

Why Deep Learning in Biomedicine

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Why deep learning in biomedicine



- Transformation of biomedicine into data science
- The future role of deep learning
- Why deep learning?
- The biggest obstacle and suggestions to overcome it

Sequencing Growth



Cost of one human genome

2000: \$3,000,000,000

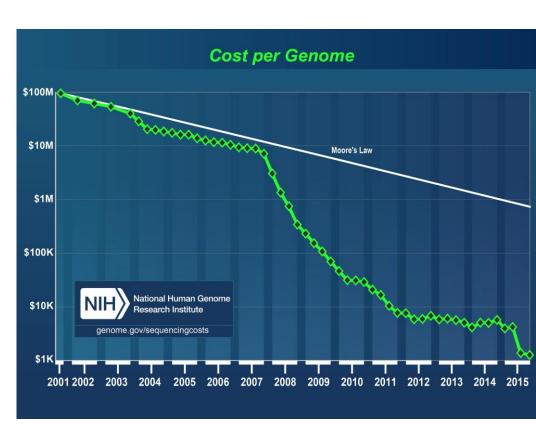
2004: \$30,000,000

• 2008: \$100,000

• 2010: \$10,000

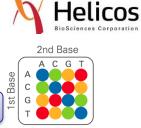
• **2015**: \$1000

• ???: \$300













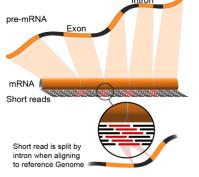




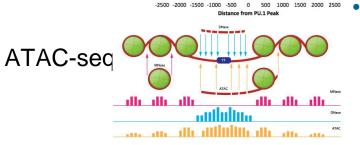
A Data-Acquisition Technology Explosion Enabled by Inexpensive Sequencing



RNAseq



DNase-seq



₽ 0.04

S 0.03

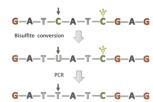
Hi-C



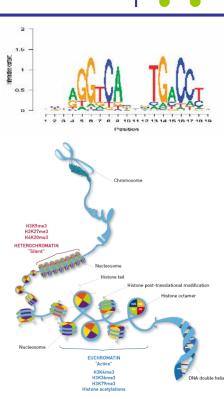
H3K4me2 (Mnase)

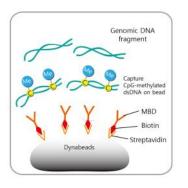
- ChIP-seq for
 - Transcription factor binding
 - Nucleosome positioning
 - **Histone Modifications**

Bisulfite treatment for methylation



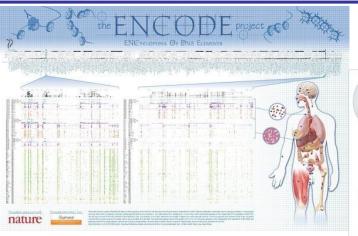
MeDIP for methylation



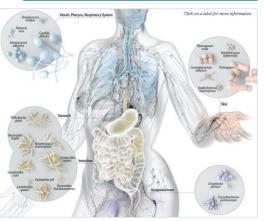


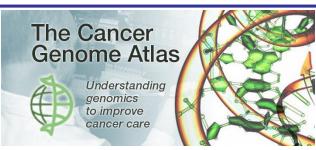
Major Data Acquisition Efforts





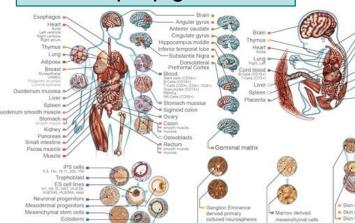
Human Microbiome













Tools

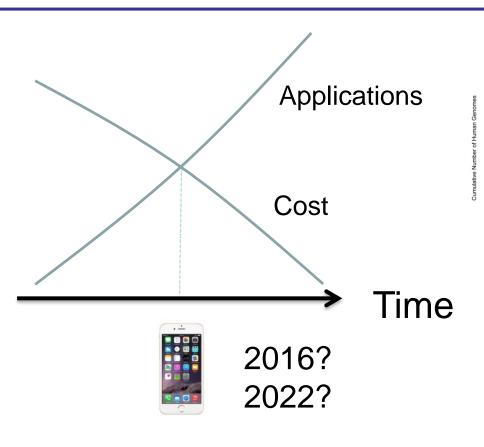
of the

1000 Genomes

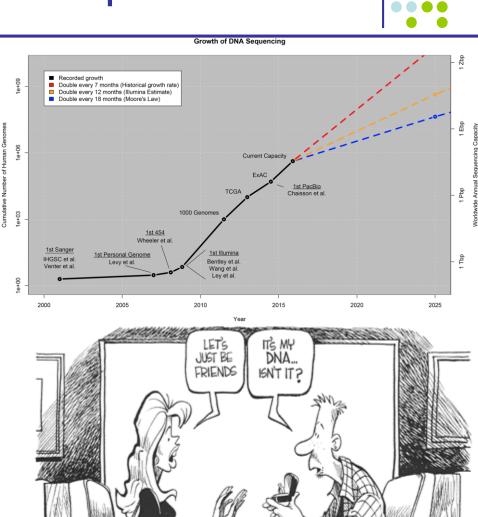


How soon will we all be sequenced?



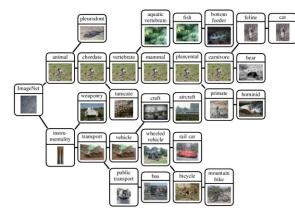


1Bn individuals by 2022-2026

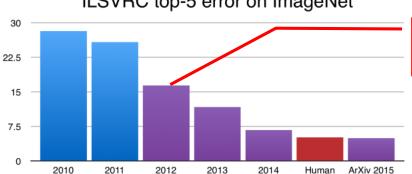


Deep Learning – major Al breakthrough





ILSVRC top-5 error on ImageNet







"man in black shirt is playing



"construction worker in orange safety vest is working on road."



"two young girls are playing with leggs toy."



"boy is doing backflip on wakeboard."



"girl in pink dress is jumping in



"black and white dog jumps over



'young girl in pink shirt is swinging on swing."



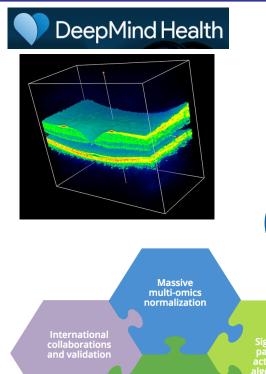
"man in blue wetsuit is surfing of wave."

Introduction of deep NNs

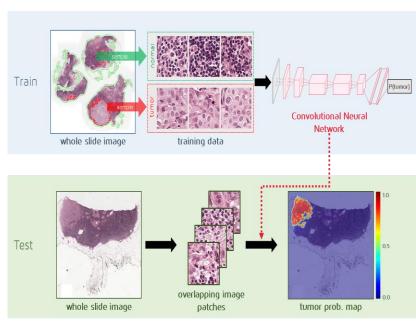


The beginning of deep learning in medicine

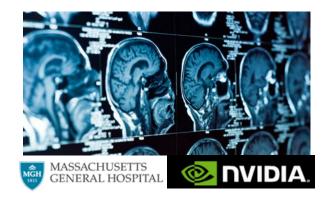








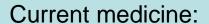




Bet: Biomedicine amenable to deep learning



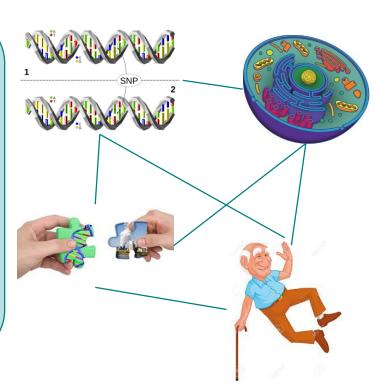




- With nurse help
- Oncology, heart, infectious diseases, dermatology, pharmacology, ..., diagnosis, treatment

Benefits:

- Improved outcomes
- Prediction, prevention
- Low cost



- Personalized health & prevention
- Powerful AI, learning from billions of examples
- New drugs & treatments, gene therapy, CRISPR, ...

OK, but why "deep learning"?



Let's digress and talk about

Boolean circuits

OK, but why "deep learning"?



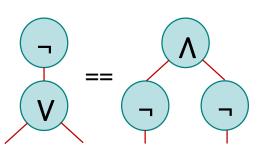
Consider circuits built of: V, Λ ,

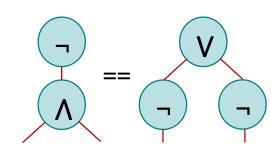
V





Can always push ¬ to bottom layer

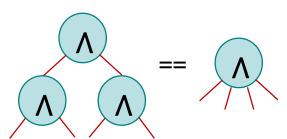




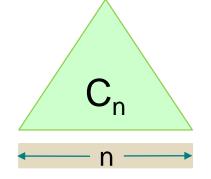
Assume alternating V,

\(\Lambda \)

Unlimited fan-in



Assume a circuit C_n that handles all inputs of length n



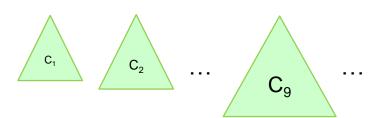
OK, but why "deep learning"?



A collection of circuits $C_1, ..., C_n, ...$ is

P-uniform: There is a poly-time Turing Machine that on input n, outputs C_n

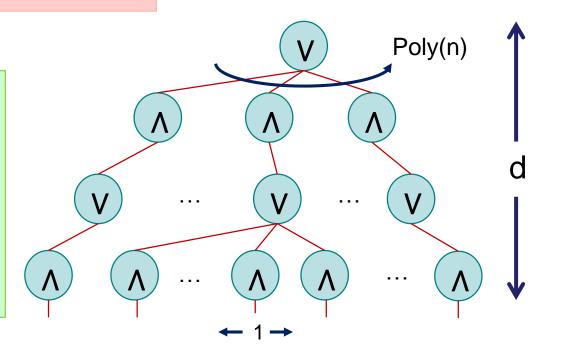
A language L is computable by a P-uniform circuit family, if and only if L ∈ P



Class AC⁰:

Unlimited fan-in circuits of constant depth d

Assume bottom layer fan-in 1



What can we compute with a constantdepth circuit?



- Start with depth-d circuit
- Collapse 1st & 2nd layers Switching Lemma

- - -

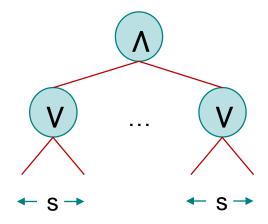
- Flatten to 2 layers without making it "too fat"
- What can we now compute?

Switching Lemma



k-DNF, n variables

s-CNF, n – t variables



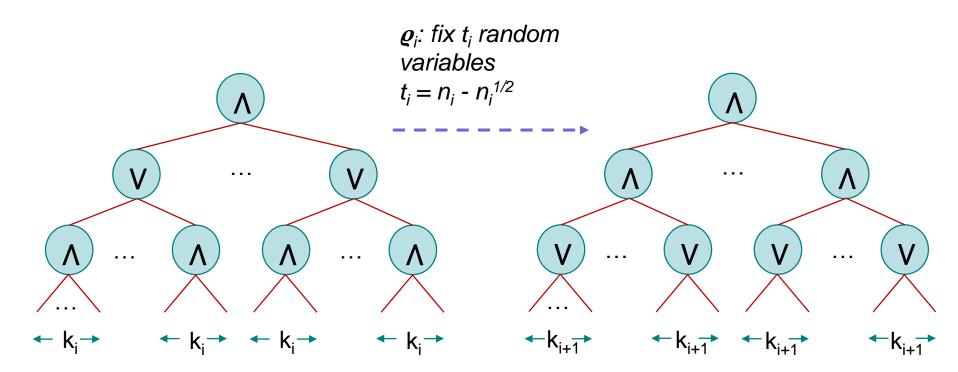
Switching Lemma (CNF ←→ DNF)

Prob [$f|_{\rho}$ not expressed as a s-CNF] $\leq q$]

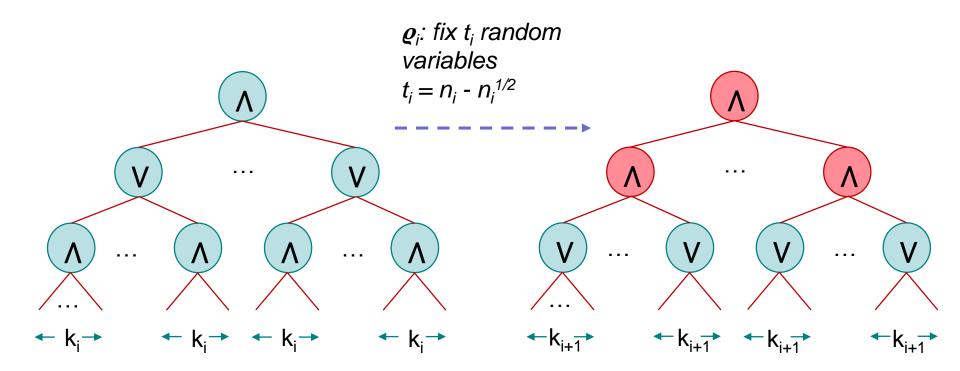
Where q is equal to the ugly:

$$q = q(k, s, t, n) = ((n-t)k^{10}/n)^{s/2}$$

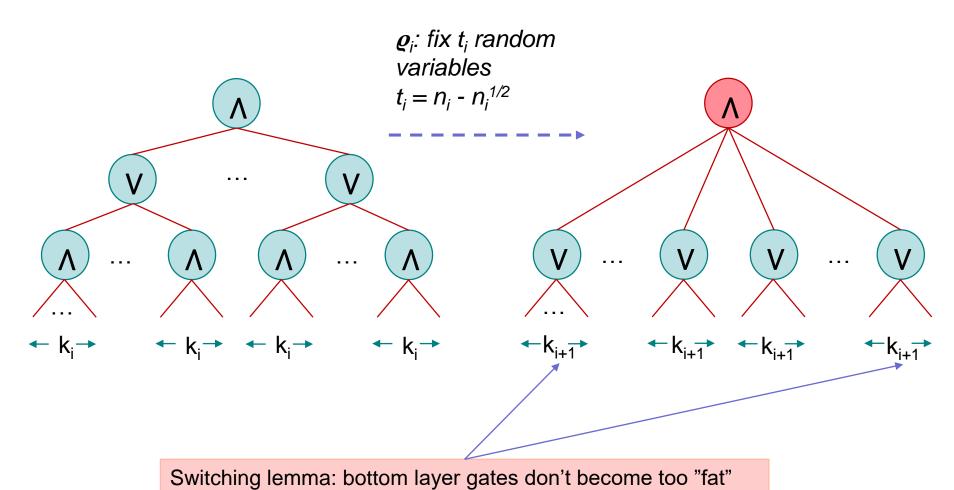




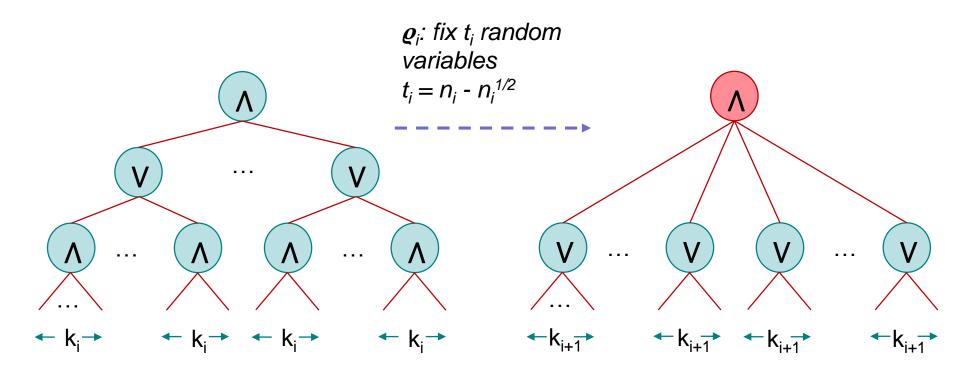












After some hairy calculations,

After d-2 steps, each applied to all layer-2 gates:

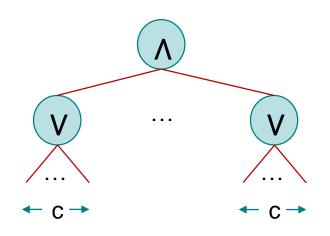
variables left: $n^{1/(2^{d-2})}$

depth: 2

fan-in: 10 b 2^{d-2} , where circuit size = n^b

Final Circuit





variables left: $n^{1/(2^{d-2})}$

depth: 2

fan-in: $c = 10b2^{d-2}$

Turn any circuit family in AC⁰ into 2-layer

If n large, #vars >> fan-in
Fix c vars, make output constant = 0

Now take $\mathbf{f} = \text{parity}(\mathbf{x}_1, ..., \mathbf{x}_n)$ \neq constant by fixing any $\mathbf{s} < \mathbf{n}$ variables

Therefore, $\mathbf{f} \notin AC^0$

A "shallow" ML method cannot compute parity unless exponential # features

How about HMM, K states? N layers, each layer receives log K bits from previous layer. Can do parity, but limited to simple functions

Shallow ML seems unable to replace general algorithms

Biggest Obstacle: wide, free data availability



Ideal:

- Millions of publicly available genomes, p
- Far-reaching anti data-based discrip

Not ideal:

- Data silos in hospitals, health care p.
- Specialized large-scale projects (1000 ge.

"I don't want to live in a world where someone else makes the world a better place better than I do."

cal records)

What about privacy?

- No cancer patient has died yet because of data privacy breach
- My Facebook page reveals much more private stuff about me than my genome or medical record

Let's make our data public – if 5% of us do, probably enough to solve medicine

Let's do so without waiting for legislation

Some examples to follow



You think recycling saves the world?

Try data sharing!

Anti-discrimination law and public awareness extends to all genetic and medical information

Data hoarders (academia, hospitals, industry) akin to polluters

