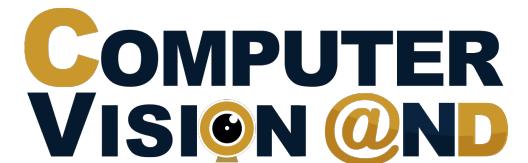


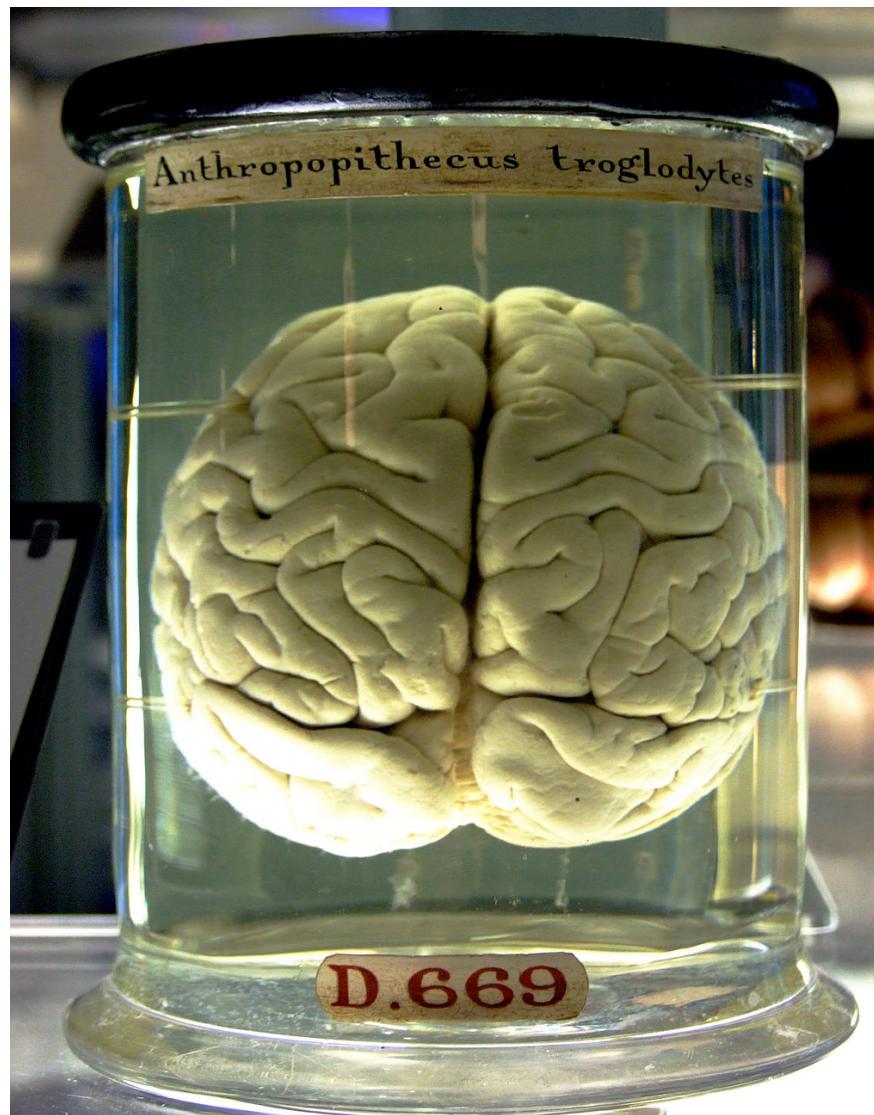
Representational Dissimilarity Analysis as a Tool for Neural Network Model Search

Walter J. Scheirer

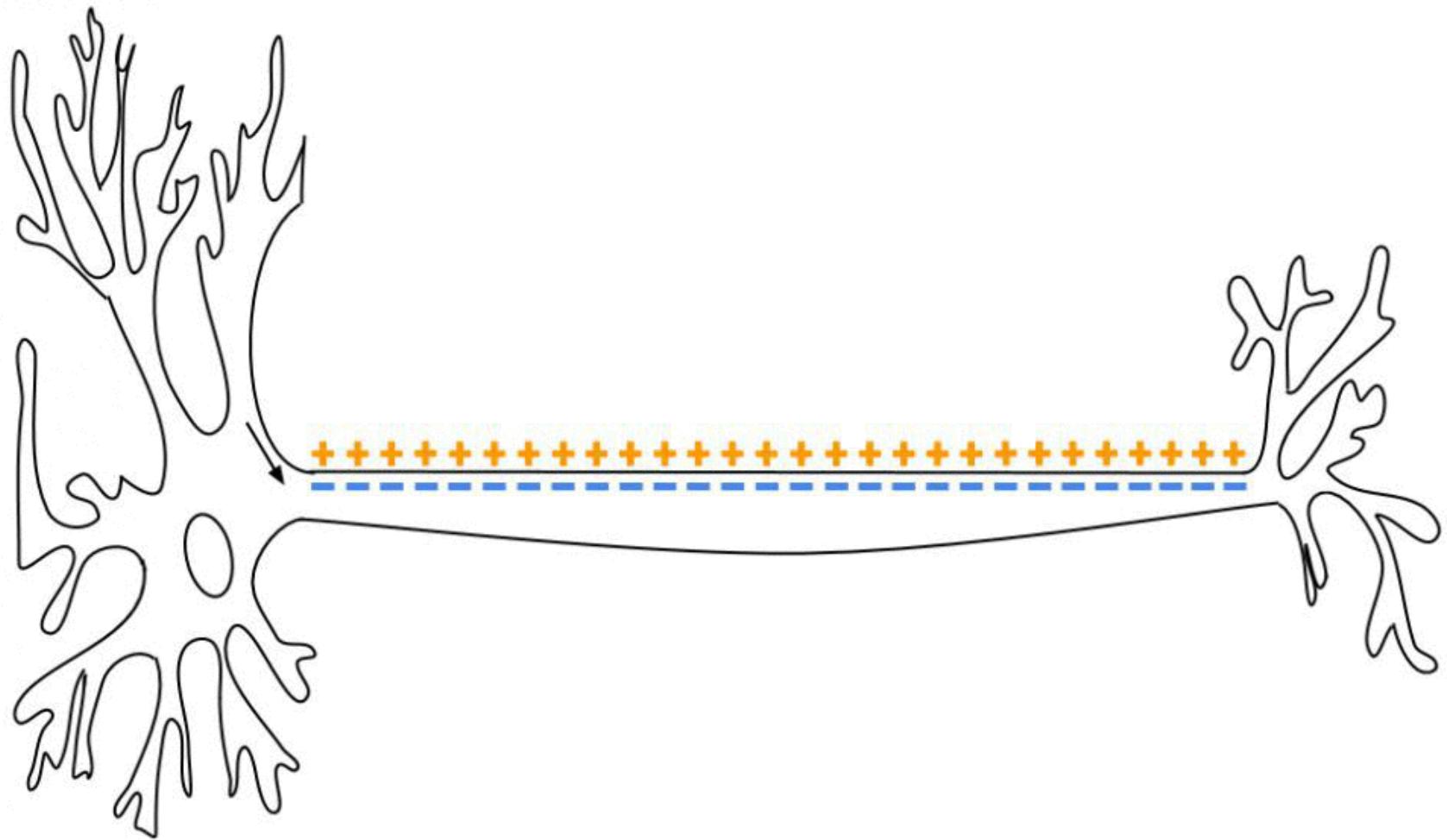
Computer Vision Research Laboratory
Department of Computer Science and Engineering



A model of vision that works:

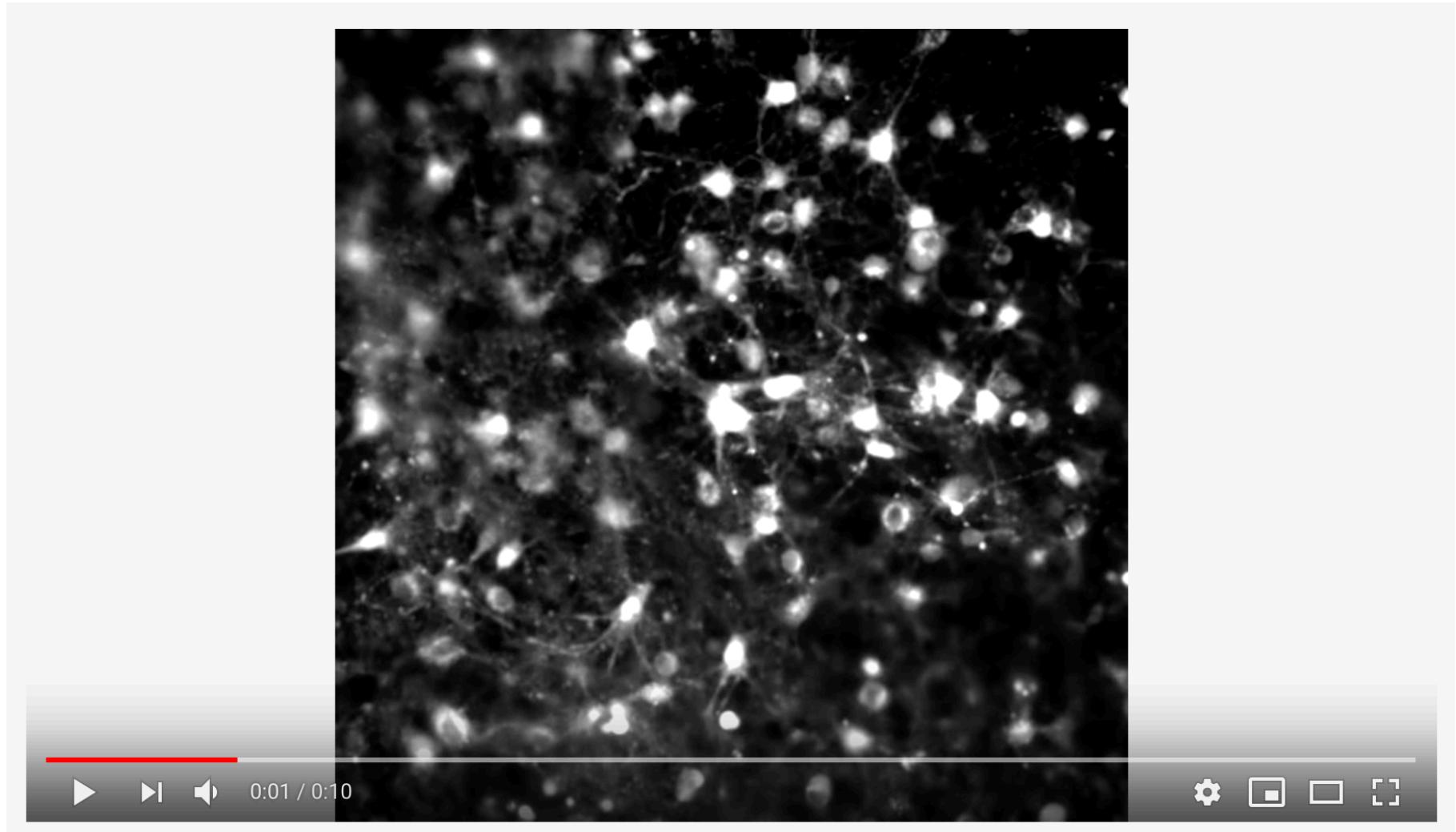


Chimp Brain in a jar CC BY 2.0 Gaetan Lee



Action Potential BY-SA 3.0 Laurentaylorj

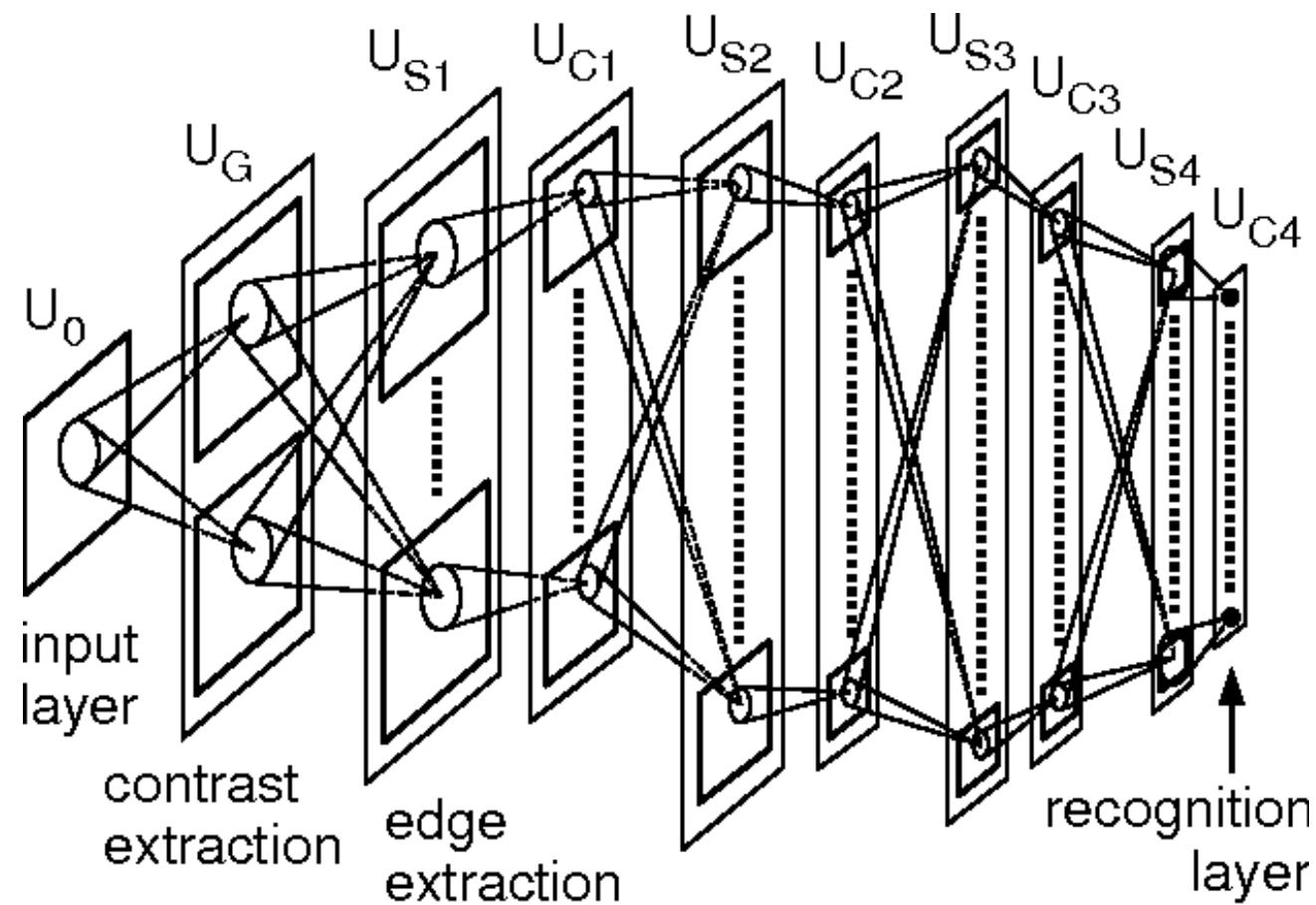
[MakeAGIF.com](#)



<https://www.youtube.com/watch?v=yy994HpFudc>

Hypothesis: networks exhibiting brain-like activation behavior will demonstrate brain-like characteristics, e.g., stronger generalization capabilities.

Fukushima 1979: Neocognitron



Is there any correspondence between activity measured in the brain and activity measured in artificial neural networks?

Monkey performing an object recognition task

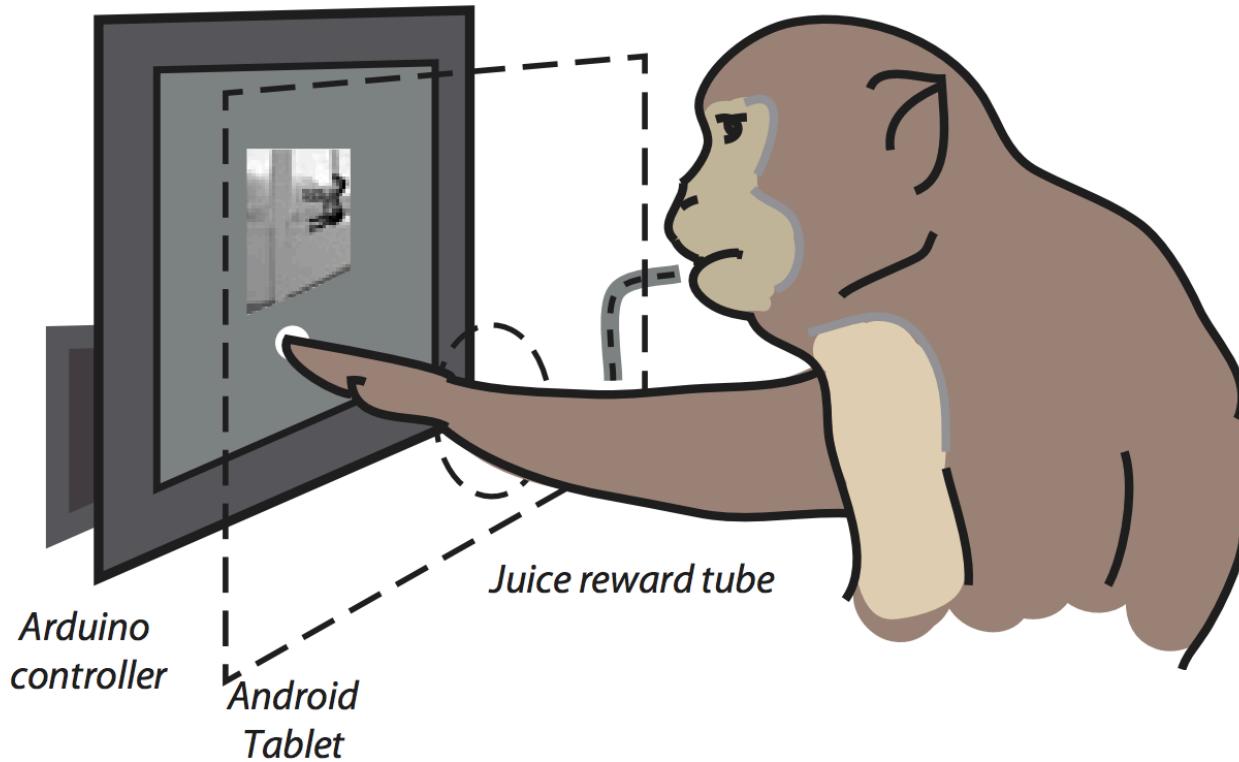
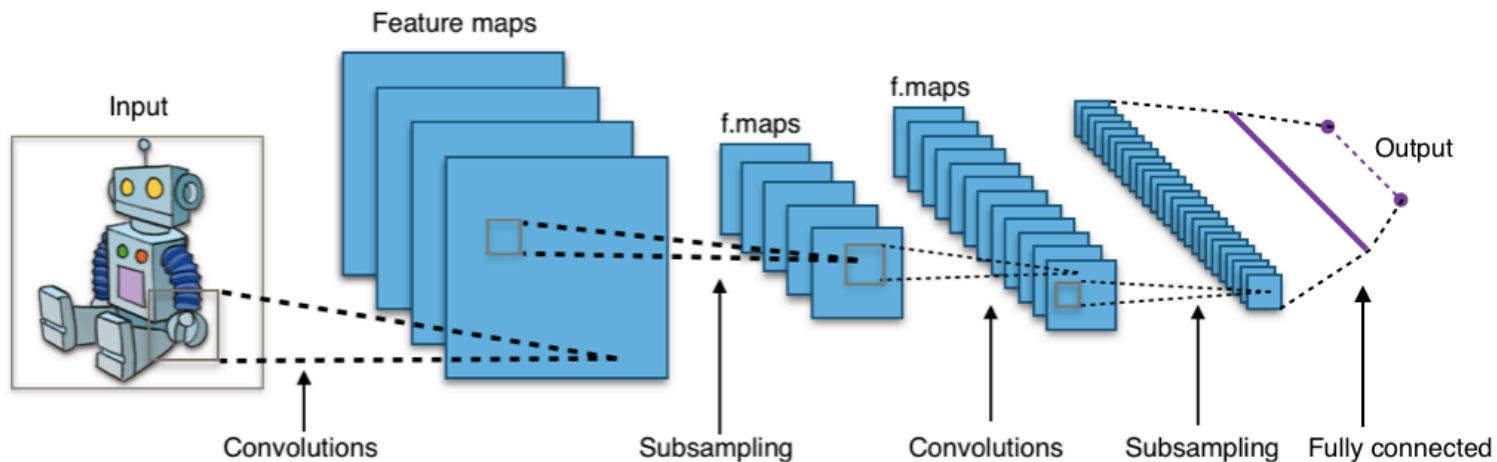


Image adapted from: Rajalingham et al. JNeurosci 2018

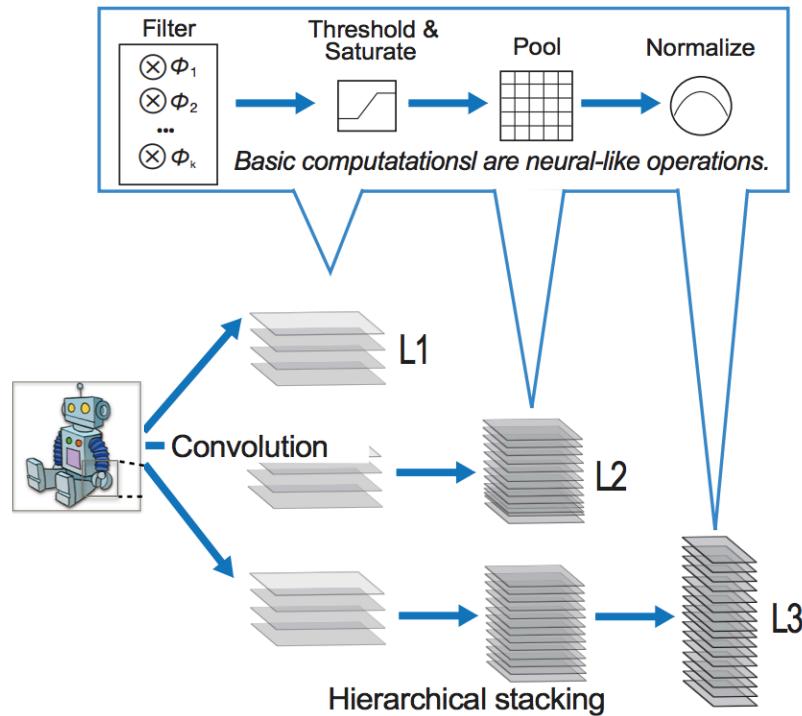
CNN for Object Recognition



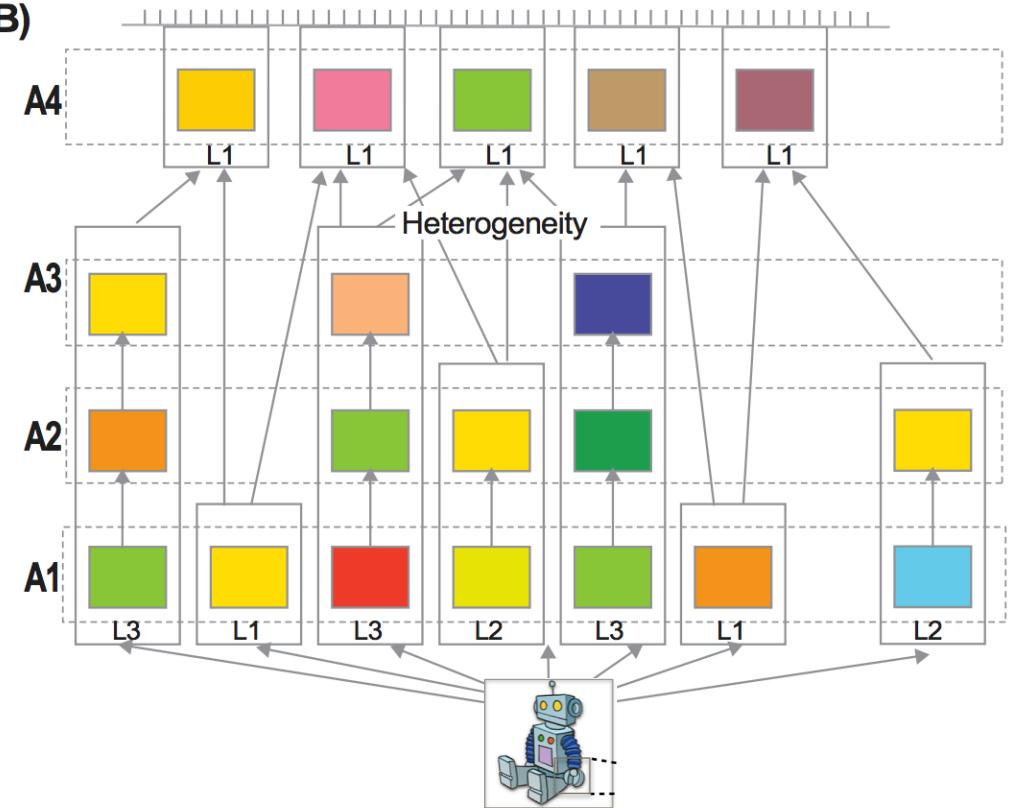
Typical CNN architecture © BY-SA 4.0 Aphex34

Heterogeneous Hierarchical CNN

A) Basic operations

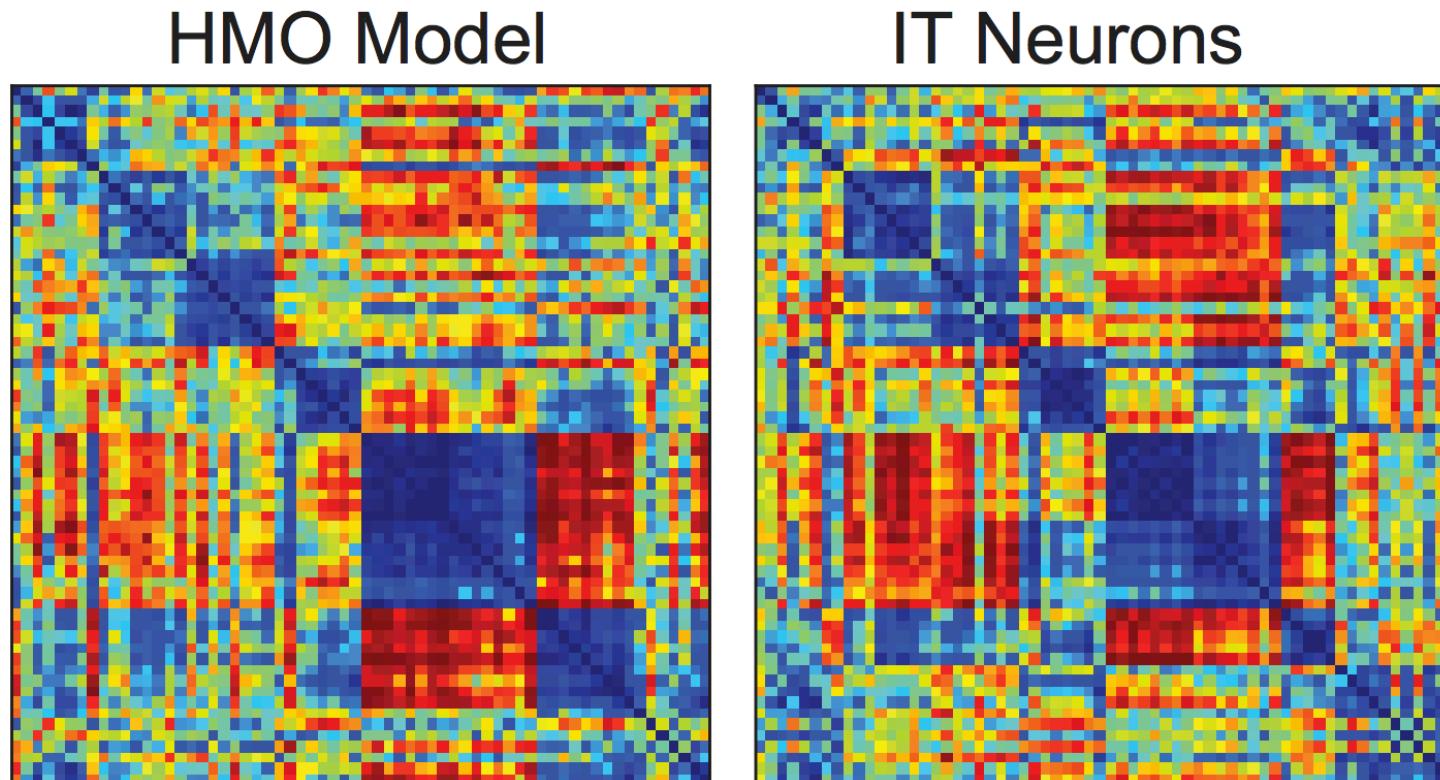


B)



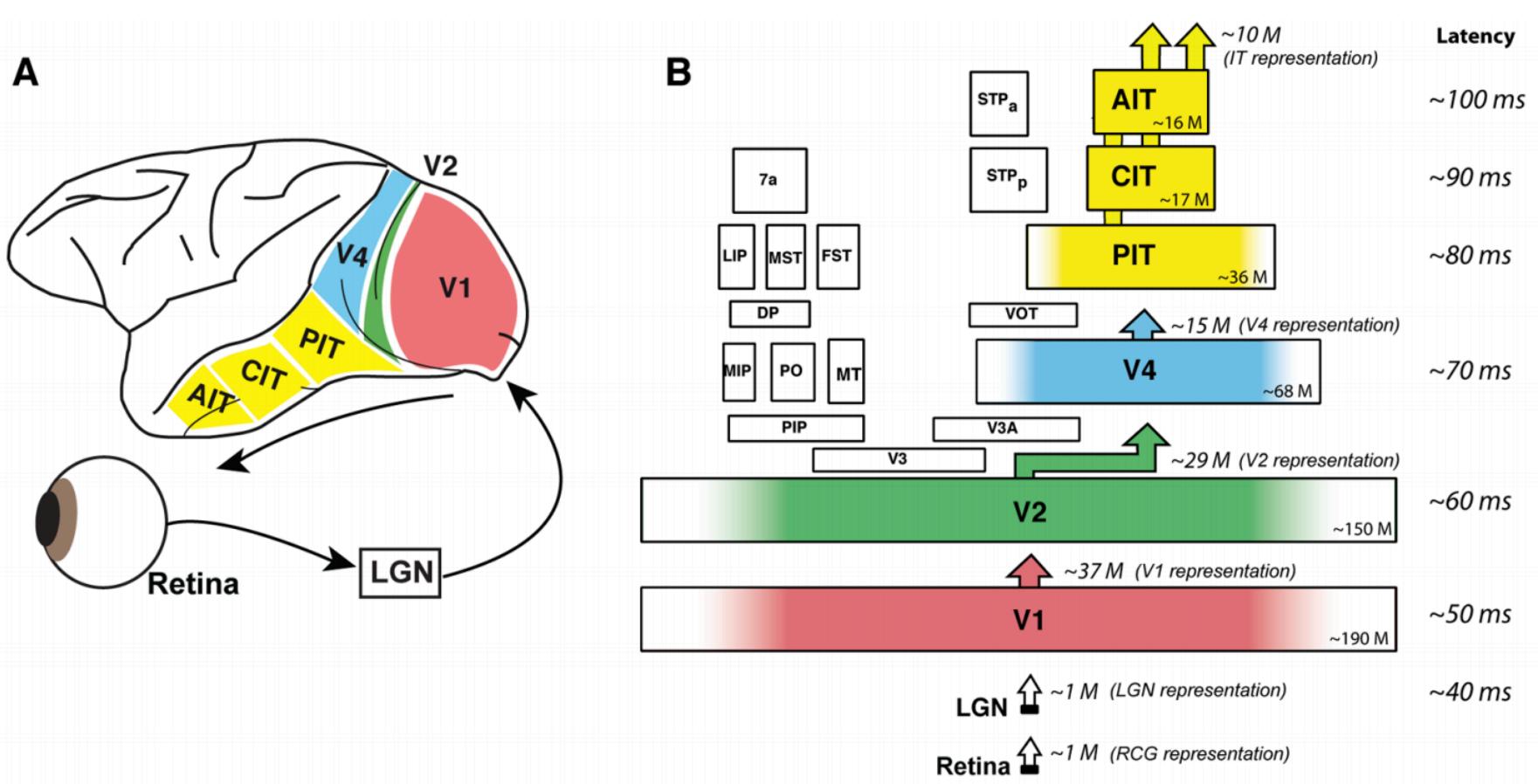
Yamins et al. NeurIPS 2013

Population Responses: Model vs. Brain



Yamins et al. NeurIPS 2013

Where in the brain is area IT?



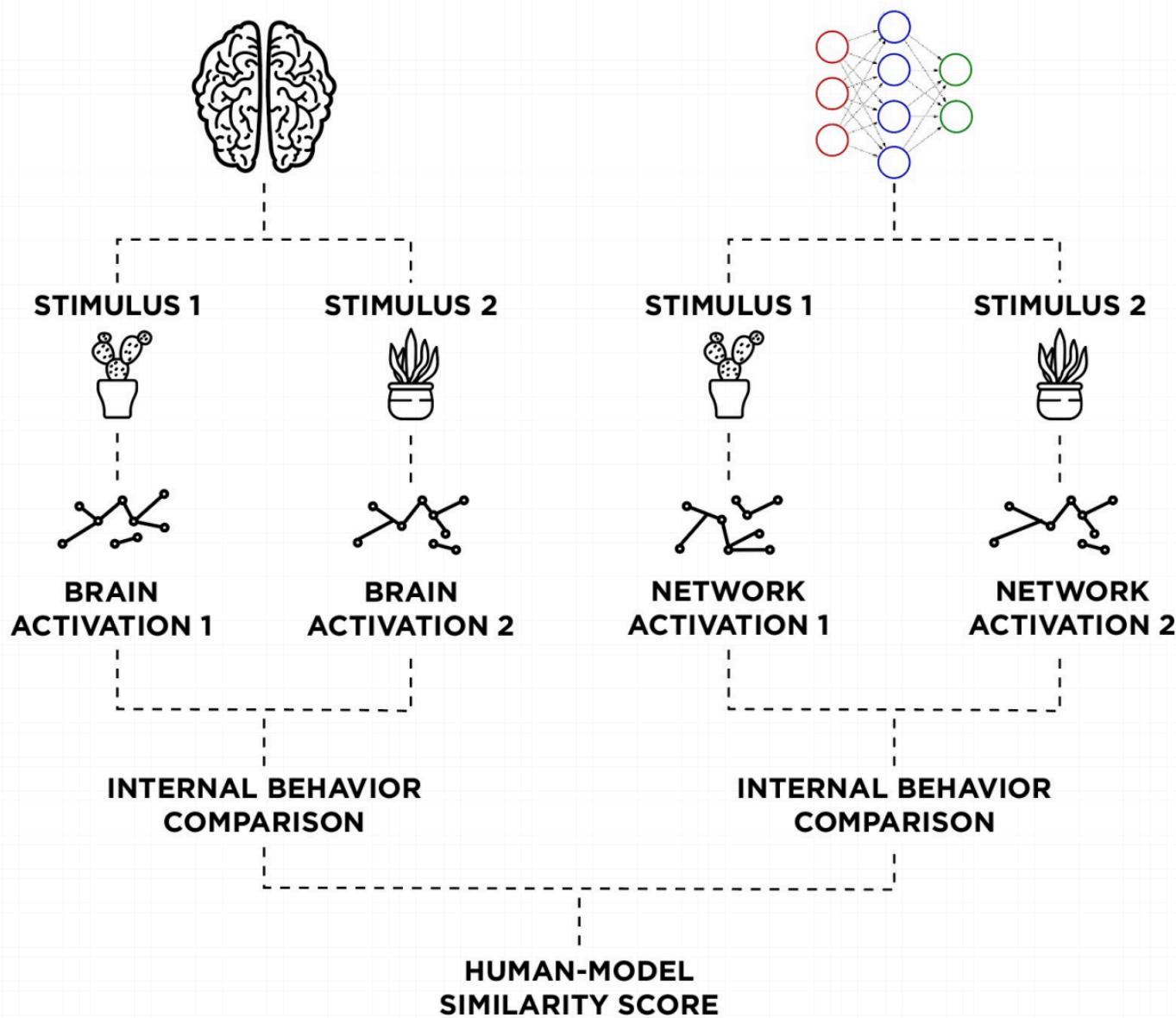
How do we compare the activity in brains
with the activity in artificial neural
networks?



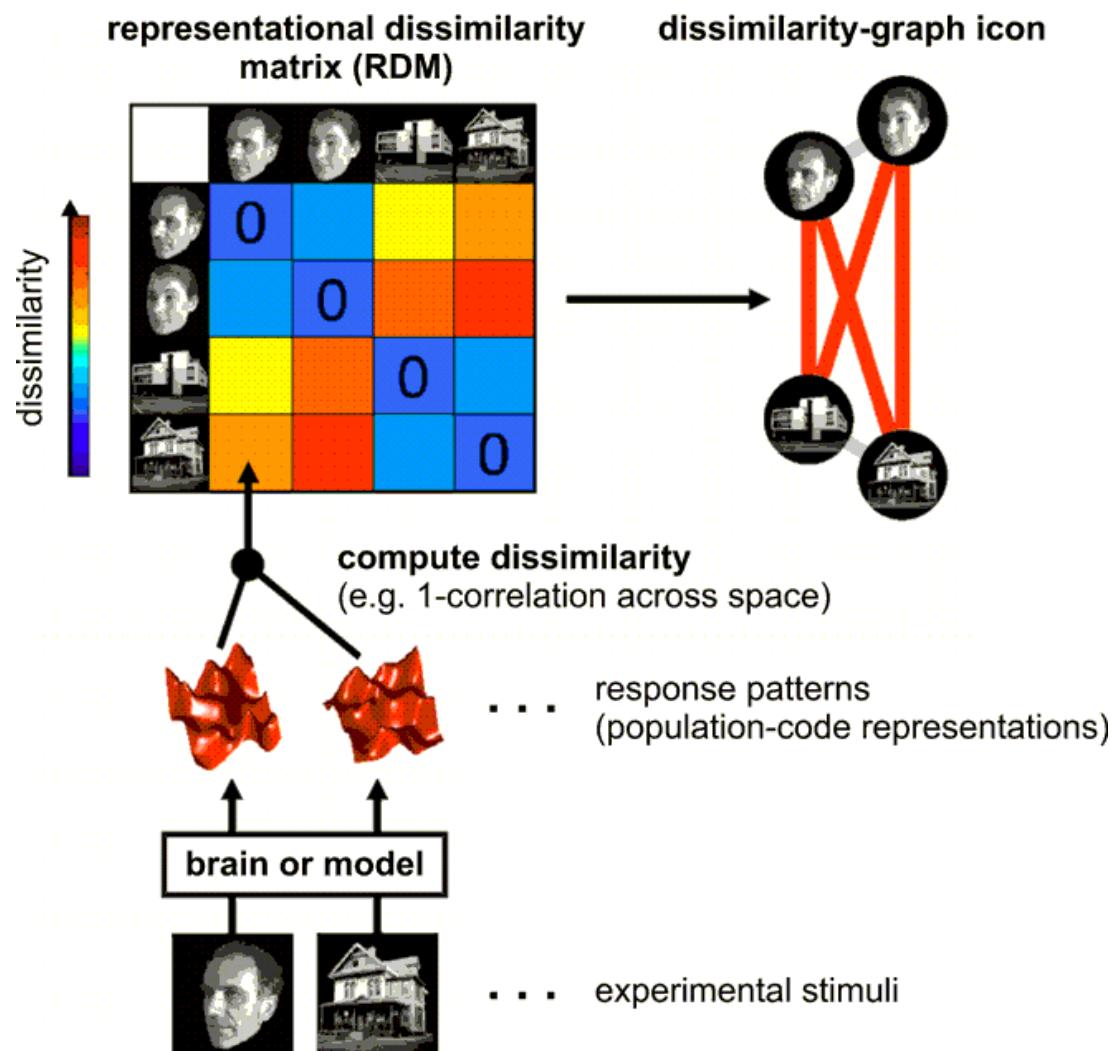
Nathaniel Blanchard

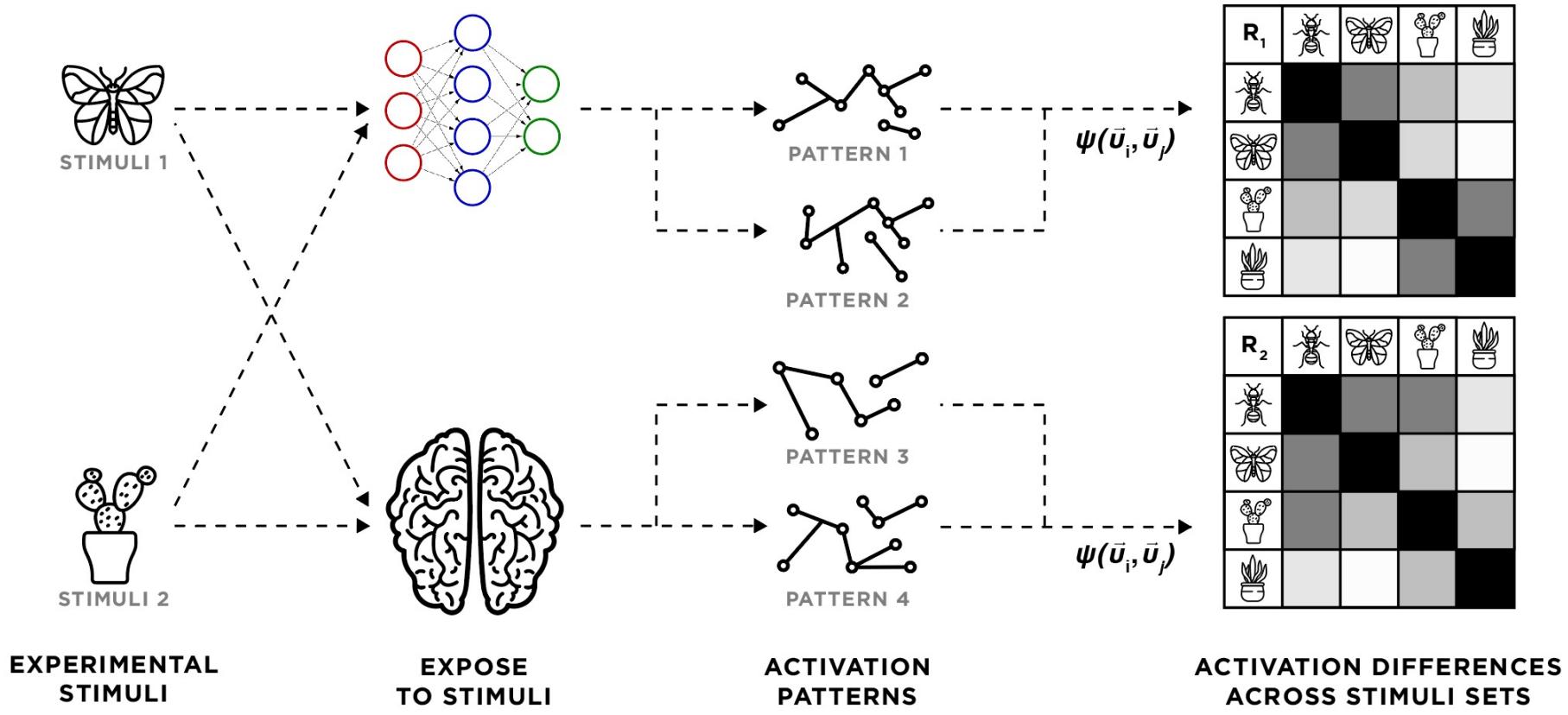
A Neurobiological Evaluation Metric for Neural Network Model Search

IEEE/CVF CVPR, 2019



Kriegeskorte et al.: Representational Similarity Analysis





RDM Step 1: Data Representation

Given a single feature f and a single stimulus s , $v = f(s)$, where v is the value of feature f in response to s . Likewise, the vector

$$\vec{v} = \begin{bmatrix} v_1 \\ v_2 \\ \vdots \\ v_n \end{bmatrix}^T = \begin{bmatrix} f_1(s) \\ f_2(s) \\ \vdots \\ f_n(s) \end{bmatrix}^T$$

can represent the feature values of a collection of n features, f_1, f_2, \dots, f_n , in response to s .

RDM Step 1: Data Representation

If one expands the representation of s to a set of m stimuli $S = s_1, s_2, \dots, s_m$, the natural extension of \vec{v} is the set of feature value collections $V = \vec{v}_1, \vec{v}_2, \dots, \vec{v}_m$, in which $s_i \in S$ is paired with $\vec{v}_i \in V$ for each $i = 1, 2, \dots, m$.

RDM Step 2: Dissimilarity

Define the dissimilarity score between any two
 $\vec{v}_i \in V$ and $\vec{v}_j \in V$:

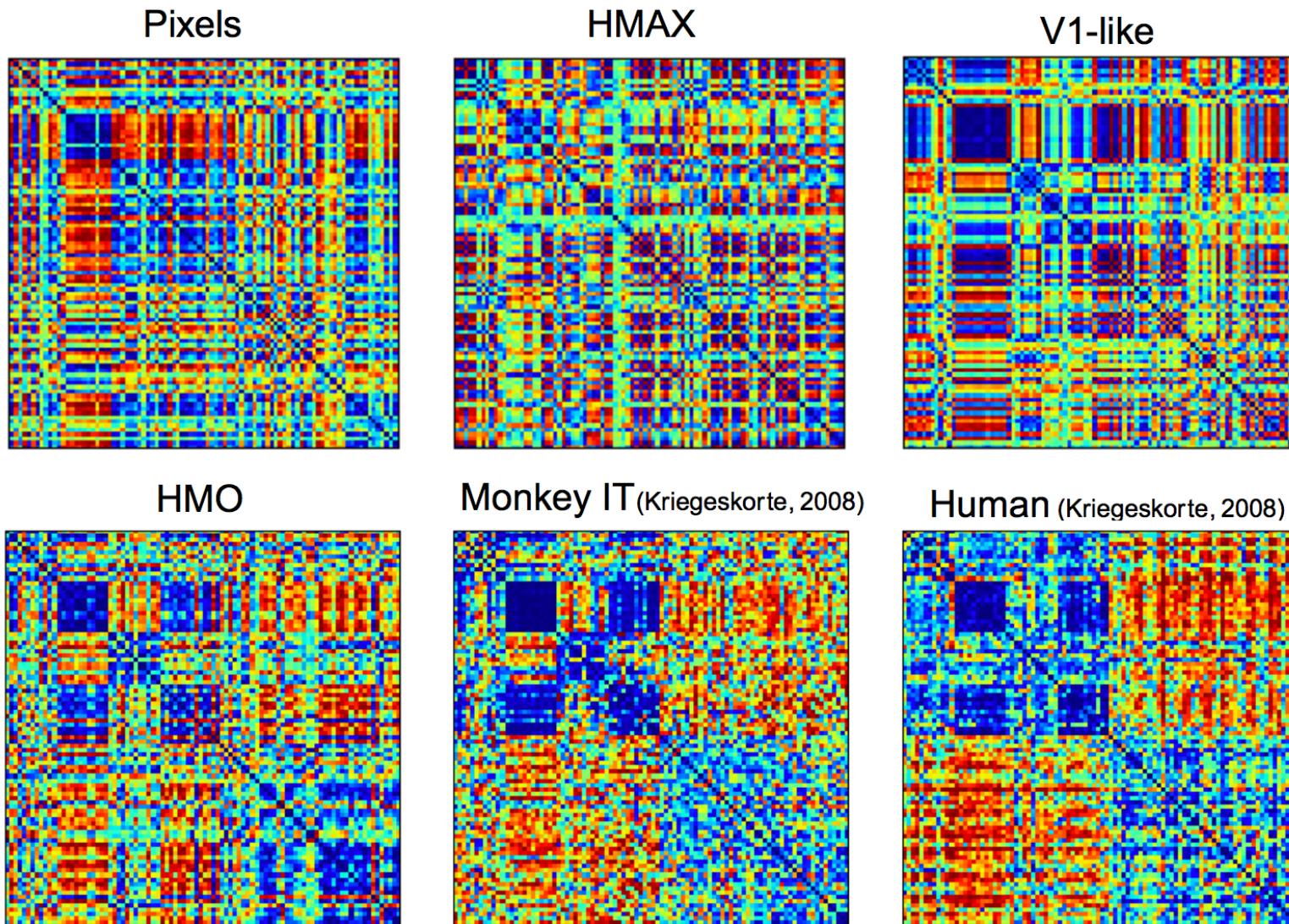
$$\psi(\vec{v}_i, \vec{v}_j) := 1 - \frac{(\vec{v}_i - \bar{v}_i) \cdot (\vec{v}_j - \bar{v}_j)}{\|\vec{v}_i - \bar{v}_i\|_2 \|\vec{v}_j - \bar{v}_j\|_2}$$

RDM Step 3: Construct Matrix

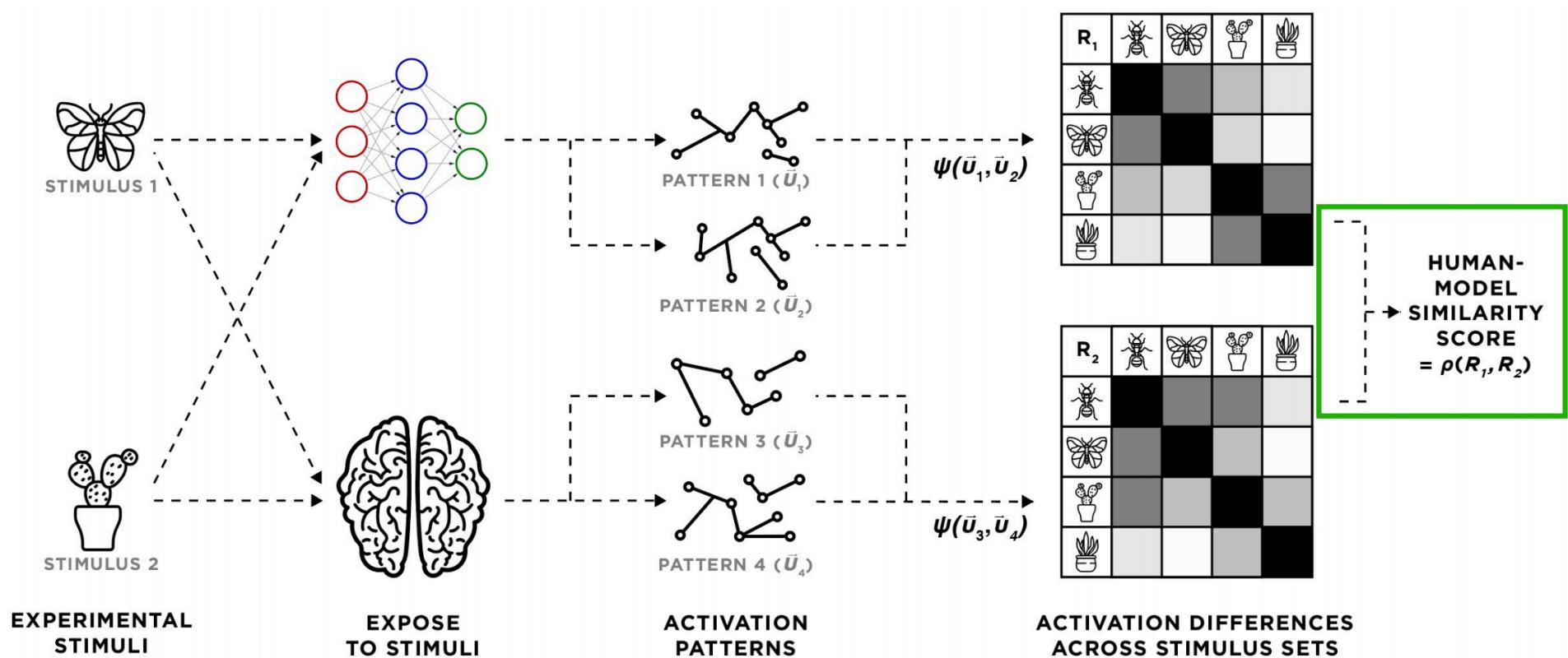
An RDM R may then be constructed from S , V , and ψ as:

$$R = \begin{bmatrix} \psi(\vec{v}_1, \vec{v}_2) & \psi(\vec{v}_1, \vec{v}_3) & \dots & \psi(\vec{v}_1, \vec{v}_m) \\ & \psi(\vec{v}_2, \vec{v}_3) & \dots & \psi(\vec{v}_2, \vec{v}_m) \\ & & \ddots & \vdots \\ & & & \psi(\vec{v}_{m-1}, \vec{v}_m) \end{bmatrix}$$

Works well for assessing biological fidelity:



Computing Human-Model Similarity



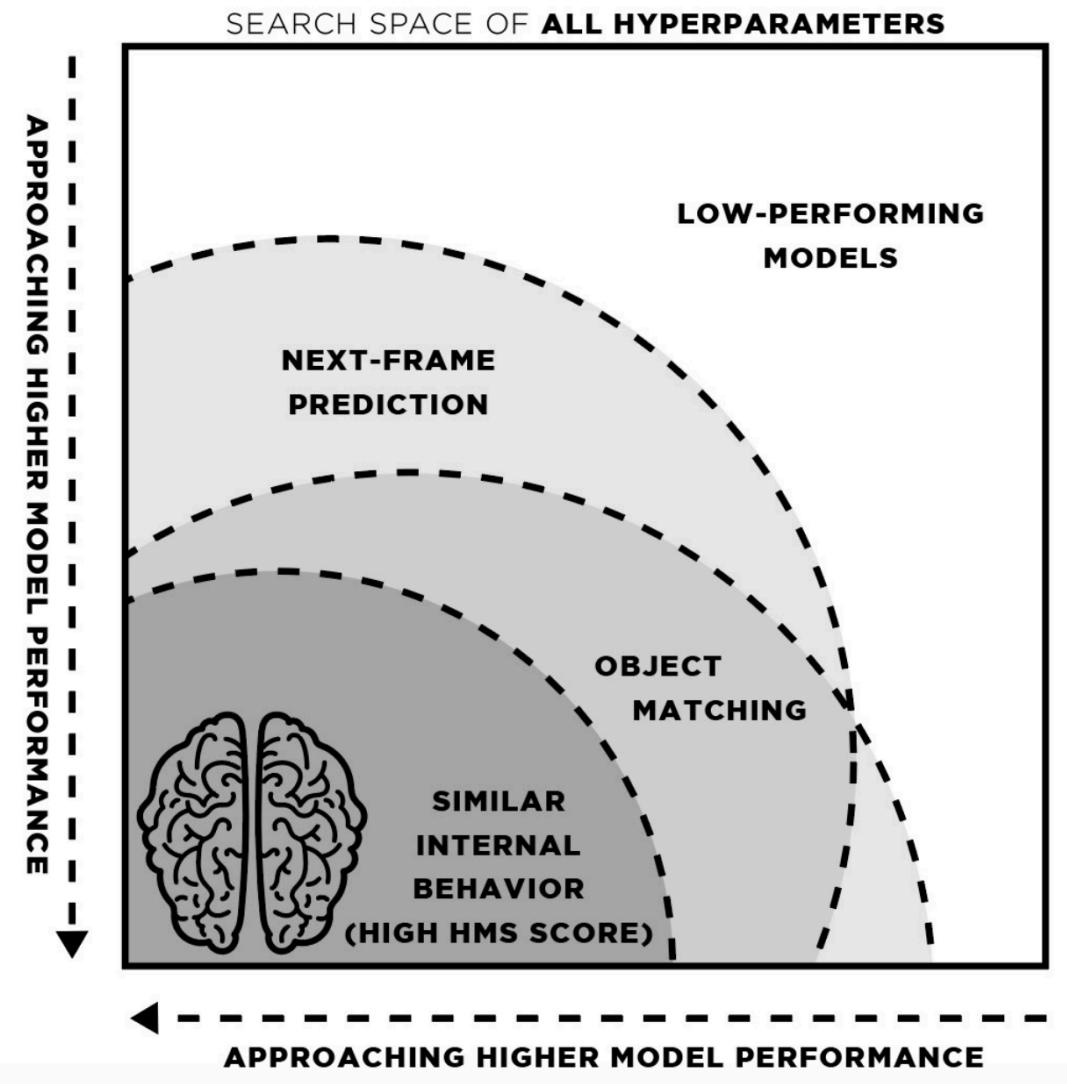
Human Model Similarity Score

$$HMS = \rho(\hat{R}_1, \hat{R}_2)$$

RDM 1 RDM 2

The diagram illustrates the calculation of the Human Model Similarity Score (HMS). At the bottom, the text "Spearman's Rank Correlation" is positioned. Above it, the formula $HMS = \rho(\hat{R}_1, \hat{R}_2)$ is centered. Two arrows point from the text "RDM 1" and "RDM 2" above the formula down to the \hat{R}_1 and \hat{R}_2 terms in the formula respectively.

AI as a search problem



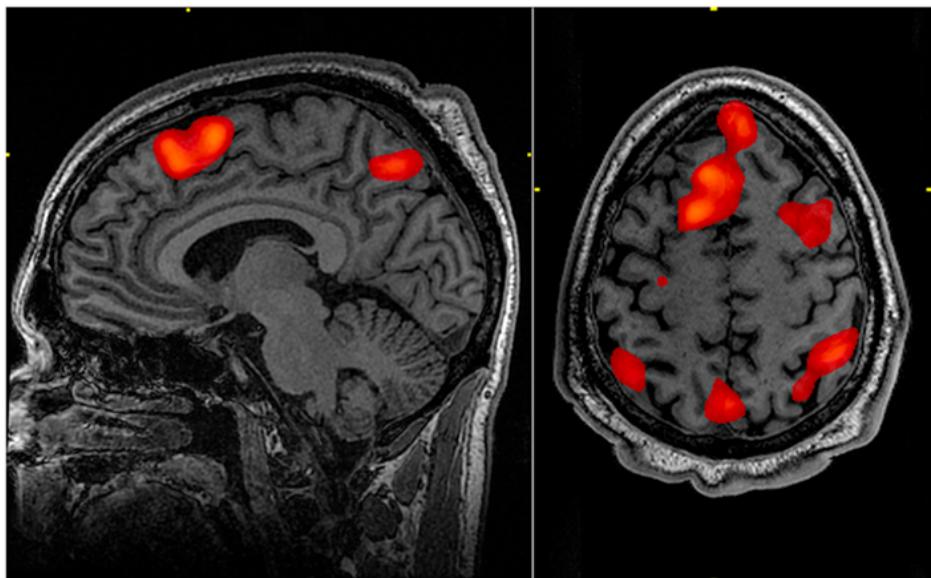
Experiments

fMRI

A direct way to measure human brain activity

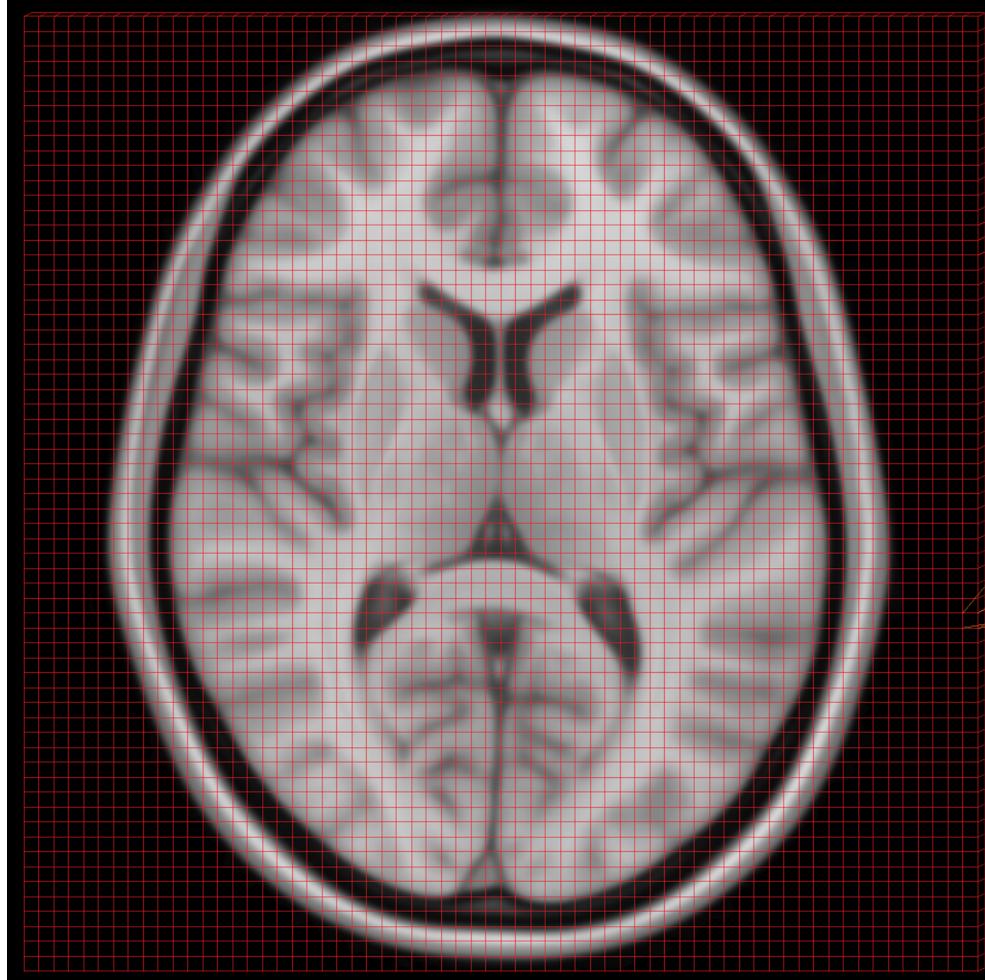
Non-invasive experimentation with humans

Uses blood flow as a proxy for neuronal activations



**Spatial resolution good enough
to identify Brodmann areas**

fMRI Data



Matrix size of slice:
 64×64

Voxel size:
3mm x 3mm x 3mm

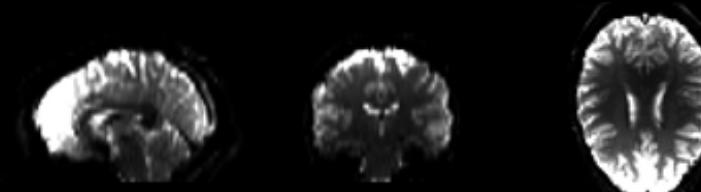
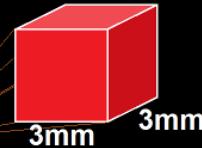


Image adapted from: <https://miykael.github.io/nipype-beginner-s-guide/neuroimaging.html>

fMRI Experimental Setup

Data collected by the Kriegeskorte lab at the University of Cambridge*

Eight RDMs were constructed from fMRI recordings of four subjects over two sessions in response to 92 random stimuli

Recordings were from measurements of $1.95 \times 1.95 \times 2\text{mm}^3$ within an occipitotemporal measurement slab (5cm thick).

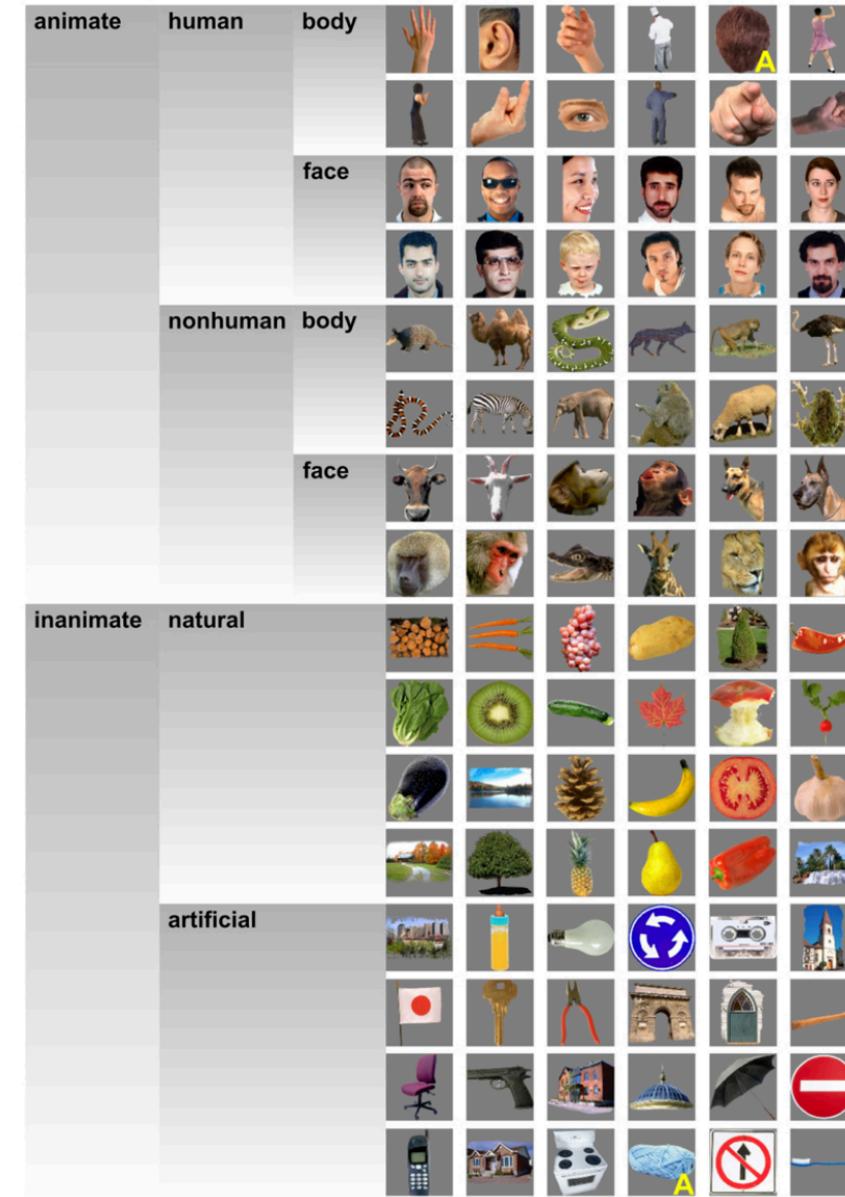
Each stimulus was displayed for 300 milliseconds, every 3700 milliseconds, with four seconds between stimuli.

Subject RDMs were averaged together into a mean human brain RDM, which reduced noise.

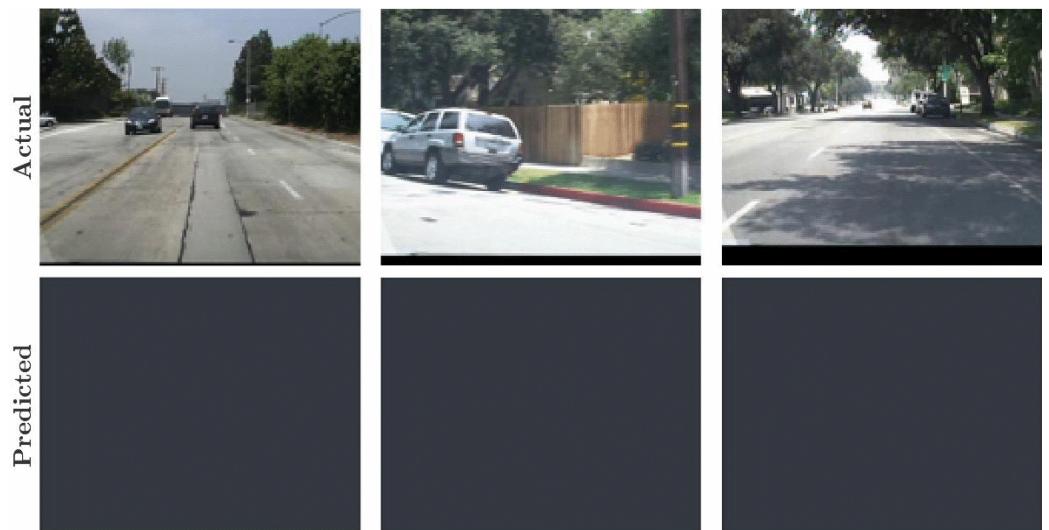
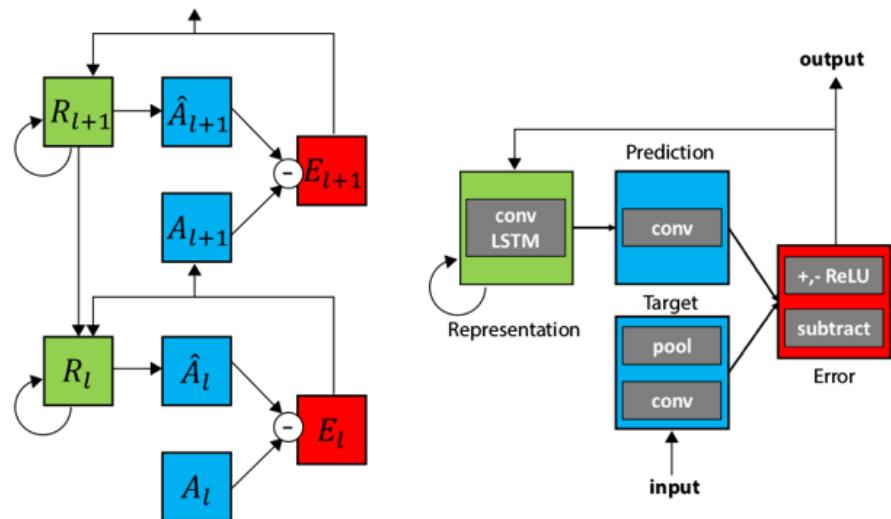


01 Siemens MAGNETOM Trio BY-SA 2.0 Image Editor

fMRI Stimuli Set



Architecture: PredNet



Predictive Coding Network Performance: 95 Nets



Tenenbaum et al. Science 2011

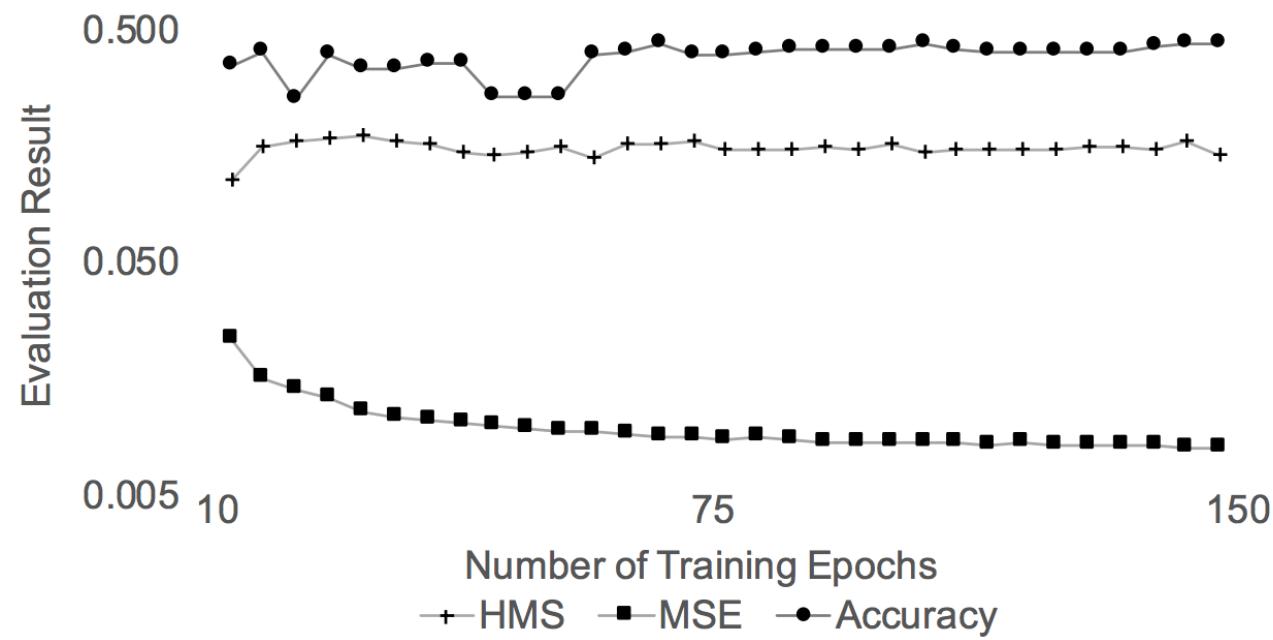
Evaluation Task	Metric	Mean (SD)	Top Ten HMS Mean (SD)
Next Frame Prediction Error	Pixel MSE	0.092 (0.148)	0.009 (0.003)
Object Matching	Accuracy	0.367 (0.134)	0.459 (0.049)
Human-Model Similarity	RDM Correlation	0.106 (0.055)	0.178 (0.011)

HMS is predictive of network performance on other metrics

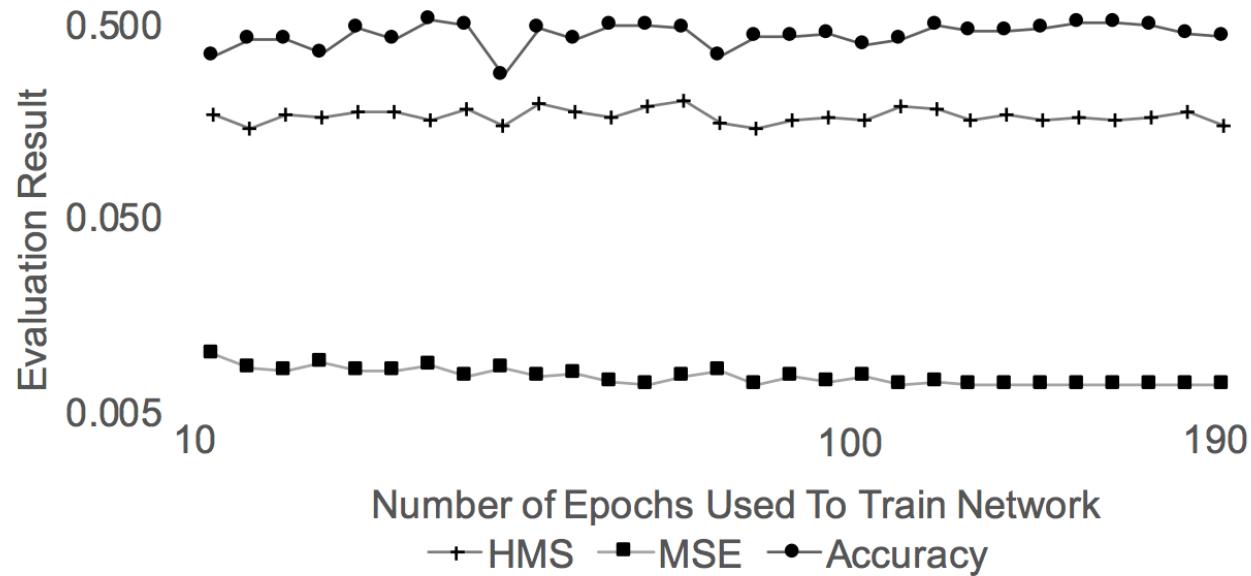
Variable	Accuracy	HMS	Learning Rate
Next Frame Prediction Error	-0.791**	-0.646**	0.635**
Object Matching Accuracy	.	0.575**	-0.517**
Human-Model Similarity	.	.	-0.452**

** $p < 0.001$

Within-Network Stability

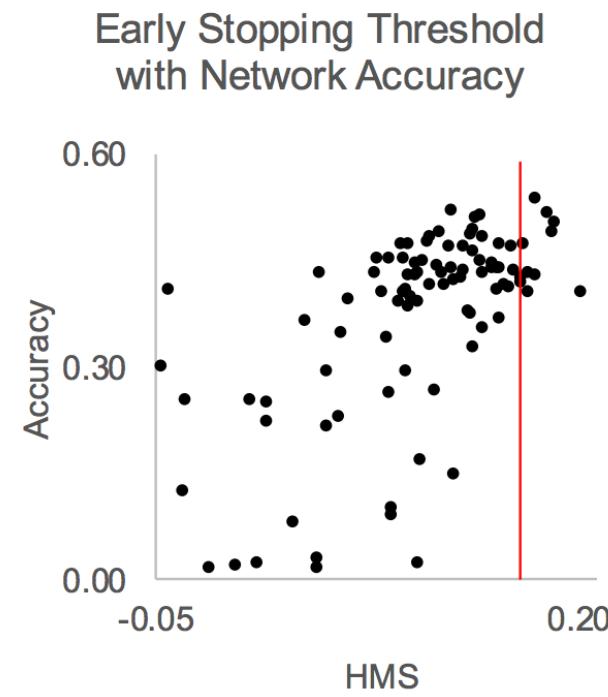
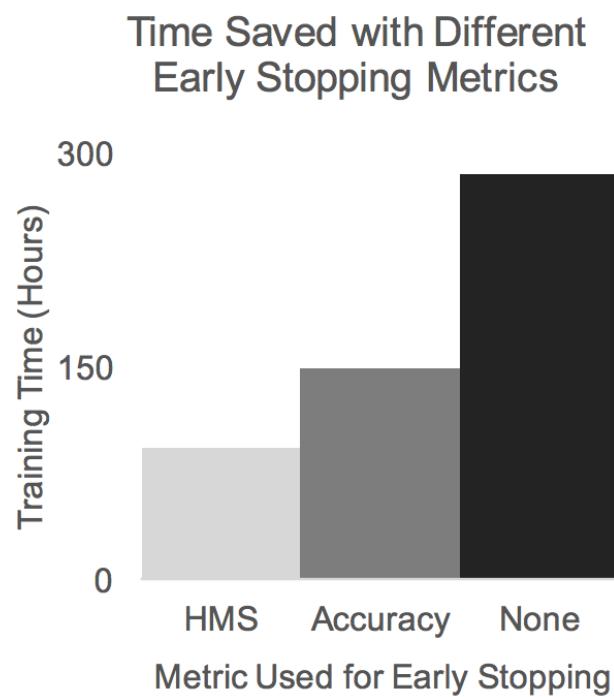


Across-Network Stability



66 Models, Mean

HMS-Driven Early Stopping

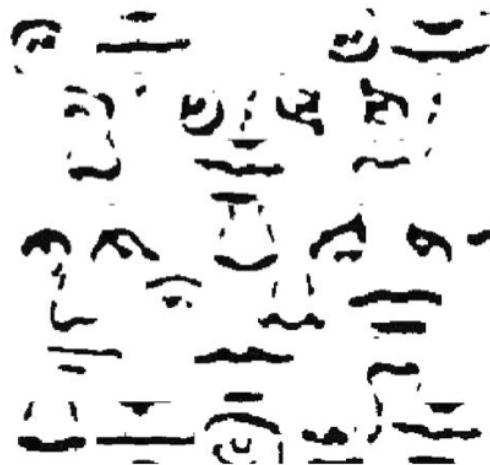


Is fMRI the best reference for this?

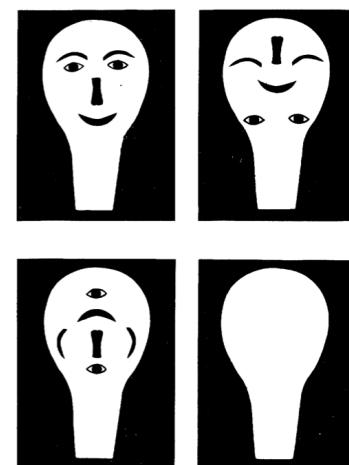
Easier: Human Behavior

Visual Psychophysics: probe psychological and perceptual thresholds through controlled manipulation of stimuli.

Careful management of stimulus construction, ordering and presentation allows for precise determination of perceptual thresholds.



Garrido et al 2011



Goren et al 1975

Thank you!