

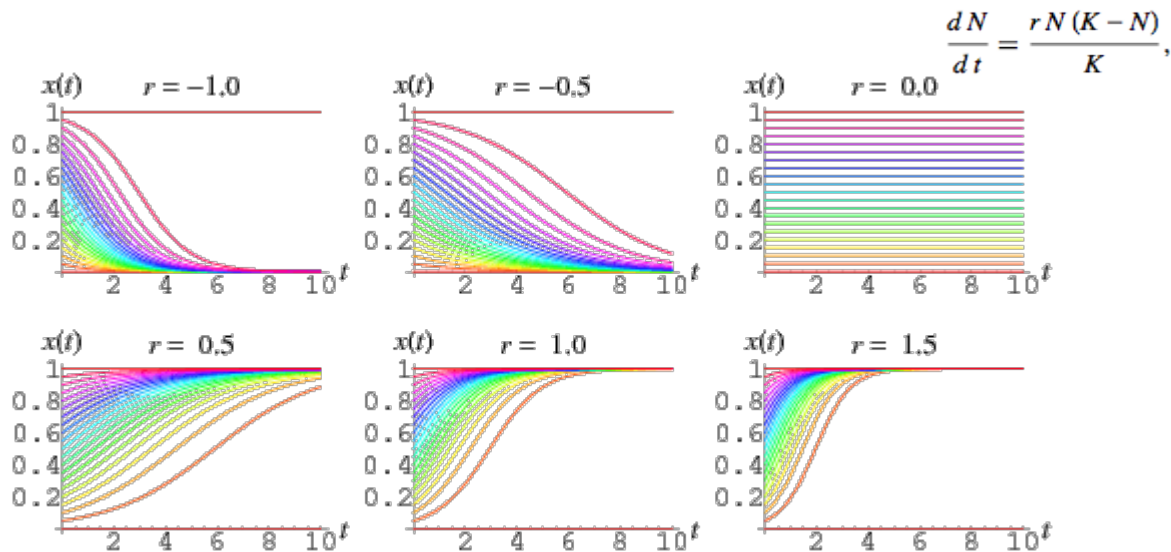


Informatics, algorithmics, and automation: machine learning in 21st century biology

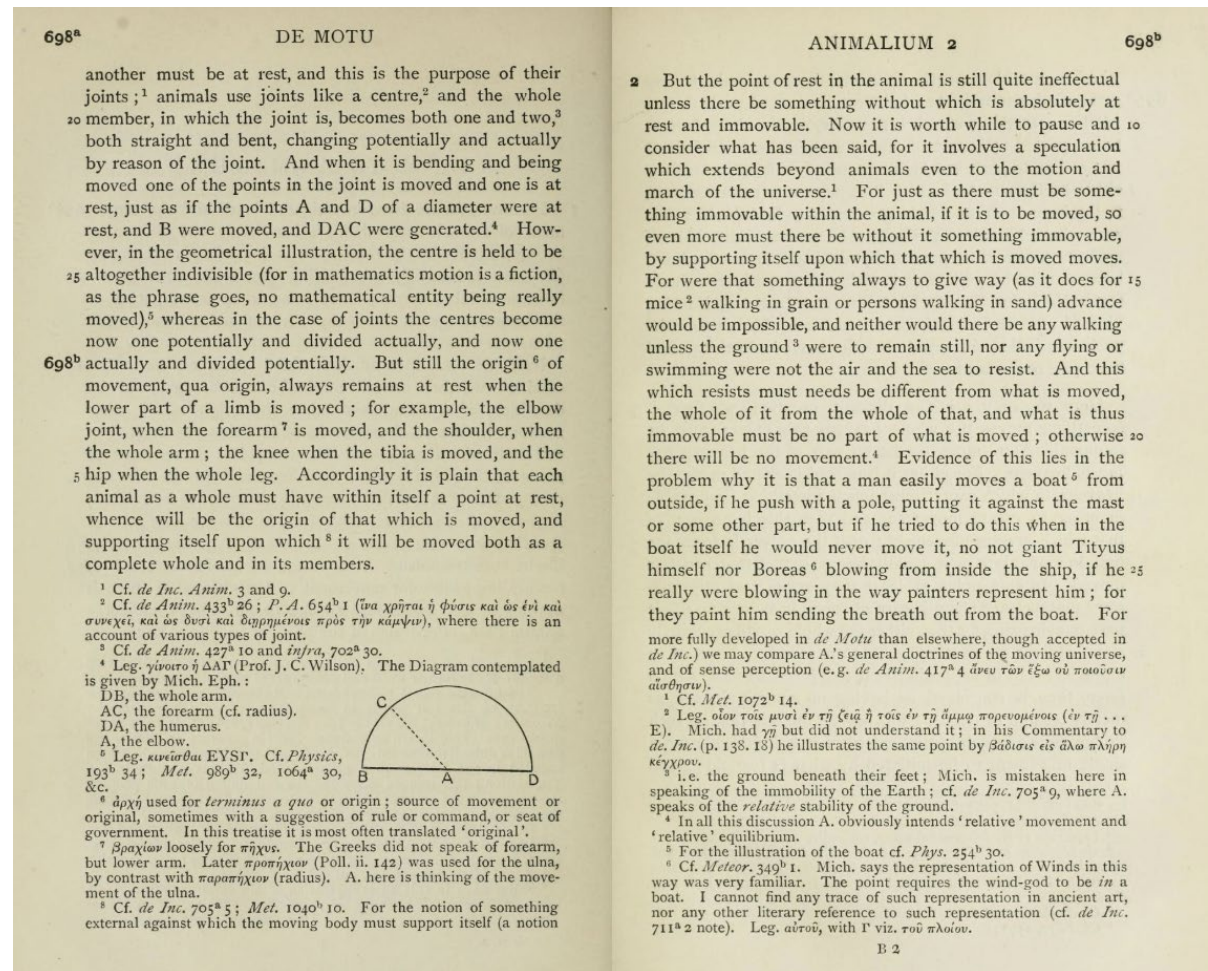
Pradipta Ray
UT Dallas

Mathematics in biology & medicine

- A very long history – as early as the Hellenic civilization
- All branches of mathematics are harnessed – from classical geometry to calculus



Plots of the logistic equation modelling population growth



On the motion of animals, Aristotle, 4th century BC (translated 1912)

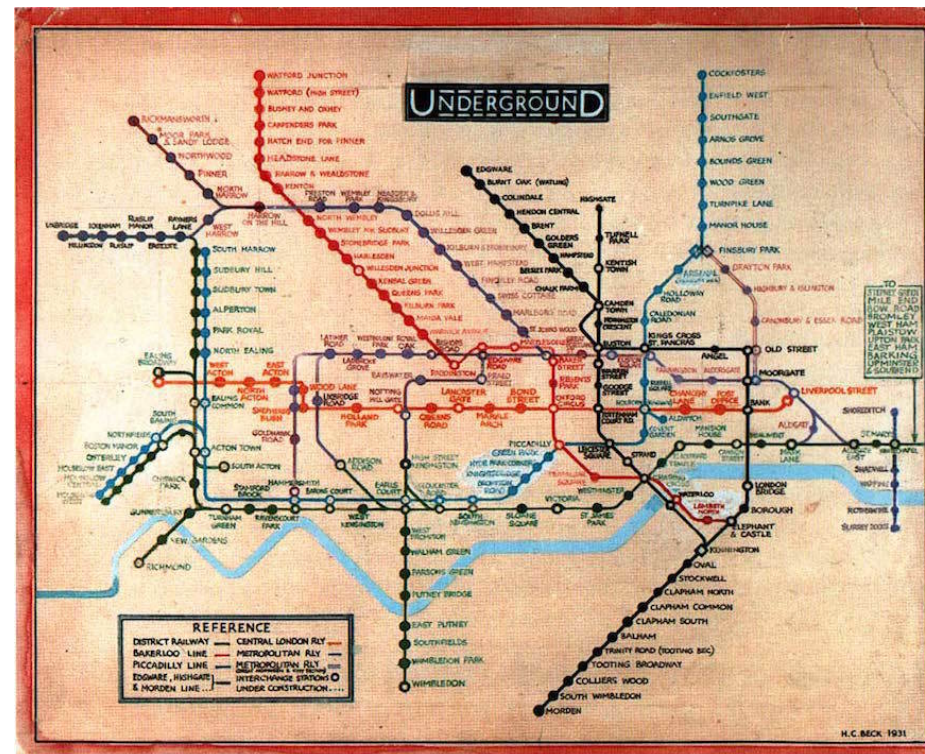
Informatics : what aspects of mathematics does it deal with ?

- **Mathematics pertaining to data**

- Automated description, modelling, visualization, prediction (and imputation), etc
- Large overlaps with Computer Science (machine learning) and Statistics community



Term first coined in 1957 though discipline dates back to ancient India / China / Greece



Harry Beck's 1931 visualization of London's tube network

Beck was a draftsman for the underground railway before being fired and voluntarily creating map & selling to ex-employer for 10 GBP !

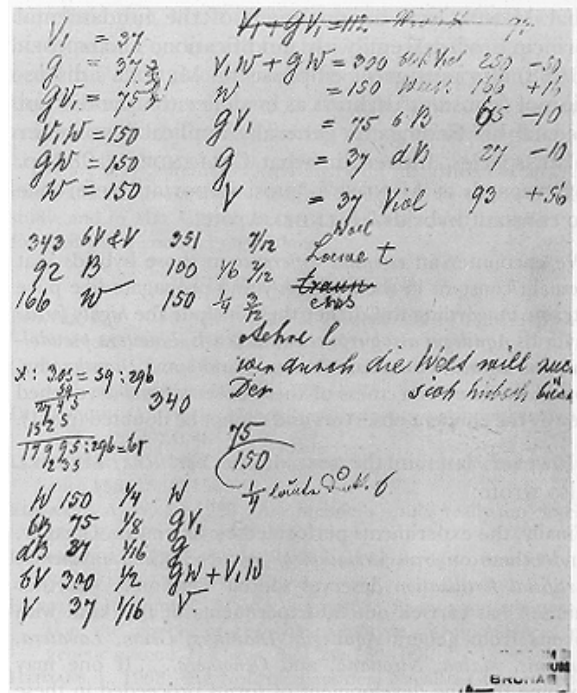
Algorithms in biology

- Algorithm : a set of instructions for performing a task / computation
 - Set of rules to navigate a car from school to home
 - Computer program for sorting number

Computational Biology

Use mathematics & CS to answer questions in biology – which traits of a pea plant are inherited independently of others and at what frequency ?

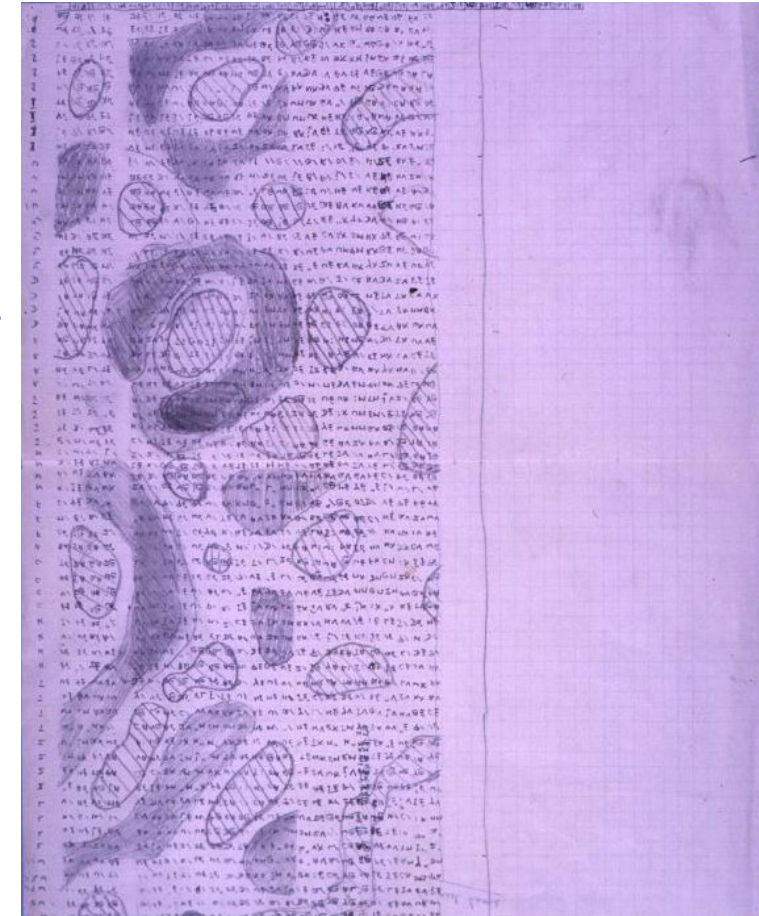
Gregor Mendel's notebook detailing his experiments with breeding peas



Computation in biology

What are the underlying instructions driving biological processes – how do molecular gradients give rise to morphological patterns ?

Alan Turing's notebook sketches for modelling how pattern formation occurs in nature



The dawn of computers

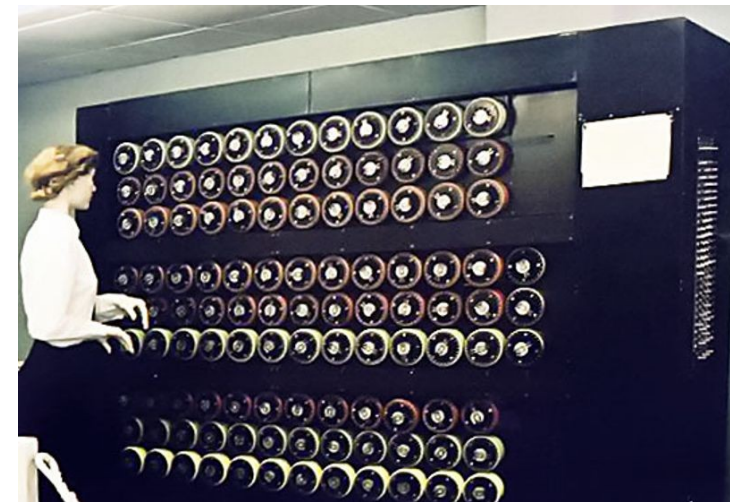
- Algorithm execution and informatics were laborious tasks before computers came about
- The invention of electronic computers and dizzying improvements in speed and memory led to most informatics tasks being performed on computers

Computers + algorithmics + informatics = Machine learning
Computers + algorithmics + informatics + biology = Bio-informatics



Wikipedia

Radhanath Sikdar, the human “computer” who first calculated the height of Mt Everest

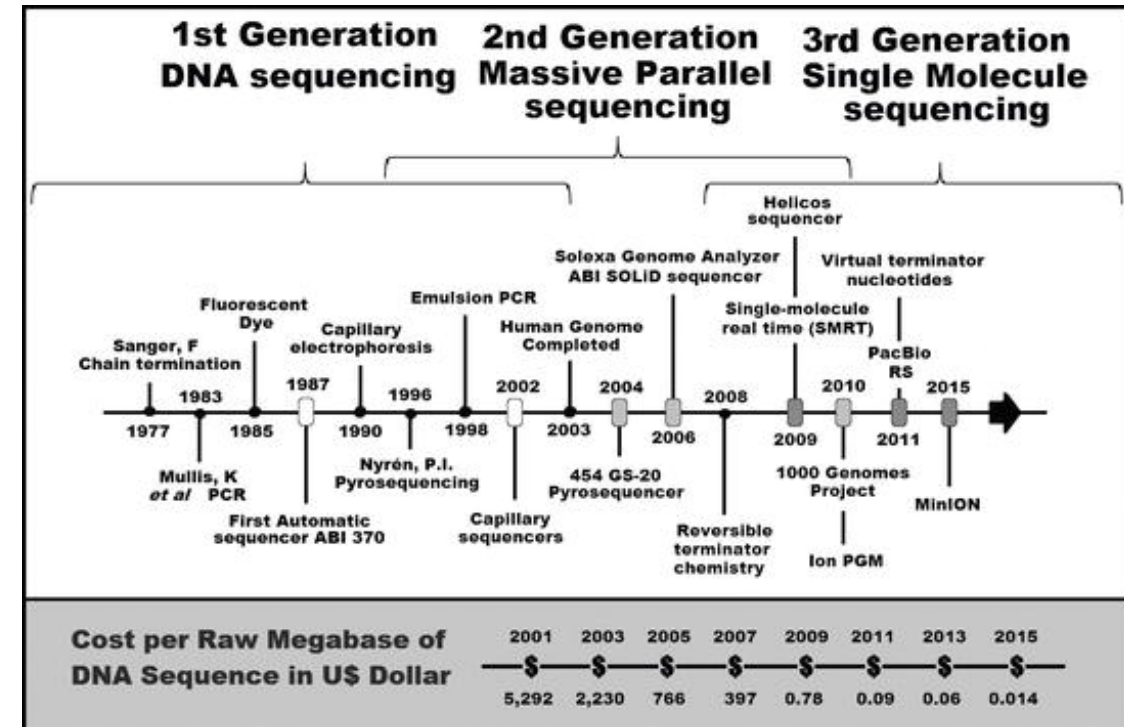
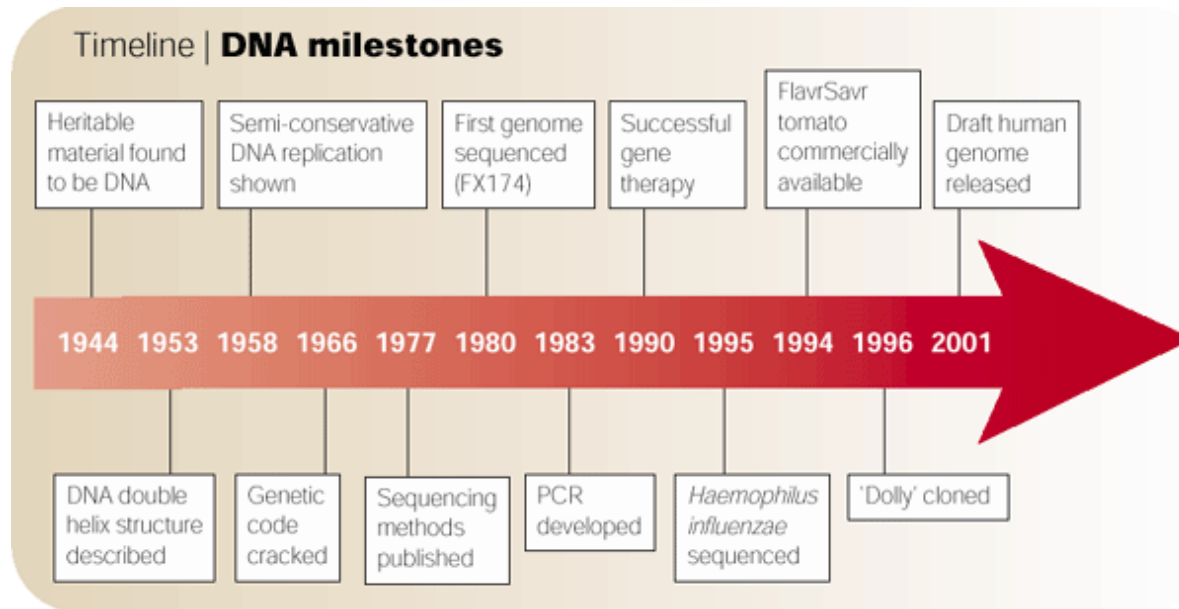


Computerhistory.org

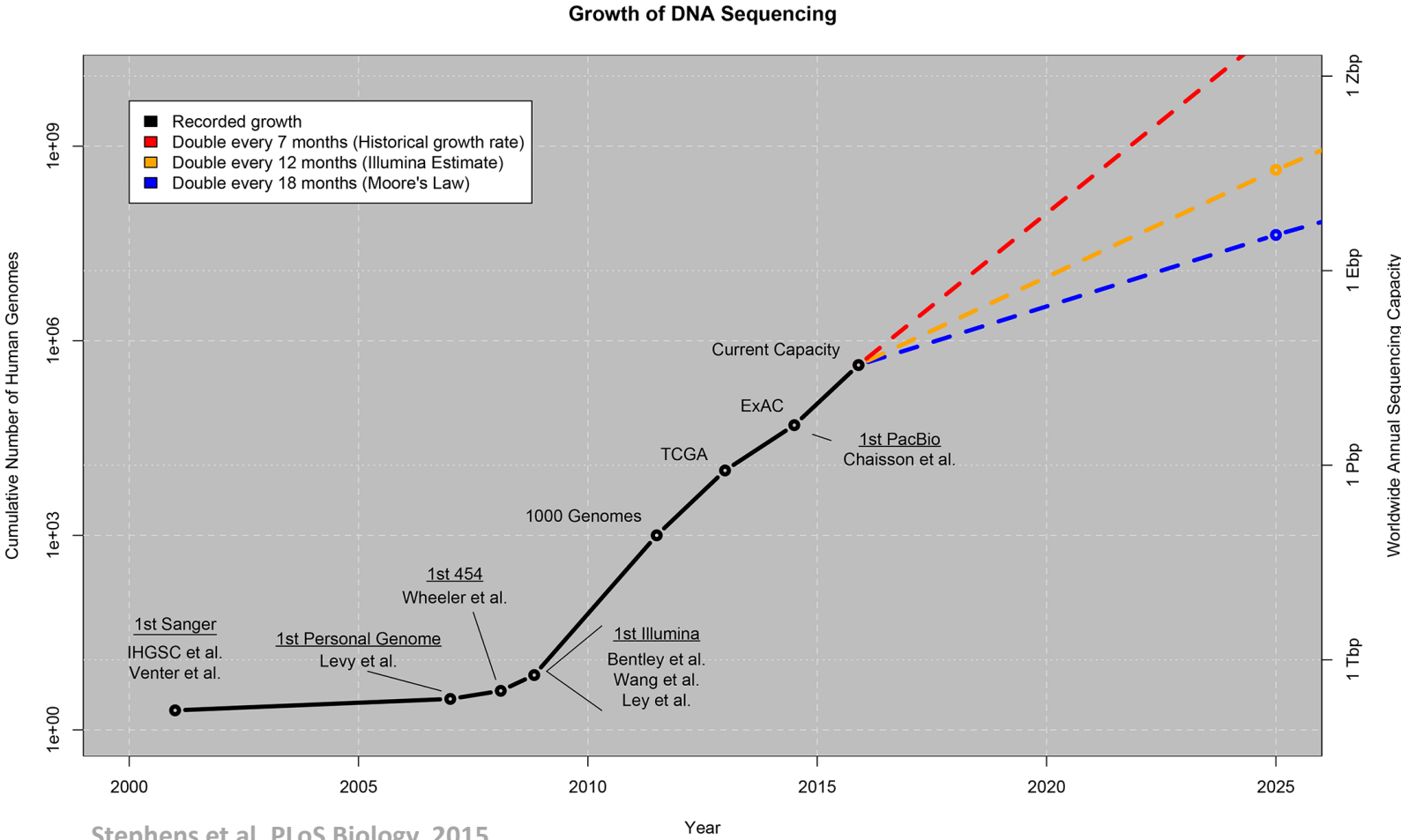
The Bombe (1942), one of the first large scale electromechanical devices built in Bletchley Park for British WW2 cryptanalysis

Nucleic acid research : understanding, sequencing, engineering

Baxter, Nature, 2003



Automation and scale in nucleic acid sequencing

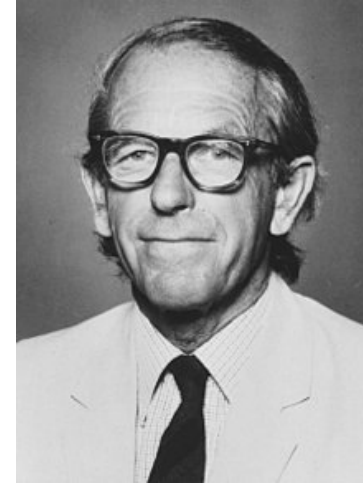


Stephens et al, PLoS Biology, 2015

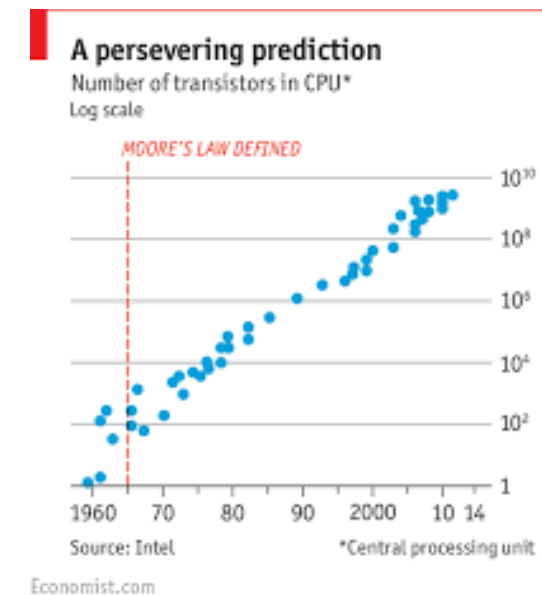
Bioinformatics : 3 decades of explosive growth

- **Rapid improvements in molecular sequencing technologies** for peptides, DNA and RNA produces large amounts of data to be analyzed
- Moore's Law : computers tend to double in computing capacity every year, making **intensive computation feasible, tractable and economical**
- Development of methodology, training a new kind of multidisciplinary scientist, investment in infrastructure – the **Human Genome Project** (1990 – 2003)

Wikipedia



Frederick Sanger : Nobel prize winner (twice!!) for studies on sequence and structure of insulin and for nucleic acid sequencing



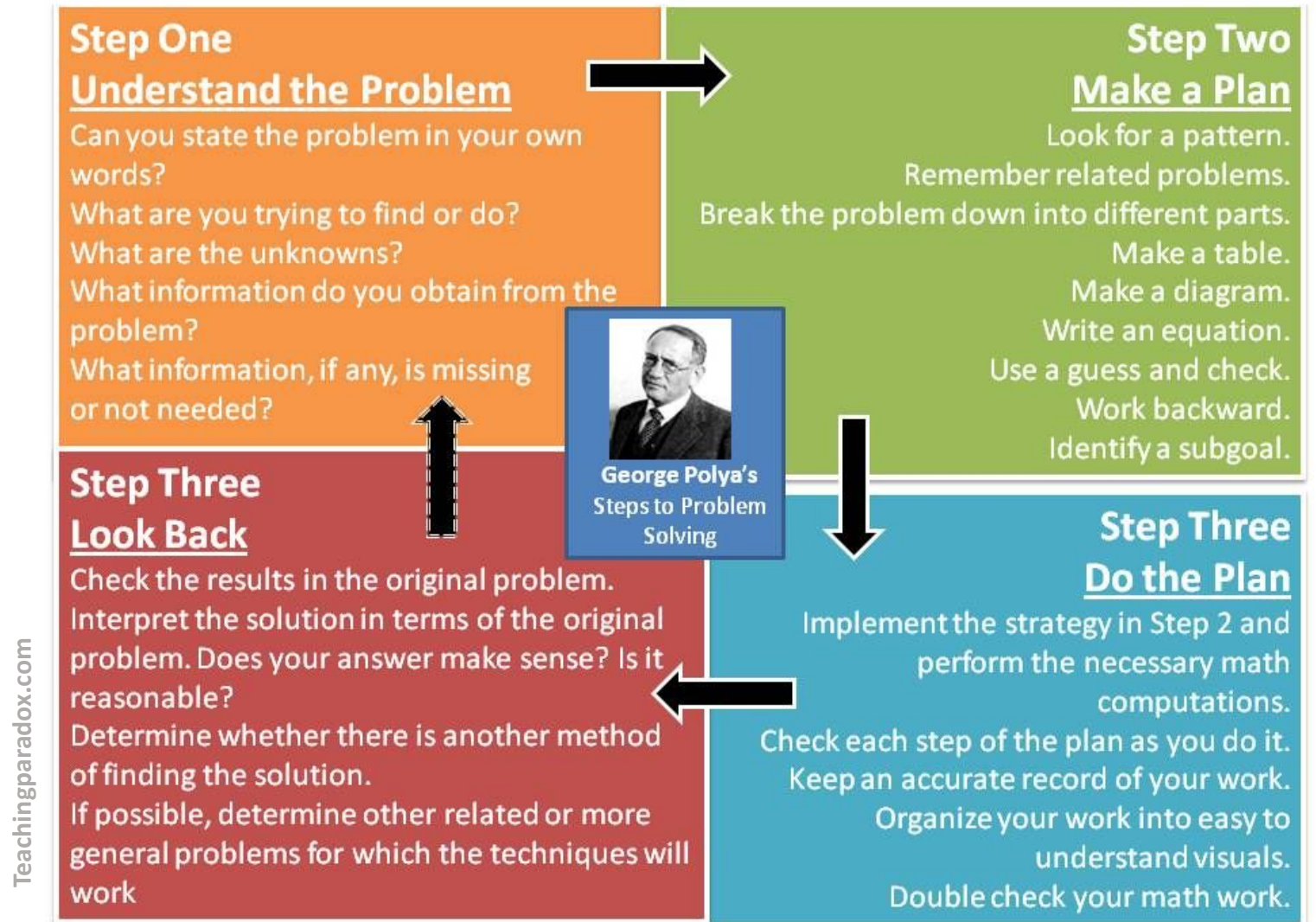
The mouse & human genome projects

- Sequenced and assembled reference drafts of the human and mouse genome
- Mapped and predicted genes in the reference sequence
- Laid the groundwork for genomics (including computational genomics) as a discipline



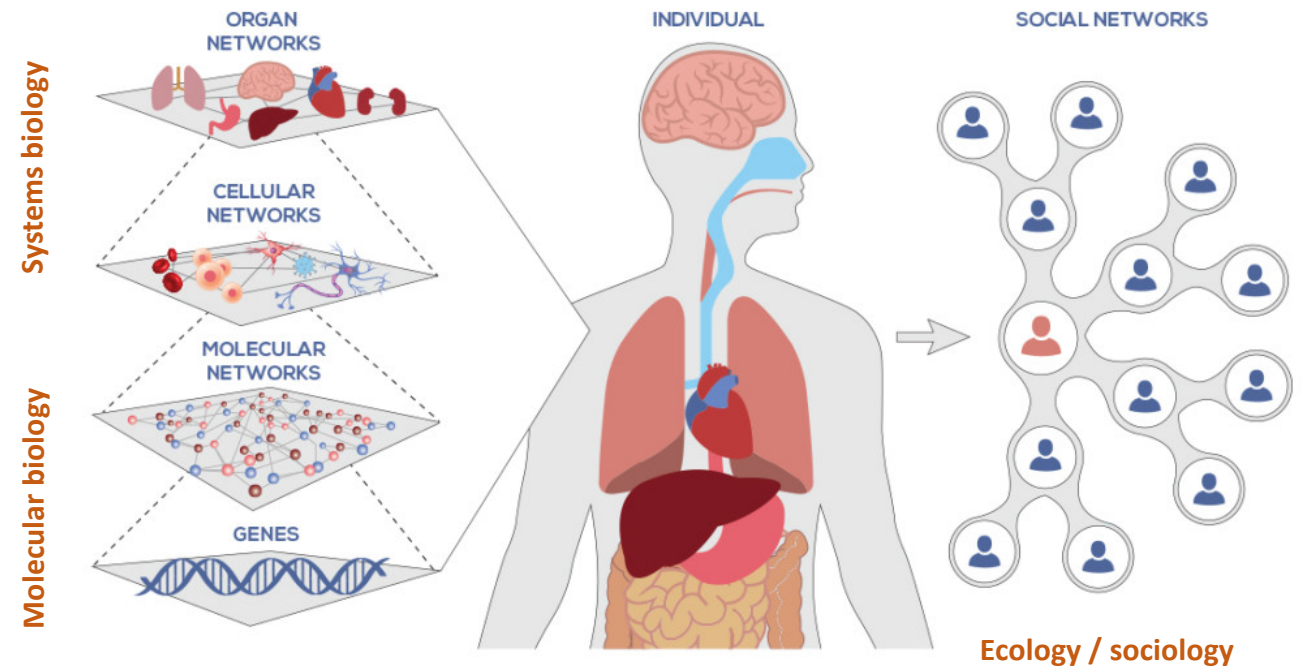
Asking the right questions

- How to solve it, written by George Polya
 - Strongly encouraged to read this book !
- Problem solving is a cyclic process : true for bioinformatics as well



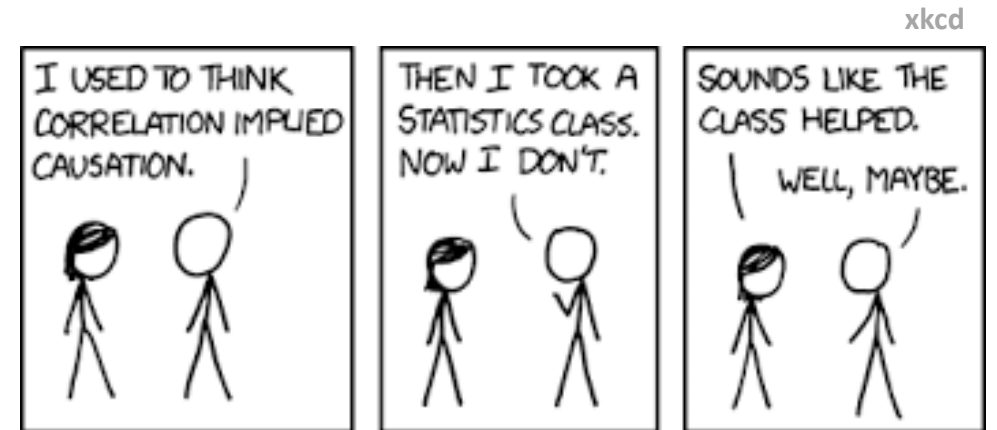
Performing analysis at the right granularity

- Given a biological problem, it is important to figure out the right granularity to perform experiments / note observations
- Some problems can be studied at multiple levels : disease - molecular pathology, systems level changes in physiology, transmission of disease



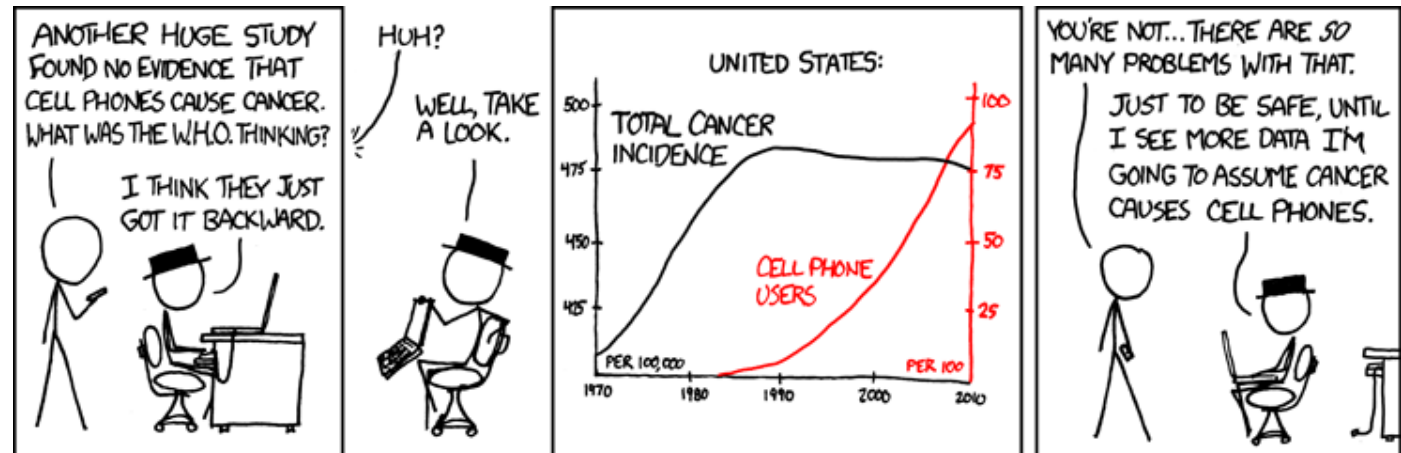
Bioinformatics as an epidemiological science

- A large amount of biology is an empirical science
 - Bioinformatics helps generate, screen and test hypotheses in high throughput fashion to help transition it to a theoretical science
- Since many bioinformatics studies are correlative, follow up studies that de-confound correlation and causation are often required
 - Perturbation studies (eg. gene knockout models)



Pitfalls of correlative studies

- Poorly set up hypotheses, bad inferential mechanisms, and low quality data can all contribute to arriving at the wrong conclusion (artifacts)
- Methodological rigor and domain knowledge are key to avoiding artifacts



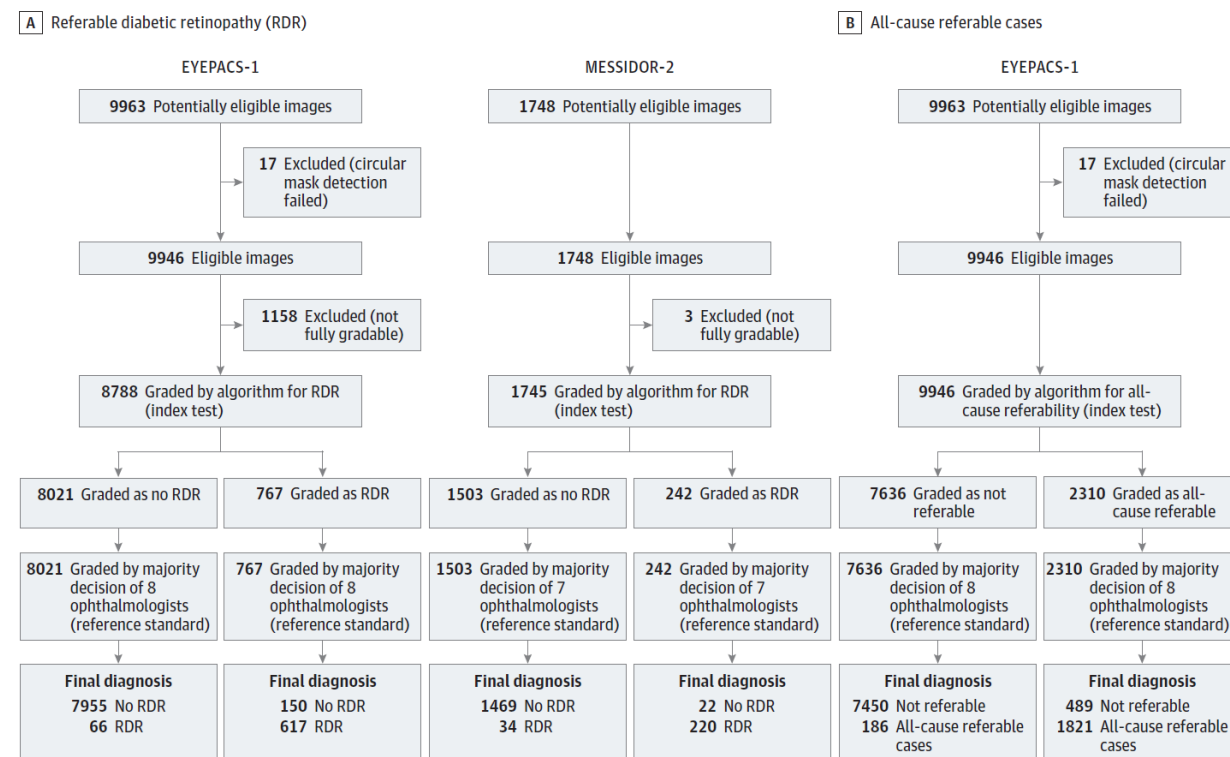
“Artificial intelligence” in biology

The perception of AI has changed over the years :

1970s : expert systems (eg. early recommender systems in medicine)

1990s : statistical machine learning (eg. genome wide association studies of diseases)

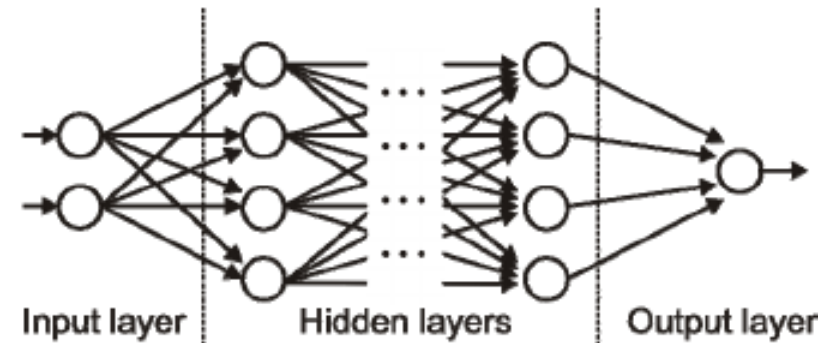
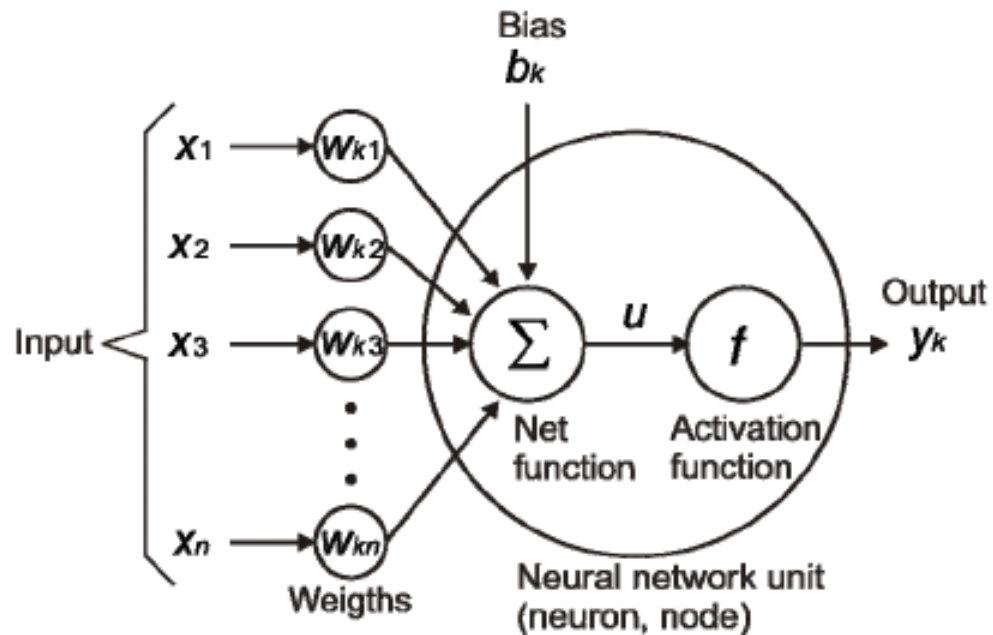
2010s : deep neural networks (eg. prediction of diabetic retinopathy from imaging)



Gulshan et al, JAMA, 2016

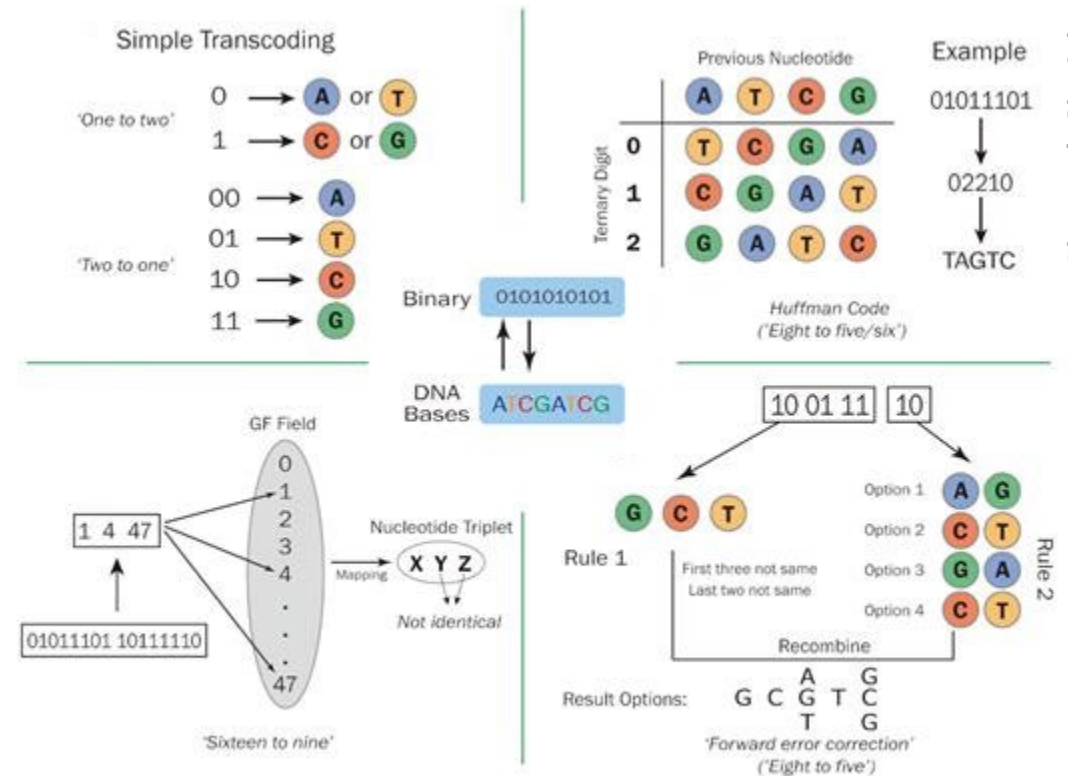
Using neuroscience to model AI

- Neural networks



Other areas of computer science – biology intersection

- **Biological computation** : how can biological processes be harnessed to improve computing – eg. DNA based data storage
- **Biomedical devices** : electronic devices used for patient monitoring, treatment and therapy
- **Synthetic biology** : creating new biochemical / biological entities



Information encoding strategies for DNA – based storage

If you'd like to perform research in bioinformatics ...

- Knowledge of coding is required
- Send email to pradiptaray@gmail.com
- Many other wonderful laboratories in UTD working on bioinformatics