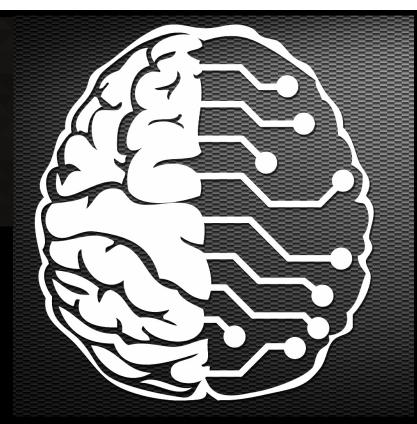
Brain Backups



Next Generation Brain Hacking, Security, and Interfaces

Prof. Russell Hanson

Locard Cybersecurity Summit Istanbul, Turkey

21 May 2016

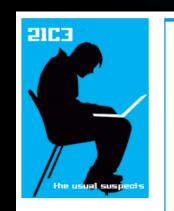


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THIS GUY WANTS TO HELP YOU DOWNLOAD YOUR BRAIN

Hanson

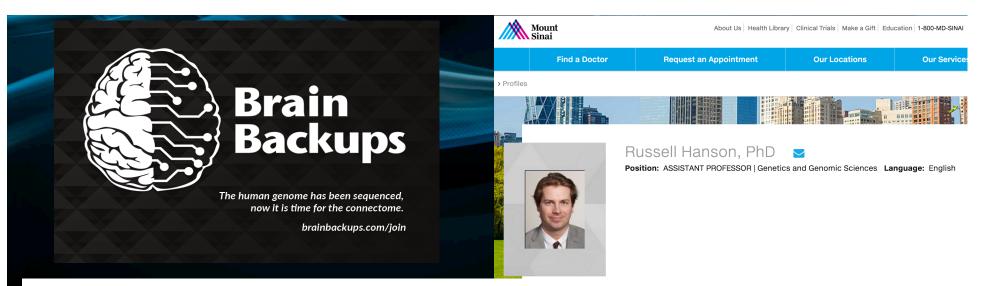






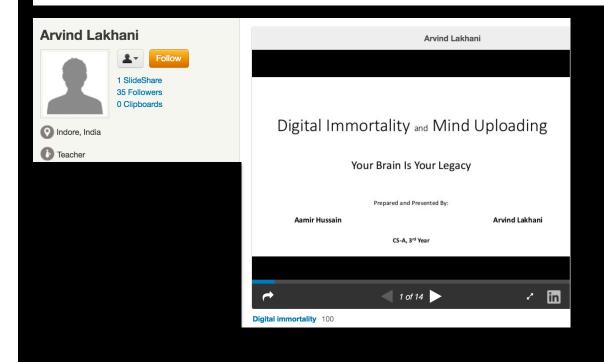


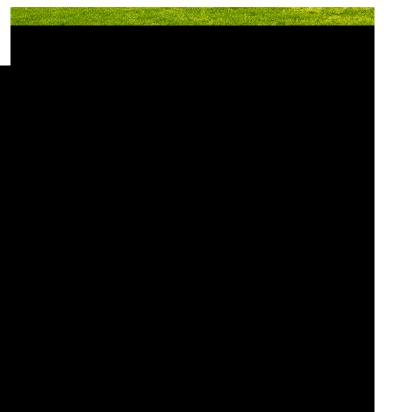




Brain Backups | Your Mind Is Your Legacy www.brainbackups.com/ -

Brain Backups, a developmental stage startup, is pursuing one of the biggest scientific and technological discoveries of our time: a map of every connection in ...





Firstly, what is a Brain Backup, and what is it good for?

- Brain Backups[®] is a trademarked name for the *connectome*, or the network of neural connections in your brain, and some metadata about those neurons.
- We see that the *connectome* has many parallels to the human genome, which has revolutionized science and medicine down to the \$1000 genome, and the \$150 gene chip (23andMe). Genomic tech is a trillion USD business.
- Why would anyone want such a thing, really?
 - <u>Health</u>: Alzheimer's, Parkinson's, dementia, psychological disorders can be better understood and treated. "Digital immortality"/"Singularity"?
 - <u>Education</u>: Image someone's brain before university and after university, use that info to transfer the knowledge to another brain, or selected pieces of that knowledge.
 - <u>Technology</u>: Use the brain image "Brain Backup" to inform a machine, minion, AI, robot, other person, whatever of a task that needs doing. Copy these minions, do more stuff, spend less time training/teaching/programming.
 - <u>Entertainment</u>? Pretty limitless.
 - <u>Business</u>: Truly limitless. The neural modem described below, if successful will enable telepathic-like interactions and communications. A multi-billion dollar business right there.



I don't believe you, this sounds like science fiction.

3600 seconds/hour * 40/60 hours = 2400 the factor from real-time human brain emulation, let's call it 2,000X (ca. 2014).

<u>Time to increase computing power 2,000X?</u> Introduced by Intel on <u>April 1, 1974</u>, the 8080 had an 8-bit architecture, 6,000 transistors, clock speeds of 2-MHz.

In <u>1985</u>, Intel 386 80386SX was available in clock speeds of 16MHz, 20MHz, 25MHz, and 33MHz.

On <u>November 20, 2000</u>, Intel released the Willamette-based Pentium 4 clocked at 1.4 and 1.5 GHz.

The Telegraph

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Supercomputer models one second of human brain activity

The most accurate simulation of the human brain ever has been carried out, but a single second's worth of activity took one of the world's largest supercomputers 40 minutes to calculate





The simulation will help scientists create more accurate models in future Photo: Alamy

By Matthew Sparkes

10:04AM GMT 13 Jan 2014



Print this article

Show me a connectome

• 'Kay.

https://github.com/openworm/CElegansNeuroML/blob/master/CElegansNeuronTables.xls

Origin	Target	Туре	Number of Connections	Neurotransmitter
ADAL	ADAR	GapJunction		1 Generic_GJ
ADAL	ADFL	GapJunction		1 Generic_GJ
ADAL	AIBL	Send		1 Glutamate
ADAL	AIBR	Send		2 Glutamate
ADAL	ASHL	GapJunction		1 Generic_GJ
ADAL	AVAR	Send		2 Glutamate
ADAL	AVBL	Send		4 Glutamate
ADAL	AVBR	Send		7 Glutamate
ADAL	AVDL	Send		1 Glutamate
ADAL	AVDR	GapJunction		2 Generic_GJ
ADAL	AVEL	Send		1 Glutamate
ADAL	AVJR	Send		5 Glutamate
ADAL	FLPR	Send		1 Glutamate
ADAL	PVQL	GapJunction		1 Generic_GJ
ADAL	RICL	Send		1 Glutamate
ADAL	RICR	Send		1 Glutamate
ADAL	RIML	Send		3 Glutamate
ADAL	RIPL	Send		1 Glutamate
ADAL	SMDVR	Send		2 Glutamate
ADAR	ADAL	GapJunction		1 Generic_GJ
ADAR	ADFR	GapJunction		1 Generic_GJ



What do I do with a connectome? Simulate it.

nest::

The Neural Simulation Tool

current release: nest v2.10.0

>> Download Release v2.10.0 (Dec 31 2015)

DOI 10.5281/zenodo.44222 license GPLv2+

NEST 2.10.0 contains 303 repository commits by 25 developers since v2.8.0. The most notable changes over v2.8.0 are:

- Support for simulations of gap junctions (see Jan Hahne et al., 2015)
- Framework for structural plasticity (see Markus Butz et al., 2013 and Markus Butz et al., 2014)
- Full support for the K computer (just in case you found one under your Christmas tree ;-))



C Elegans Neurorobotics Timothy Busbice interintelligence@gmail.com http://www.connectomeengine.com



You are your connectome



Without a brain, no non-plant organism larger than a single cell would be able to respond to its environment in any way other than that dictated by physics and simple, binary responses. The entire sum of who you are resides in the activity of your brain.

Only recently have we had any ability to **understand** the complexity of the brain. The Human Connectome Project Consortium is elucidating neural circuits or pathways in the brain and sub-organ structure, and interconnectivity between brain regions, to understand the design and function of the connectome.

Quantifiably, a connectome is a 3 dimensional mapping of all the "wired" neural connections within a brain. Living connectomes are highly dynamic – an individual's varies continuously throughout their lifetime. Your connectome today is different from when you were a child – and its structure is directly related to your previous connectomic configurations.





Question: How big is a connectome (in bytes)?

The human brain contains about ~100 billion nerve cells, or neurons. On average, each neuron is connected to other neurons through about 10 000 synapses. The actual figures vary greatly, depending on the local neuroanatomy.

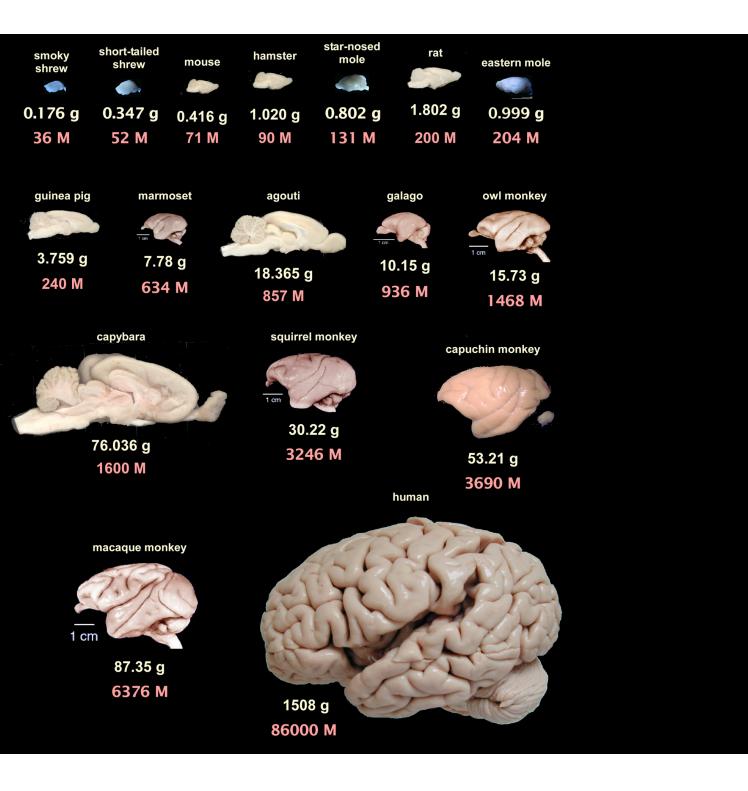
100 000 000 000 neurons * 10 000 synapses = 1 000 000 000 000 000 1 terabyte = 1 099 511 627 776 bytes

1 000 000 000 000 000/1 099 511 627 776 = 909.49 terabytes

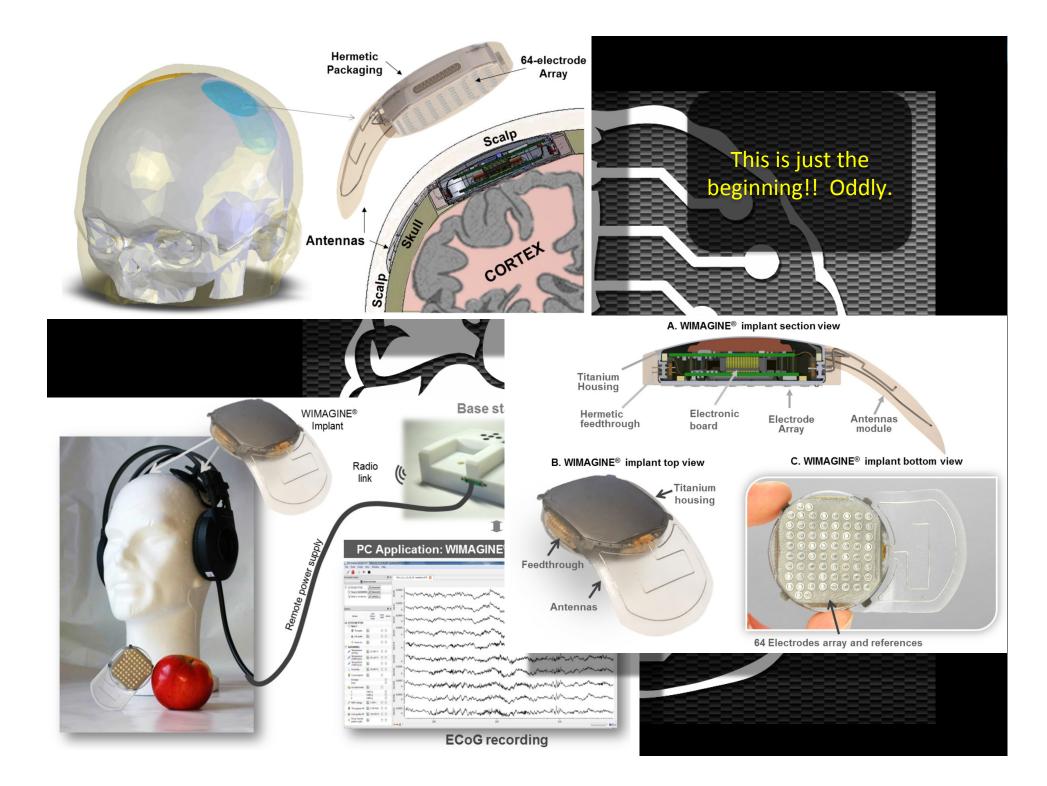
1 terabyte HD storage = \sim \$28.31 \rightarrow storage for all human neurons and synapses \sim \$28.31 * 909.49 = \sim \$25,747.66



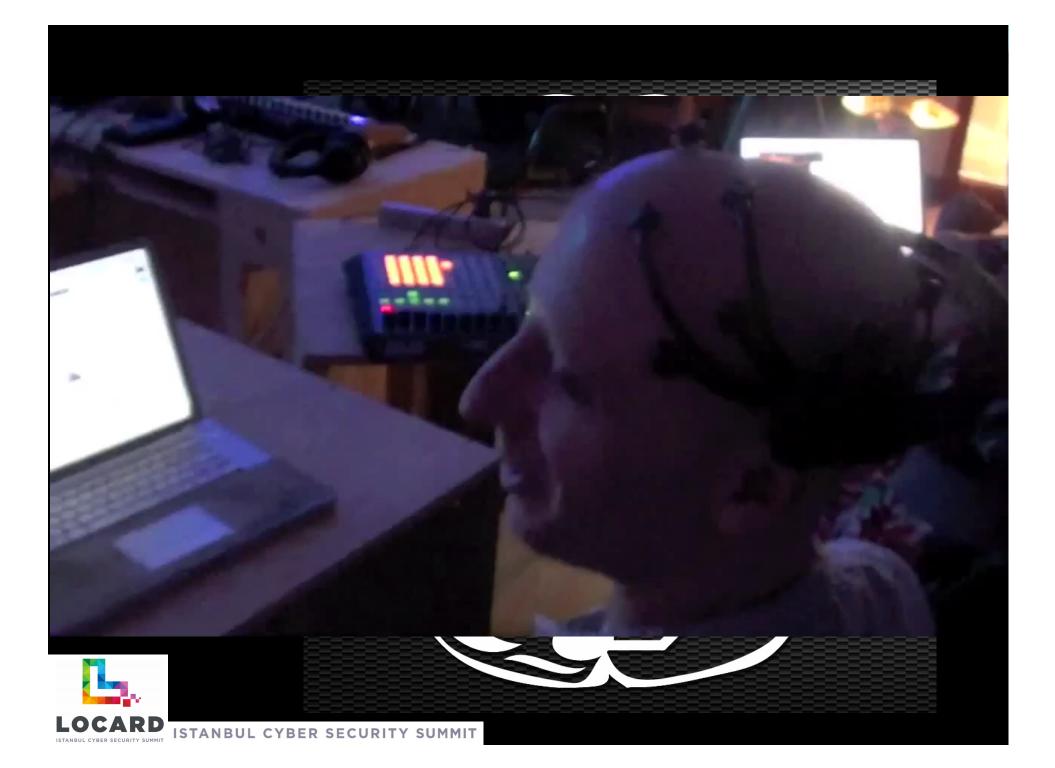
Connectivity: Assuming avg. 500 inputs per neuron, adjacency list is avg. 37•500=18,500 bits≈2kB per neuron. Neuronal type: Assume 10^3 cell types => 10 bits. Configuration: Assume each input synapse has 10^3 states => additional 5,000 bits. Total 3kB•2^37=384TB. Assume ~50% achievable compression ratio. Estimate: 200-300TB.









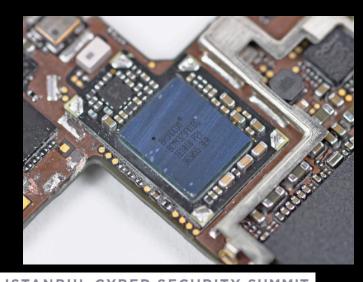


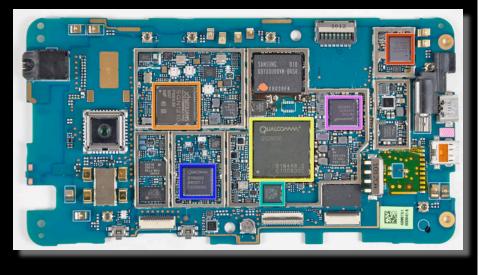
How to make a Wifi connected brain – seems useful, right?

[root@brain ~]# ifconfig brain0 up

Should one use motor cortex? ECOG? Nanodevices like our nanoparticles?

Broadcom BCM4329 chip that powers Wi-Fi, bluetooth on iphone 4, HTC EVO





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<u>Brain Modem</u>: DARPA NESD \$60M – read from 1,000,000 neurons stimulate 100,000 neurons



Neural Engineering System Design Solicitation Number: DARPA-BAA-16-09 Agency: Other Defense Agencies

Office: Defense Advanced Research Projects Agency Location: Contracts Management Office

Notice Details	Packages
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Interested Vendors List

Original Synopsis Jan 21, 2016

11:58 am

Return To Opportunities List

Solicitation Number: DARPA-BAA-16-09 Notice Type: Presolicitation

Synopsis:

Added: Jan 21, 2016 11:58 am

DARPA seeks proposals to design, build, demonstrate, and validate a neural interface platform capable of recording from more than 1,000,000 neurons and stimulating more than 100,000 neurons in proposer-defined regions of the human auditory, visual and somatosensory cortex. The complete system must demonstrate high-precision detection, transduction, and encoding of neural activity.



MOTOR CORTEX and sensory cortices

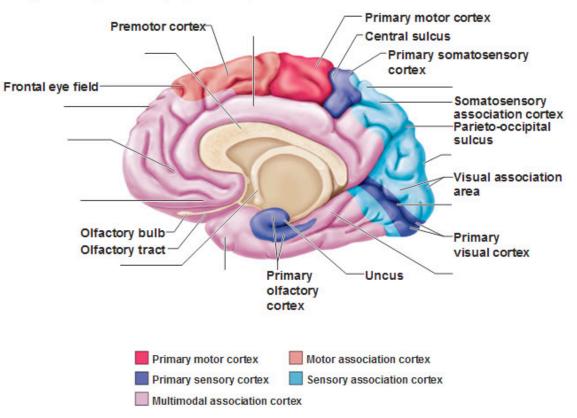


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Functional Areas of the Cerebral Cortex

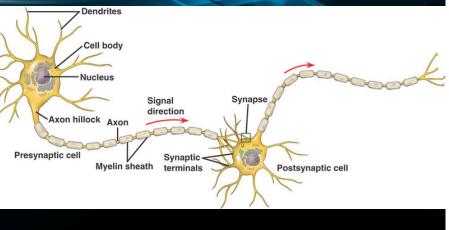
(b) Parasagittal view, right hemisphere



Fundamental components of a neural network

system

- Synaptic weight
- Neurotransmitters
- Long term potentiation (LTP)
- Long term depotentiaion (LTD)



To model a network of LT neurons, assume that their activities at time *t* are given by the *N* variables, $x_1(t)$, $x_2(t)$..., $x_N(t)$ which take on the values 0 or 1, that is, a neuron is either active ("1") or silent ("0"). Then the activities at time t + 1 are given by

Nanoconnectomic upper bound on the variability of synaptic plasticity

Thomas M Bartol Jr^{1*}, Cailey Bromer¹, Justin Kinney^{1,2†}, Michael A Chirillo³, Jennifer N Bourne^{3‡}, Kristen M Harris^{3*}, Terrence J Sejnowski^{1,4*}

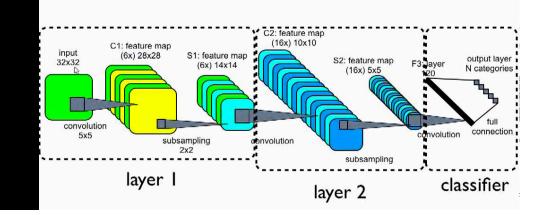
Abstract Information in a computer is quantified by the number of bits that can be stored and recovered. An important question about the brain is how much information can be stored at a synapse through synaptic plasticity, which depends on the history of probabilistic synaptic activity. The strong correlation between size and efficacy of a synapse allowed us to estimate the variability of synaptic plasticity. In an EM reconstruction of hippocampal neuropil we found single axons making two or more synaptic contacts onto the same dendrites, having shared histories of presynaptic and postsynaptic activity. The spine heads and neck diameters, but not neck lengths, of these pairs were nearly identical in size. We found that there is a minimum of 26 distinguishable synaptic strengths, corresponding to storing 4.7 bits of information at each synapse. Because of stochastic variability of synaptic activation the observed precision requires averaging activity over several minutes.

$$x_{i}(t+1) = H\left(\sum_{j=1}^{N} W_{ij}x_{j}(t) - \theta_{j}\right)$$
 (E-1)

where *H* is the Heaviside step function defined by H(u) = 1 for $u \ge 0$ and H(u) = 0 otherwise, W_{ij} is the strength or weight of the synapse between neuron *i* and the presynaptic neuron *j*, and θ_j is the threshold of neuron *i*. For a network of *N* neurons, the synaptic weights W_{ij} form an $N \times N$ matrix, and the thresholds θ_j an *N*-dimensional vector.

Key differences between classical ML/AI/deep learning and biological brain

- Specific networks for specific functions, significance of connectivity between these regions
- "Supervisor"/"teacher" to say when done, to move on to next task
- Highly optimized yet optimization procedure unknown
- Only 86 billion neurons, energy consumption 12 watts
- Highly integrated with peripheral nervous system (somatic nervous system and autonomic nervous system)
- Human intelligence while a general intelligence also performs many distinctly human functions



Convolutional Neural Networks

Connectome Mapping Today

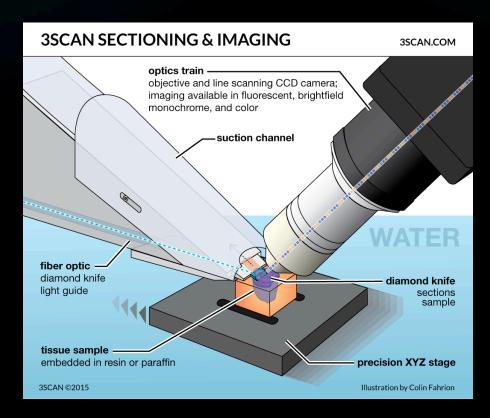


The current methodology of mapping the connectome today relies on imaging of the neurons in a brain, in one way or another. All current methods are:

Highly Destructive Highly Invasive or Very Low Resolution

Don't destroy the thing you want to image!!





Using genome sequencing for connectome sequencing

Moore's Law

Jul-09

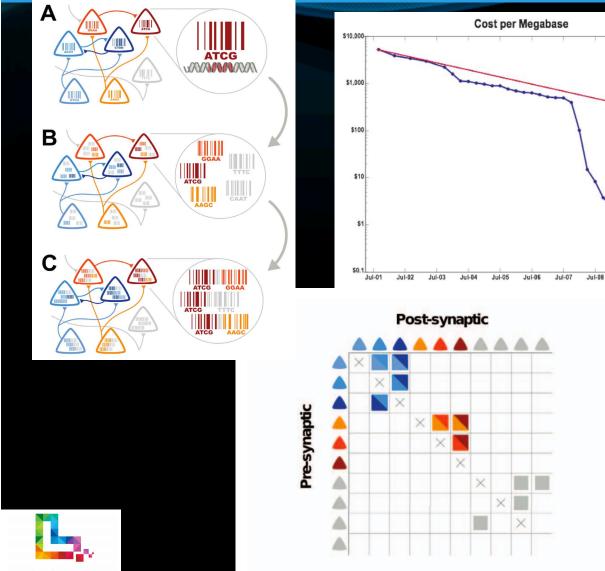
Jul-10

ATCG

GAD

Jul-11

Sequencing Cost



В

ATCG TYN AMAR XYXXXX

SOM

DOP

CTAG

It's still

destructive!!

8888

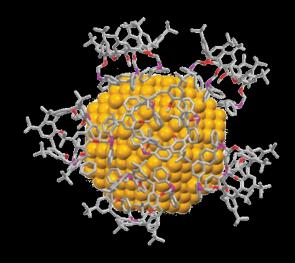
LOCARD **ISTANBUL CYBER SECURITY SUMMIT** ISTANBUL CYBER SECURITY SUMMIT

Novel, non invasive, *in vivo* Imaging Methodology

Non-destructive non-invasive imaging of neural targets to map tissue structure and visualize dynamic biochemical processes at sub-second timescales from 1mm to 300nm. Nanoparticle imaging-agents may be "barcoded" for MRI and dual mode/ spectral CT.

Design: Ligand + Contrast-providing agent = Specific targeted contrast particle/agent

AptaMark: Targeted RNA aptamers + gold nanoparticles = Specific targeted contrast particle/agent



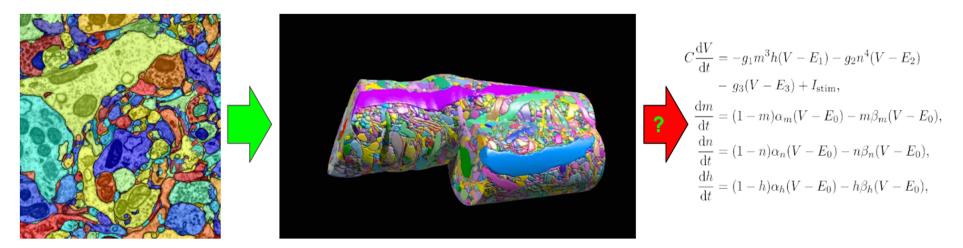




Quoting Randal Koene...



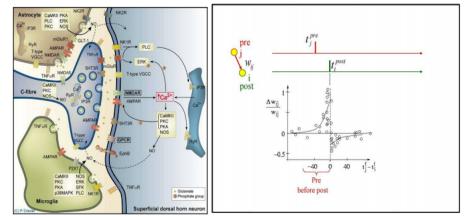
Where are we now - on reconstructing brains?



We lack information to infer function from structure

We **can** do direct functional system identification (see hippocampal prosthesis)

Demands better recording & stimulation



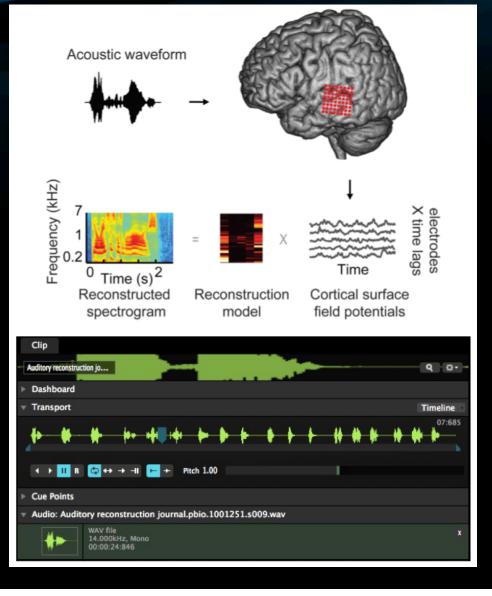
evolutionary patchwork vs the 'language' of the brain

Decoding the visual cortex



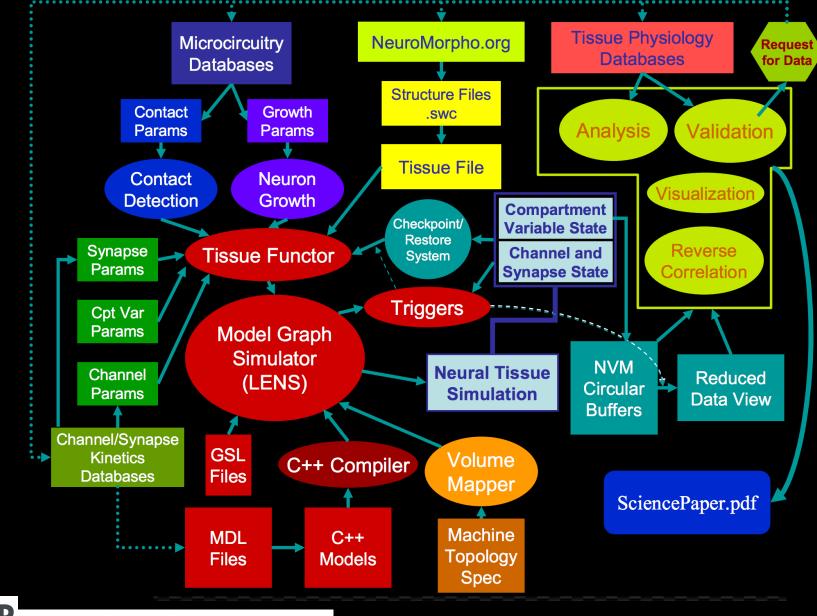
Decoding the auditory cortex







Neural Tissue Simulator Workflow



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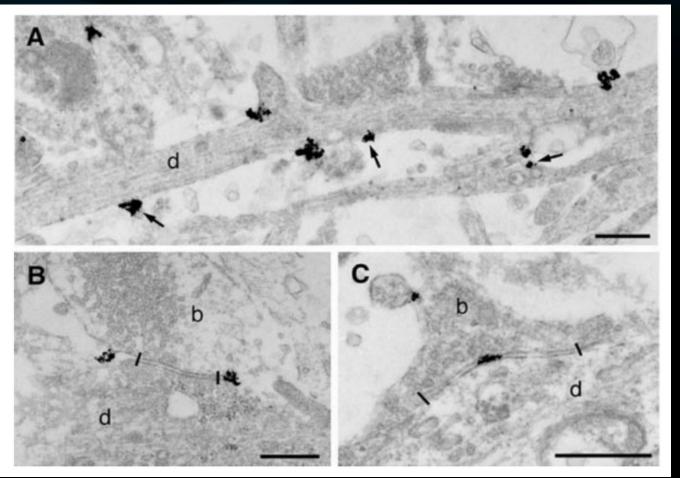


GlyR-a1 subunits imaged using primary antibody (mAb2b), biotinylated anti-mouse Fab fragments, and streptavidin-coated quantum dots.



EM resolution vs. nanoCT

Fig. 4. Transmission EM detection of QD-GlyRs. QD-GlyRs detected on the dendritic surface and associated with extrasynaptic membranes (arrows) (**A**), at the periphery of synapses (**B**), and within the synaptic cleft (**C**). d, dendrites; b, synaptic boutons. The edges of the synaptic clefts are outlined in (B) and (C). Scale bars, 500 nm.





Neurological Imaging Targets



Uniprot accession Subtype Туре number Adrenergic For memory encoding: α1A, α1b, α1c, α1d α1A ADRA1A P35348 ADRA1B a1b P35368 AMPA-R a1c ADRA1C Q7KYZ9/Q6LD06 Exclusive glutamate, excitatory, α1d ADRA1D P25100 α2a, α2b, α2c, α2d α2a ADRA2A P08913 Na+ influx ONLY, hetero OR a2b ADRA2B P18089 homo-tetramer, FAST ADRA2C a2c P18825 a2d adra2da/adra2db Q8JG70/Q8JG69 NMDA-R **β1** β1, β2, β3 ADRB1 P08588 Glutamate and glycine receptor, <u>62</u> ADRB2 P07550 inhibitory, Ca2+ and Na+ influx, ß3 ADRB3 P13945 GluN1 GluN2 heterotetramer ---Dopaminergic D1, D2, D3, D4, D5 D1 always 2 GluN1 + either GluN2 or DRD1 P21728 GluN3. Has Mg+ in core. SLOW D2 DRD2 P14416 D3 DRD3 P35462 D4 DRD4 P21917 D5 DRD5 P21918 GABAergic GABAA. GABAB1a, GABAB15. GABAB2, GABAC GABAA GABAB1a GABRA1 P14867

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Imaging method (size/time)	Contrast agent	Ligand	Delivery	Indication/ application	Shortcoming
X-ray CT (200-400nm/ 10 mins)	15 nm Au NP	GluR1 aptamer	Insulin, aptamer	Synaptic weights, GluR1 densities, brain mapping	Radiation, expense of machine
<i>u</i>	KI	None	None	Tissue stain	Non-specific
NIR (400nm/ ms)	Upconverti ng NP	Aptamer/ antibody	Aptamer	Surface receptor densities	Toxicity of NPs (?), slower response time
u	Au nanorod	Aptamer/ antibody	Aptamer	Voltage sensitive	Orientation
MR	Paramagne tic NP	Aptamer/ antibody	Aptamer	Region activity, blood flow, tau, synuclein	1-5mm voxel
Fluorescence microscopy (400nm/ms)	15 nm Au NP	GluR1 aptamer	None, surface accessible	Targeting quality control	Serial sectioning, shallow depth, destructive
Electron microscopy (EM)(10 -40nm/∞)	Osmium tetroxide	None	Histological stain	High-res imaging	No biological metabolites, no protein density info

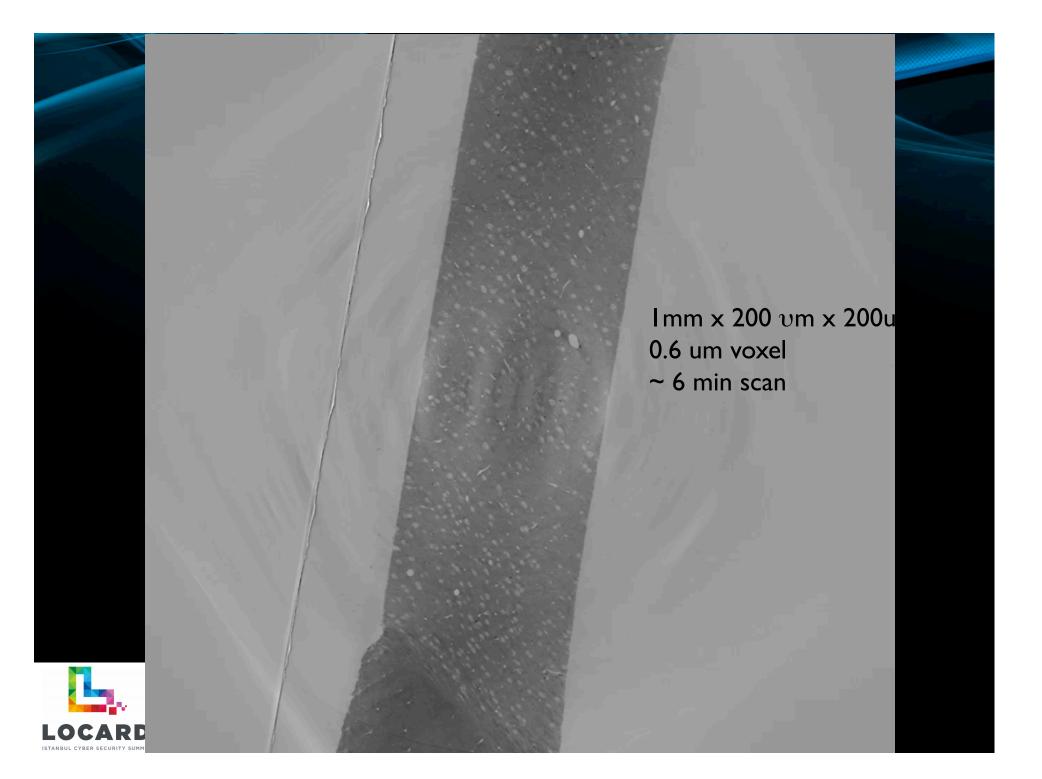
Technical Specifications & Configurations





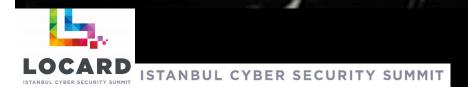
	phoenix nanotom s	phoenix nanotom m		
X-ray tube type	Proprietary open high-power nanofocus X-ray tube, optimized for long-term stability			
	Optional X-ray tube cooling	Internal X-ray tube cooling		
Max. voltage / power	180 kV/15 W			
Target	Tungsten on beryllium (optional tungsten on CVD diamond)	Tungsten on CVD diamond for up to 2 times faster data acquisition at the same high image quality level		
	Transmission target type, rotatable for multiple use (other target materials, e.g. molybdenum on request)			
Filament	Tungsten hairpin, pre-adjusted plug-in cartridges for fast and easy exchange			
Geom. magnification (3D)	1.7 x - 250 x	1.5 x - 300 x		
Detail detectability	Down to 200 nm (0.2 microns)	Down to 200 nm (0.2 microns)		
Min. voxel size	Down to 500 nm (0.5 microns)	Down to 300 nm (0.3 microns)		
Detector type	High-Contrast Detector HCD 120-50, 12 bit / 16 bit, 3 x virtual detector enlargement (max. 6,900 pixel detector width)	Temperature-stabilized high dynamic GE DXR, 14 bit/16 bit, 1.5 x detector enlargement (max. 4,600 pixel detector width)		
Pixels	2,300×2,300	3,072×2,400		
Pixel size	50 µm	100 µm		
Manipulation	Granite based 5-axes manipulator with vibration insulation, precision rotation table on air bearing			
Variable focus detector distance	from 200 mm to 500 mm	from 220 mm to 600 mm		
Max. sample diameter	<1mm to 120mm	<1mm to 240mm		
Max. sample height/weight	150 mm/2 kg (4.4 lbs.)	250 mm/3 kg (6.6 lbs.)		
Sample travel length Y/Z	150mm/300mm	250 mm / 400 mm		
Rotation	0° - 360° x n			







Nanorobots



Ex vivo EM imaging: synapses





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Dvorsky's Complaint #1: Brain functions are not computable

"Is" and "are" are complicated words, semantically.... Computability, in the sense it's used in Dvorsky's complaint, is a mathematical tool used for modeling certain systems. So his claim would be "brain functions cannot be effectively modeled using computable models."

This is an interesting hypothesis, but there is certainly no scientific evidence for it. Furthermore, within the confines of current science, there is no possible way to gather solid scientific evidence for it. The problem is that all scientific data ever gathered, constitutes one large but finite set of bits (i.e a finite set of finite-precision numbers). Any finite set of bits can be modeled computationally. Of course, someone can claim a non-computable model is "better" than any computational model, for a given finite set of bits. But this then becomes a subjective claim, based on aesthetics, or intuition.

Perhaps some future discipline, going beyond the bounds of science as we know it today, will formulate some new sense in which brains fundamentally cannot be computationally modeled. But this vague possibility seems a rather threadbare excuse for rejecting mind uploading.



Ethical implications of brain imaging?

"My personal view about the 'ethical implications' is that it is unethical to NOT permit tetraplegic patients or other injured parties to receive next-generation neural interfaces. And regarding connectome imaging -- again my personal view is that it is unethical to NOT permit patients or other interested parties to image their connectome, just like withholding genetic/genomic data from an oncologist/cancer patient is presently unconscionable. If one is afraid of knowledge, one's head is truly in the sand."

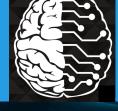


Outstanding problems, areas for outstanding contributions!

- Implanted CPU with database of neural codes
- Deep learning to improve interfaces using ephys spikes to sensorimotor cortex
- AI/ML to trace neurons/axons in image stack data
- <u>Neural Modem</u>: In the next 3-4 years DARPA wants a device that reads from 1,000,000 neurons stimulates 100,000 neurons. Cochlear implant uses only 4 electrodes.







Thanks to the Locard team! And the kind people of Turkey! B)



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