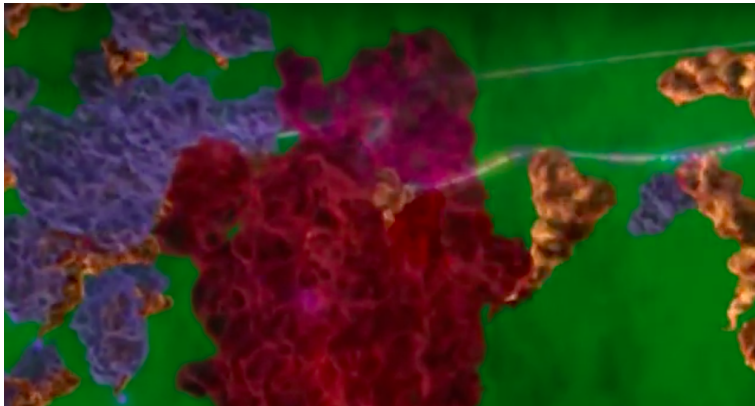


BIO LOGIC: Biological Computation

Kay Kirkpatrick, Math 490

2019

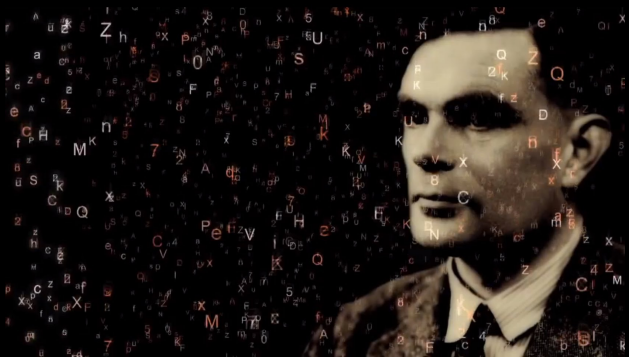
BIO LOGIC: Biology and Computation



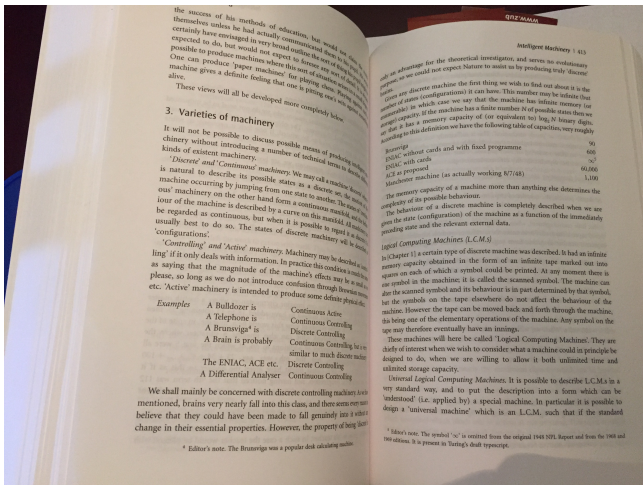
Kay Kirkpatrick, Math 490

Urbana, Miami and Mascouten lands

Movie still from the Yonath lab; next slide from Cambridge U.



Turing 1948: Intelligent Machinery



Courtesy Copeland, Complete Turing

Turing 1948: Intelligent Machinery zoom-in

'Controlling' and 'Active' machinery. Machinery may be described as 'controlling' if it only deals with information. In practice this condition is much the same as saying that the magnitude of the machine's effects may be as small as we please, so long as we do not introduce confusion through Brownian movement etc. 'Active' machinery is intended to produce some definite physical effect.

Examples

| | |
|-----------------------------|--|
| A Bulldozer is | Continuous Active |
| A Telephone is | Continuous Controlling |
| A Brunsviga ⁴ is | Discrete Controlling |
| A Brain is probably | Continuous Controlling, but is very similar to much discrete machinery |
| The ENIAC, ACE etc. | Discrete Controlling |
| A Differential Analyser | Continuous Controlling |

Turing's Mistake

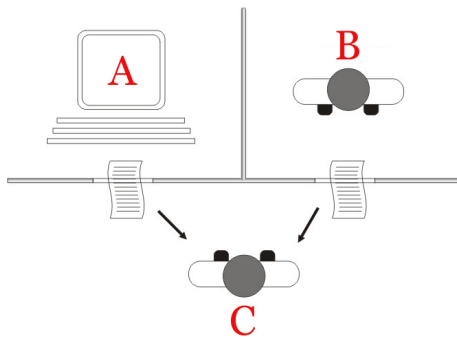
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The brain processes information AND is active:

ions, neurotransmitters, neuromodulators, hormones, remodeling synapses, movement, fields.

His 1950 Imitation Game relies on this mistake

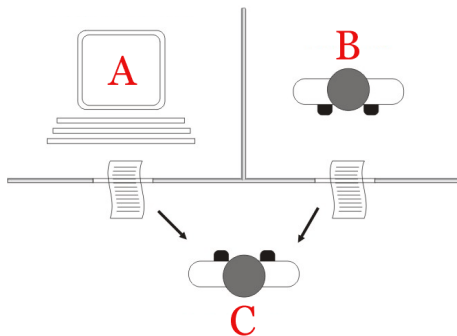
courtesy wikipedia



Turing Test involves just info input & info output.

His 1950 Imitation Game relies on this mistake

courtesy wikipedia



Turing Test involves just info input & info output.

Same with the Chinese Room Argument objection.

Need new: Active + Information Machines that...

Process info AND produce definite physical effects

Examples: Brains, organs, robots, ribosomes ...

Need new: Active + Information Machines that...

Process info AND produce definite physical effects

Examples: Brains, organs, robots, ribosomes ...

Computers? (require user)

Artificial neural networks? (some act)

A + I Machines may be better than Turing machines:

Brains, etc., compute and act simultaneously.

Robots have a communication bottleneck.

A + I Machines may be better than Turing machines:

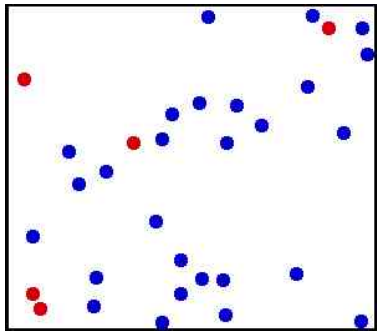
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Statistical mechanics might help us figure out the brain

Statistical mechanics might help us figure out the brain

microscopic first principles \rightsquigarrow zoom out \rightsquigarrow MACROSCOPIC STATES



Courtesy Greg L and Digital Vision/Getty Images

Outline

1. What is the microscopic unit of computation?
2. Unit properties: information & computation.
3. Consequences of this new approach.

What are microscopic computational units in the brain?

Neurons are mesoscopic & complicated.

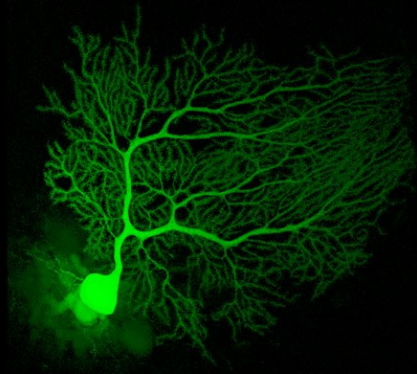
McCulloch-Pitts and von Neumann claimed:
neuron \simeq vacuum tube (wrong)

What are microscopic computational units in the brain?

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If they were right, the iPhone could model a brain.



S. Grant: a synapse is a computer $>$ vacuum tube

So a neuron $\geq 10^4$ synapses \gg vacuum tube

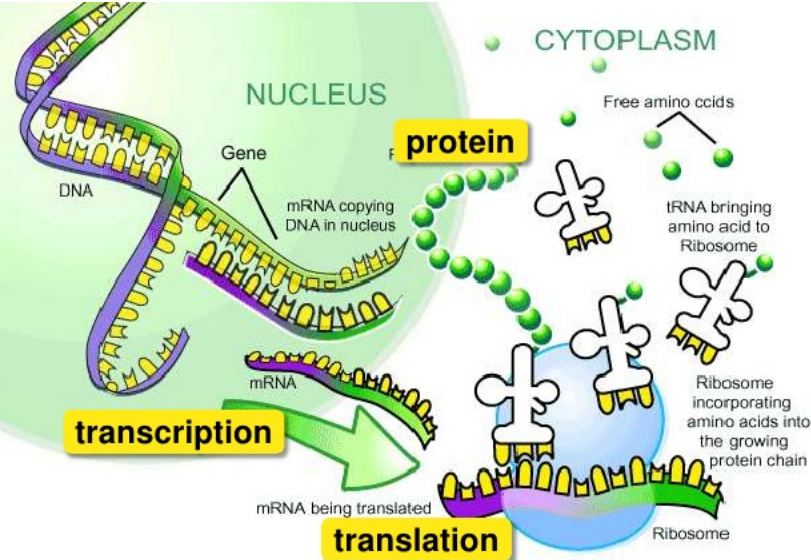
Instead?

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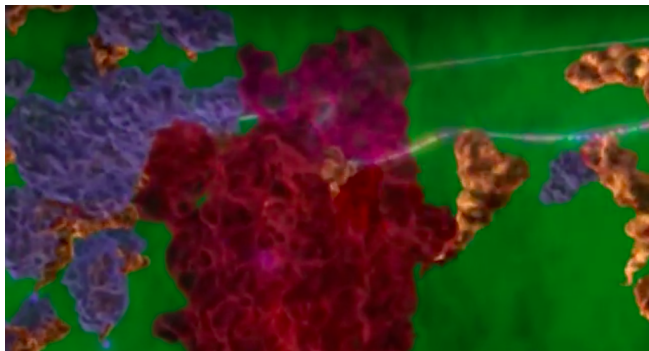
Instead? Proteins have similar complexity to vacuum tubes, and ribosomes help make proteins.

Ribosomes synthesize proteins from mRNA genes



How to model a ribosome mathematically? (Take 1)

Detailed: Molecular Dynamics. But movie:



Reductive: TASEP. No information. But theory.

How to model a ribosome, Take 2

Caetano-Anollés & Caetano-Anollés 2015:
As a Universal Turing Machine in the cell.

As a finite-state automaton.

How to model a ribosome, Take 2

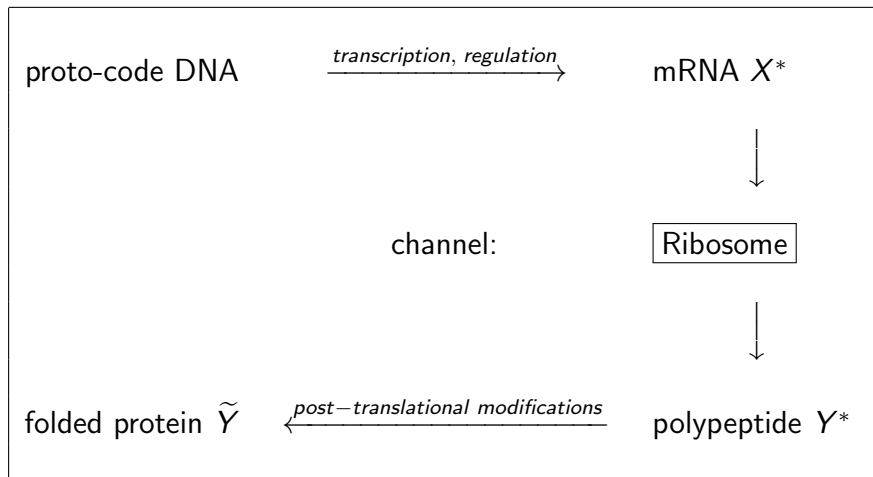
Caetano-Anollés & Caetano-Anollés 2015:
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Reductive and info/computational, not active.

AIM above pure info, but first info details...

The ribosome as an information channel



The ribosome operates under its Shannon capacity

Theorem (Inafuku-K.-Osuagwu 2019):

Ribosome's Shannon capacity is ≥ 266 bits per second. Cf: observed rates $\simeq 24$ to 120 bits per second.

Proof uses the uniform distribution.

This is why ribosomes are $\sim 99.99\%$ accurate.

Enzymes operate under their Shannon capacities

Theorems (D. Inafuku 2019): Theoretical capacities & observed rates (units are bits/sec):

| Enzyme name | Cap. \geq | Actual rate |
|--------------------|-------------------------------|--------------------|
| DNA polymerase | 1554.2 | 1498 |
| RNA II polymerase | 191.9 | 160 |
| KcsA ion channel | $3 * 10^7$ | $2 * 10^7$ |

Notation to get beyond pure computation

Input mRNA tape X^* is string of codons in $\Sigma := \{A, C, G, U\}^3$

$\Sigma^* := \{\text{words with letters in } \Sigma, \text{ incl. empty word } \epsilon\}.$

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Y^* contains info and is also a physico-chemical structure.

Turing defined a 2-tape automatic a-machine

Consists of six parts: $M := \langle Q, \Sigma, \Delta, \delta, q_0, F \rangle$

- ▶ State space Q
- ▶ Σ input alphabet, and input program $X^* \in \Sigma^*$
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- ▶ Initial state q_0
- ▶ Final/accepting states $F \subset Q$
- ▶ Transition function (rules about updating state/tape)
 $\delta : (Q \setminus F) \times \Sigma \rightarrow Q \times \Sigma \times \Delta \times \{L, R\}$

We define an automatic biochemical abc-machine

An **abc-machine** $M_{T,A} = \langle Q, \Sigma, \Delta, \delta, q_0, F, T, A \rangle$ has:

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- ▶ Δ output **blocks**, and Y^* is **block-string** output
- ▶ **T** a random stopping time, when program X^* **halts**
- ▶ **A** an auxiliary machine, optimizing a bio-chemical energy functional, e.g., protein-folding

Diagram for the ribosome as an abc-machine $M_{T,A}$

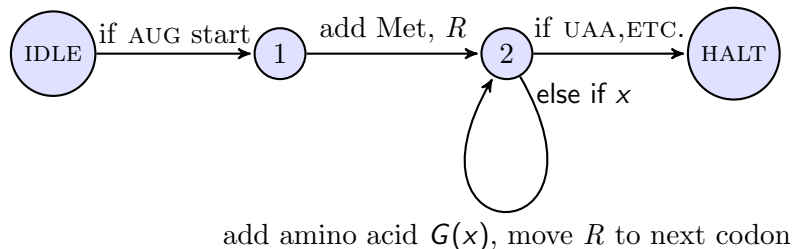


Figure: The (basic!) transition function δ diagram for a ribosome.

If the RNA degrades at time $\mathbf{T}(X^*, \omega)$, transition to HALT.

Auxiliary machine \mathbf{A} folds the polypeptide: $Y^* \rightsquigarrow \tilde{Y}$

The ribosome is more than just a Turing Machine (TM)

Observation (K.-Osuagwu 2018): $M_{T,A}$ is equivalent to a 2-tape TM with 2 oracles: protein-folding machine **A** for output \tilde{Y} , and stopping time $\{t < \mathbf{T}\}$.

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The ribosome $M_{T,A}$ is an A+I machine.

Protein product \tilde{Y} is an A+I machine too.

(Modified input pieces can act too: e.g., ATP, GTP.)

Better than TMs on Decidability/Computability

Abc-machines avoid the Halting Problem by stopping at **T**.

Better than TMs on Decidability/Computability

Abc-machines avoid the Halting Problem by stopping at **T**.

A non-deterministic Turing Machine (NDTM) can approx. but not simulate **T**, which is truly random & non-computable.

Abc-machines are possibly higher complexity than TMs

If protein folder **A** is NP-complete, then it is in class **NP**.

Self-avoiding random walk model of **A** is **NP**-complete
(Berger and Leighton 1998)



Consequences for the brain

Individual computational units are more than TMs.

Consequences for the brain

Individual computational units are more than TMs.

Conjecture: Brain is more than a TM. Embodiment.

When can we model the whole brain? (Weak AI)

The brain has $\sim 10^{11}$ cells (neurons and glia).

Each cell has $\sim 10^6$ ribosomes and $\sim 10^7$ proteins.

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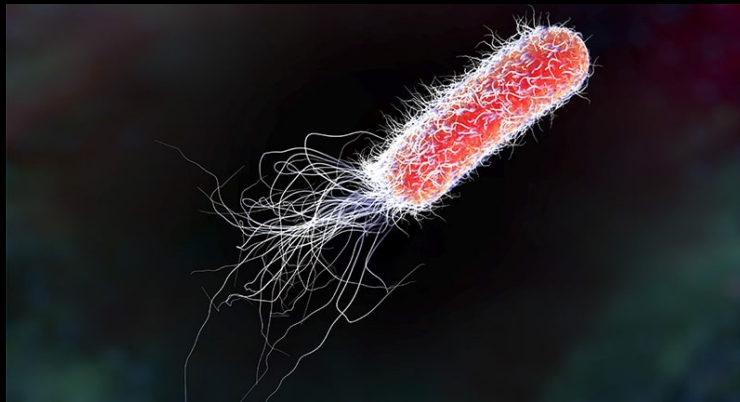
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2-week weather modeling may require zettascale.

Why are artificial “neural networks” successful?

Maybe not because they're like brains, but like something else...

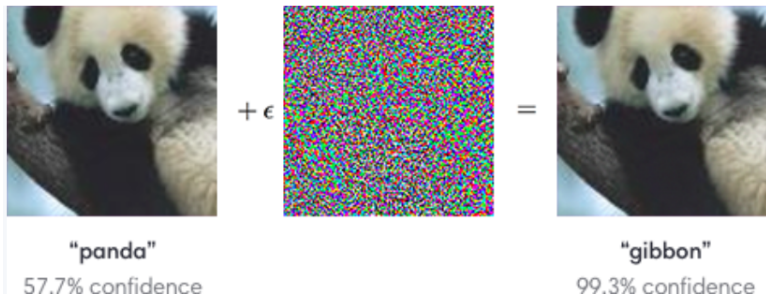


Maybe artificial neural networks are successful...

Because they're like biochemical networks in *E. coli*, which can basically do Stochastic Gradient Descent.

But artificial neural networks fail spectacularly

An adversarial example:



Courtesy Open AI

Key takeaway: “Neural network” is a misnomer

Nodes (neurones) are simpler than real neurons.

Better: “Biologically inspired” networks.

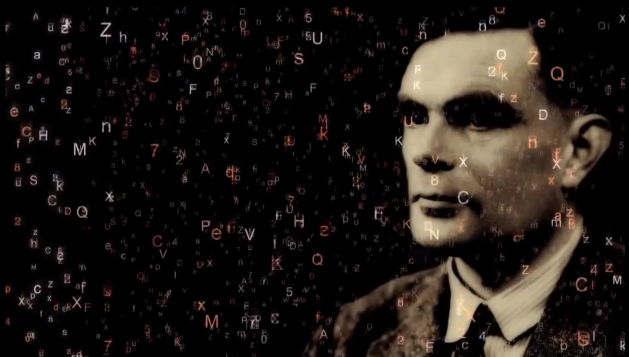
More takeaways

Biological computation is more sophisticated than we thought.

More takeaways

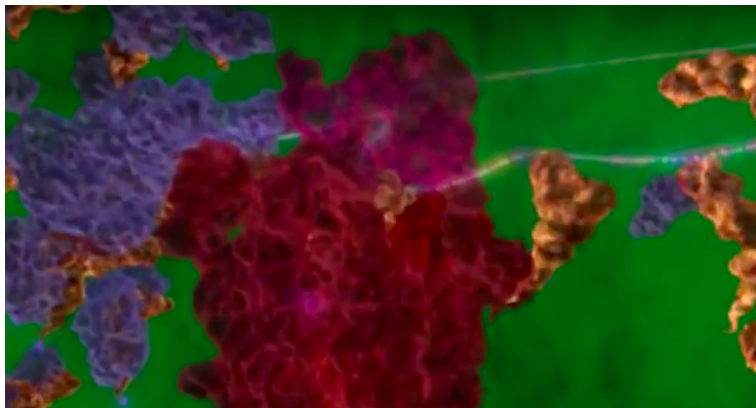
Biological computation is more sophisticated than we thought.

True AI may be harder than we thought.



Thanks

NSF CAREER Award DMS-1254791, Simons Sabbatical Fellowship



faculty.math.illinois.edu/~kkirkpat/

What about DNA computing?

