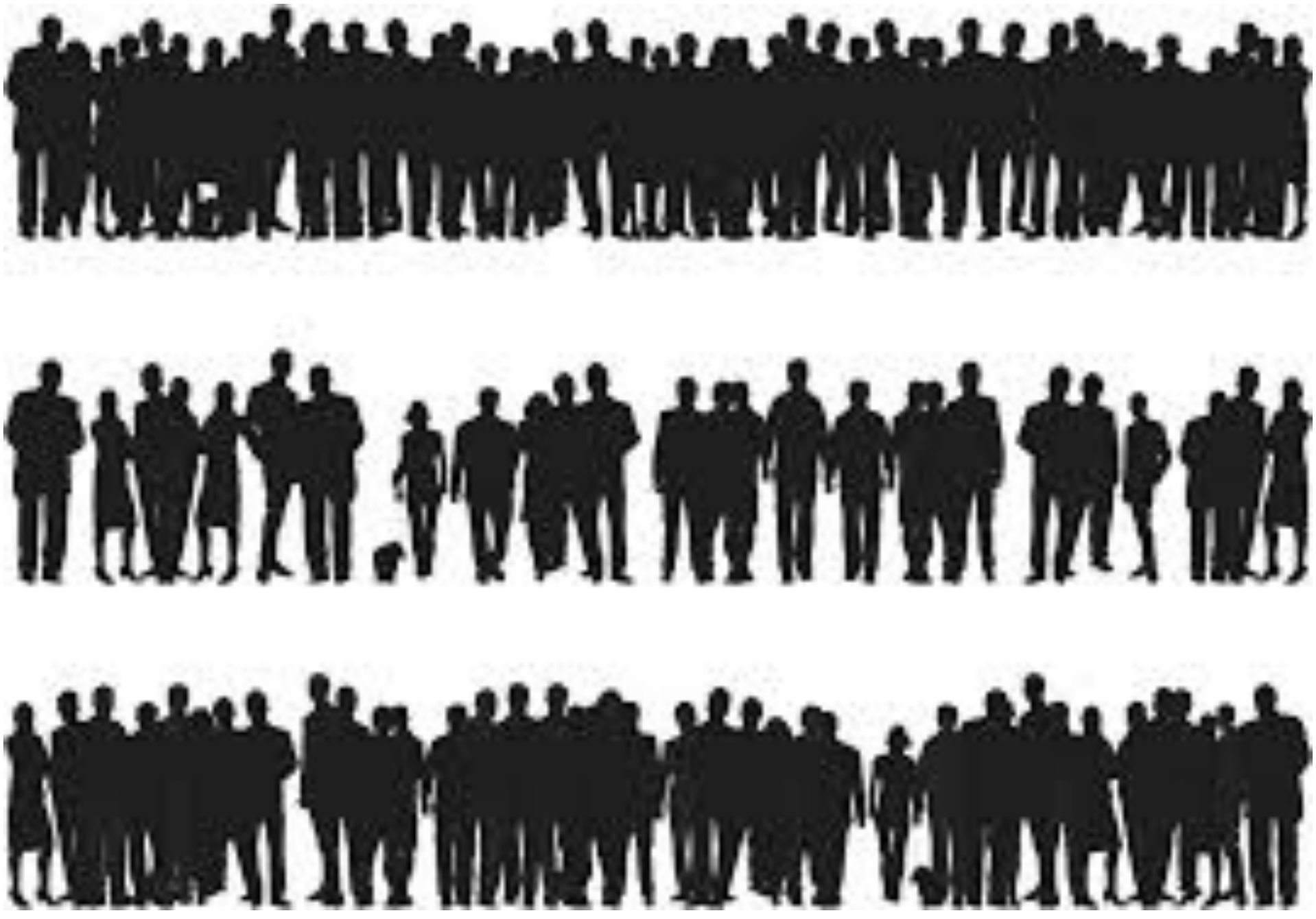
Mendelian traits (rare diseases)

Dark ages of human genetics

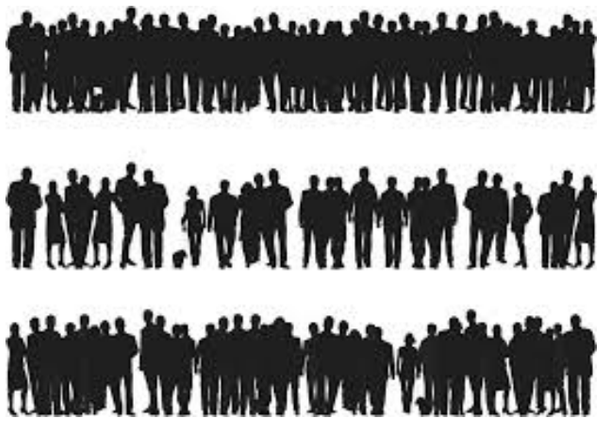


Comlex traits (common diseases)

Precision Medicine



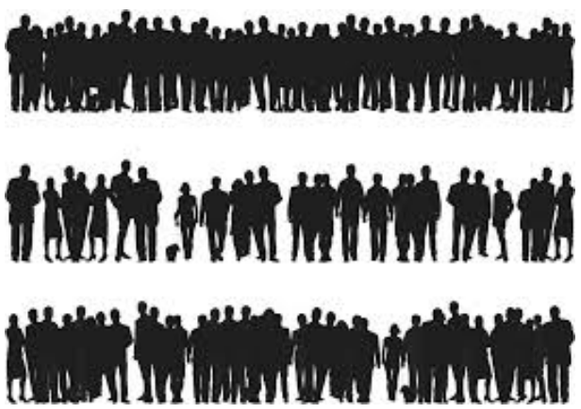
“Experiments of nature” that protect can guide selection of drug targets



Lower risk for disease

Examples of protective mutations

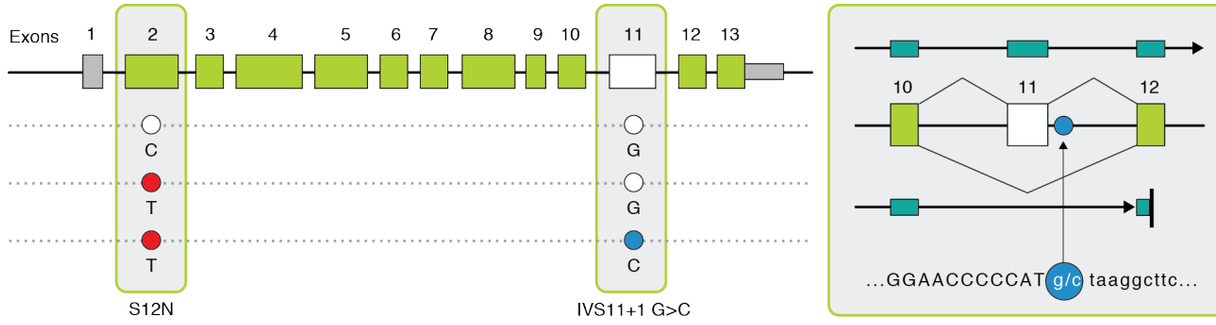
PCSK9 for LDL and MI
Nav 1.7 for pain
CARD9 for Crohn's disease and ulcerative colitis
RNF186 for ulcerative colitis
CCR5 for HIV



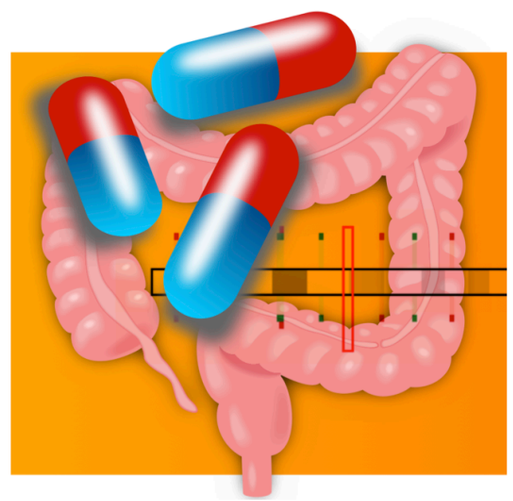
Lower risk for disease

Rare, strong acting alleles provide interpretation of the GWAS results

- Splice variant in *CARD9* cause premature truncating of protein and ***strongly protects*** against the development of Crohn's disease and ulcerative colitis ($p < 10^{-16}$).



- Protective genetic variants reveal process that is:
  - **safe** (naturally occurs in healthy adults)
  - **effective** (proven to reduce risk of disease).



Population medical data combined with **human genetic data** empowers novel **data science technologies**

Global Biobank Engine (pre-alpha)

Search for a gene or variant or region or phenotype coding (coming soon)

Examples - Gene: F5, Transcript: ENST00000367797, Variant: 1:169519049, RS ID: rs6025, Region: 10:114686614-114786614

Genetic Association Results

Note: We present summary statistic results from the UK Biobank hospital in-patient health-related outcomes summary information data (Data-Field 41202); computational grouping of phenotypes with cancer (Category 100092) registry, death registry data (Category 100093), algorithmically-defined outcomes (Category 42), and verbal questionnaire data (Category 100071); and manually curated grouping of phenotypes.

Browseable phenotypes

Cancer

Family History

Up next

October 10-18, 2017

- Imputation results upload with (Neale Lab).

Recent News

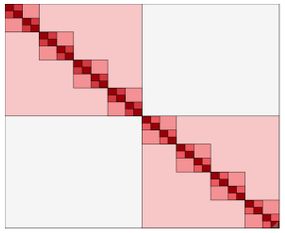
October 18, 2017

- Release of genetic parameters app I: genetic correlation.

October 18, 2017

- Release of decomposition app.

https://biobankengine.stanford.edu



Global Biobank Engine

Global Biobank Engine | About | Downloads | Terms | Contact | Power | FAQ & News archive

Select Language

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Browseable phenotypes

- Cancer
- Family History
- Verbal and ICD disease grouping
- Quantitative measures

Up next

August 10-18, 2017

- Imputation results upload with (Neale Lab).

Recent News

September 1, 2017

- Check out quantitative measures II.

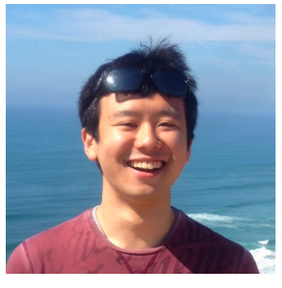
August 24, 2017

- GBE Power App available.

August 18, 2017

- Support for gene-based results: top five phenotypes per gene (independent effects model; more documentation soon).

August 13, 2017



Course projects

Topics will be proposed during next week's lecture

Data are organized by Yosuke

Goal : To implement and apply techniques learned in the class to big biomedical datasets

Data available in the course

1. **UK Biobank**

   Summary statistic data for over 100 diseases

2. ENCODE

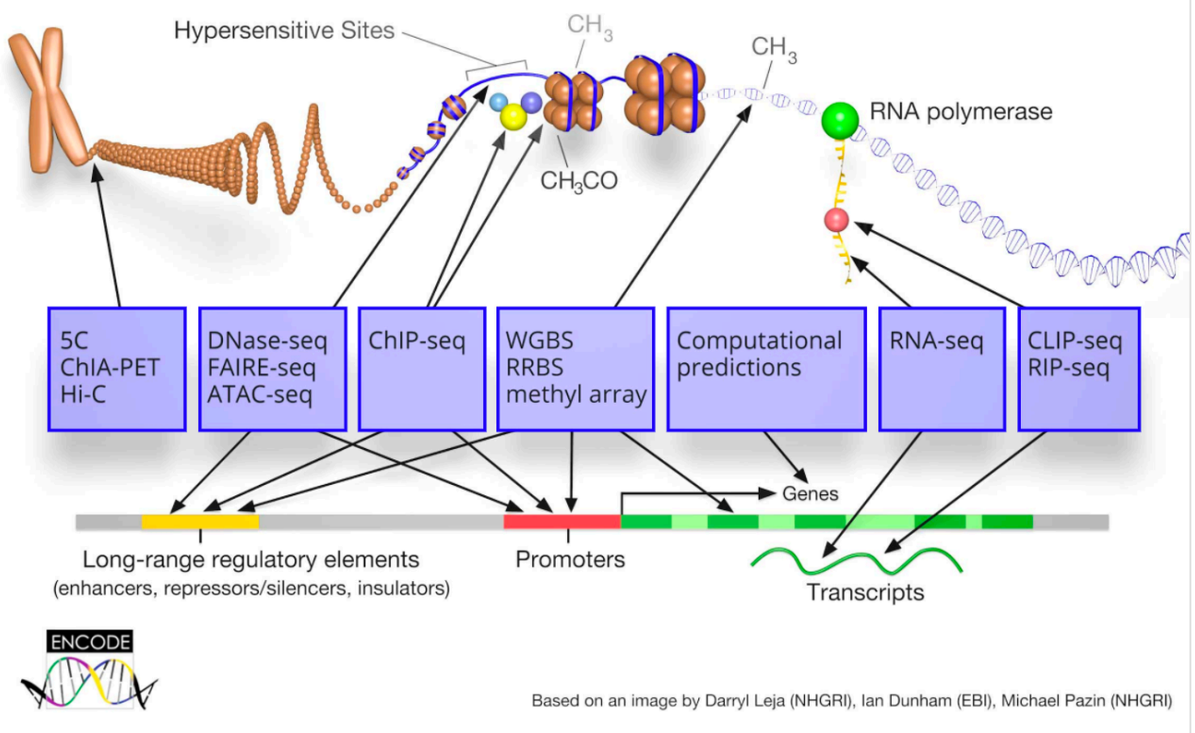
Data available in the course

1. UK Biobank

   Summary statistic data for over 100 diseases

2. ENCODE

ENCODE: Encyclopedia of DNA Elements



The ENCODE (Encyclopedia of DNA Elements) Consortium is an international collaboration of research groups funded by the National Human Genome Research Institute (NHGRI). The goal of ENCODE is to build a comprehensive parts list of functional elements in the human genome, including elements that act at the protein and RNA levels, and regulatory elements that control cells and circumstances in which a gene is active.

Get Started

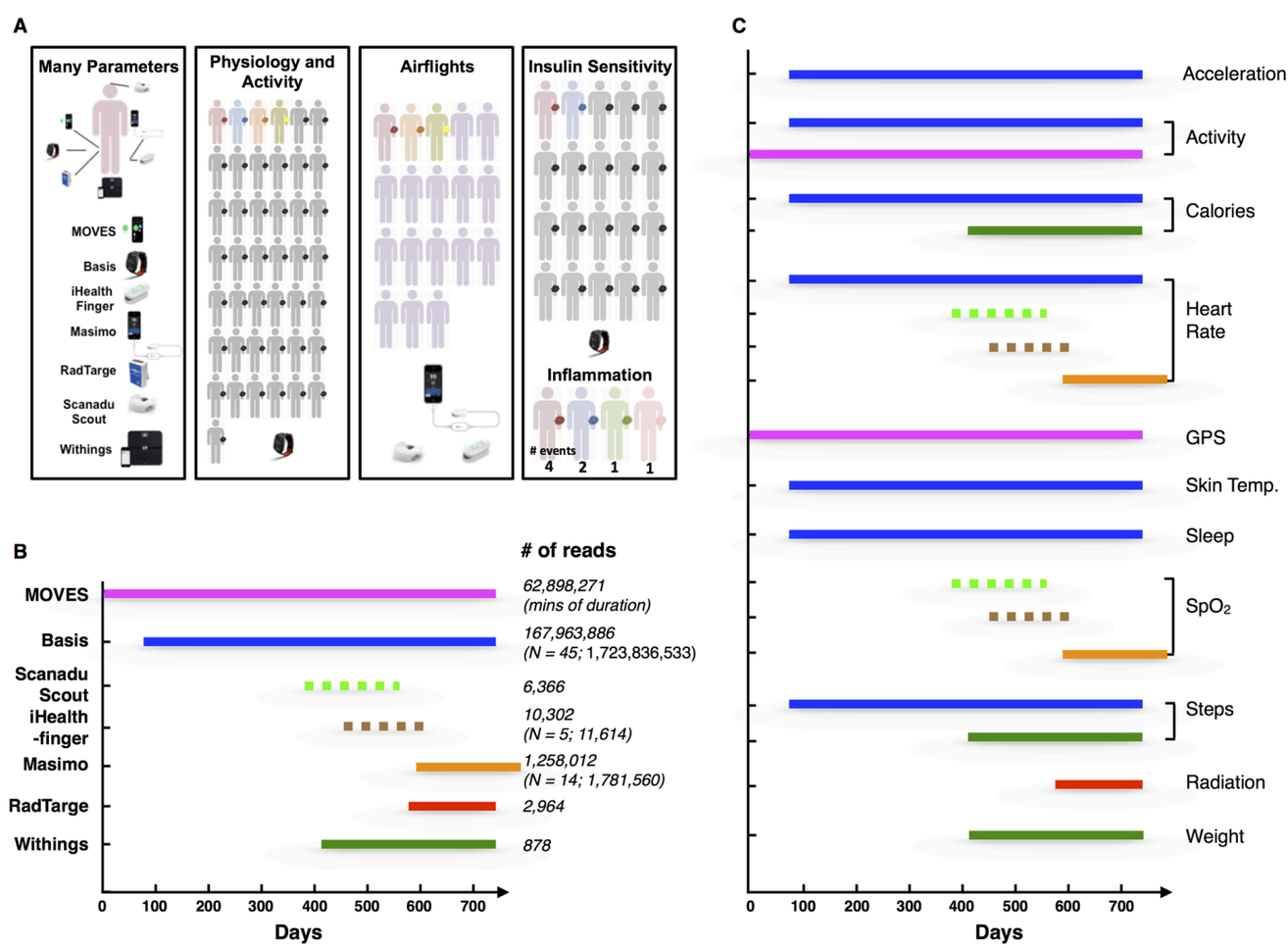
Data available in the course

1. UK Biobank

   Summary statistic data for over 100 diseases

2. ENCODE

3. Wearable Biosensor



Data available in the course

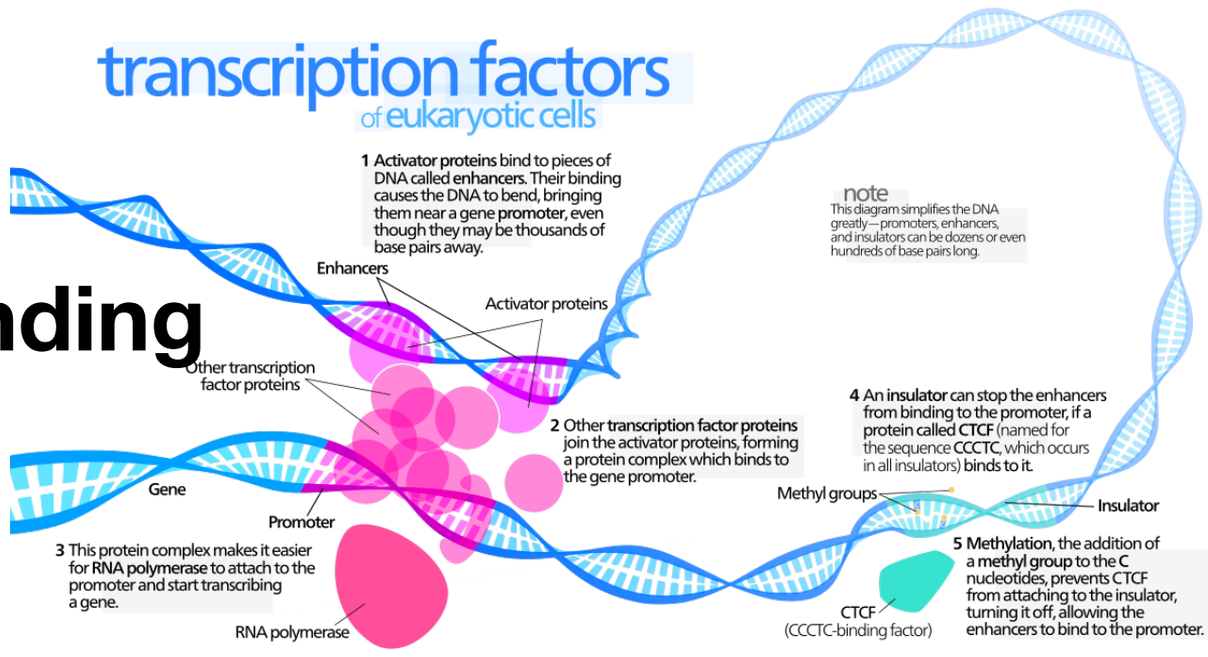
1. UK Biobank

    Summary statistic data for over 100 diseases

2. ENCODE

3. Wearable Biosensor

4. Transcription factor binding



50 years of Data Science

Ten to twenty years ago, John Chambers, Bill Cleveland and Leo Breiman urged academic statistics to expand its boundaries beyond the classical domain of theoretical statistics

50 years of Data Science, David Donoho 2015

50 years of Data Science

Chambers: more emphasis on data preparation and presentation

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50 years of Data Science

Breiman: more emphasis on prediction

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50 years of Data Science

"For a long time I have thought I was a statistician, interested in inferences from the particular to the general. But as I have watched mathematical statistics evolve, I have had cause to wonder and to doubt… All in all I have come to feel that my central interest is in data analysis, which I take to include, among other things: procedures for analyzing data, techniques for interpreting the results of such procedures, ways of planning the gathering of data to make its analysis easier, more precise or more accurate, and all the machinery and results of (mathematical statistics) which apply to analyzing the data."
— *The Future of Data* Analysis, John Turkey 1962

50 years of Data Science, David Donoho 2015

The Six Divisions of Greater Data Science

1. Data exploration and preparation

50 years of Data Science, David Donoho 2015

The Six Divisions of Greater Data Science

1. Data exploration and preparation

2.Data representation and transformation

50 years of Data Science, David Donoho 2015

The Six Divisions of Greater Data Science

1. Data exploration and preparation

2.Data representation and transformation

3. Computing with data

STAN in this course

50 years of Data Science, David Donoho 2015

The Six Divisions of Greater Data Science

1. Data exploration and preparation

2.Data representation and transformation

3. Computing with data

4.Data modeling

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The Six Divisions of Greater Data Science

1. Data exploration and preparation

2.Data representation and transformation

3. Computing with data

4.Data modeling

5. Data visualization and presentation

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The Six Divisions of Greater Data Science

1. Data exploration and preparation

2.Data representation and transformation

3. Computing with data

4.Data modeling

5. Data visualization and presentation

6.Science about Data Science

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The Next 50 years of Data Science

Open Science takes over

Reproducibility

Documented workflows

Science as data

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Topics you will learn in this course

Poisson Models

Rank based methods - permutation

Monte Carlo, Markov Chains

Splines, Fourier analysis, PCA

Bayesian multilevel modeling

Bayesian mixture models

Bayesian prediction

Distributed computing with privacy preserving schemes

Convolution Neural Networks, LSTM

Course website

Syllabus
Reading materials
Lecture notes https://canvas.stanford.edu/courses/66507
Problem Sets
Data links

Sample STAN programs
Application to data https://biods215.github.io/
Github repository

Github repository for the course

https://biods215.github.io/