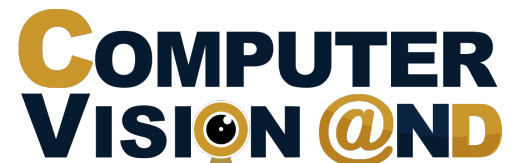


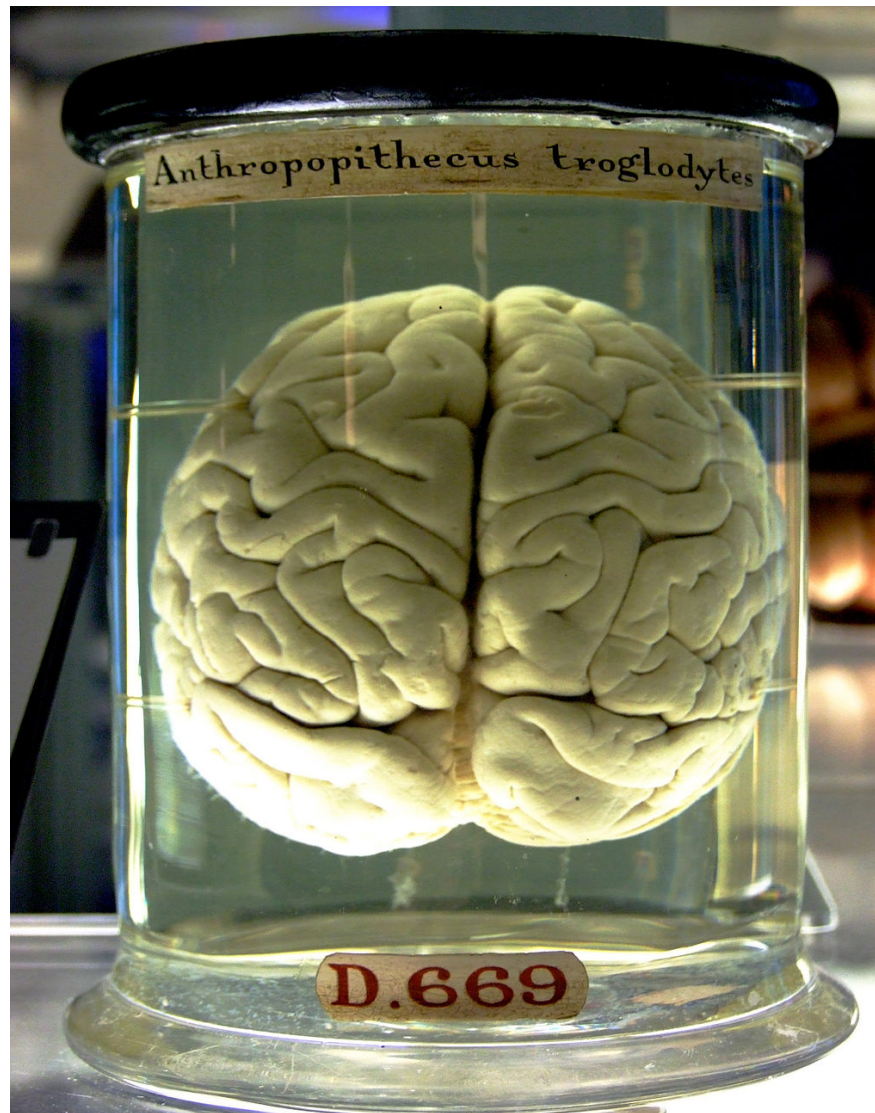
# Representational Dissimilarity Analysis as a Tool for Neural Network Model Search

Walter J. Scheirer

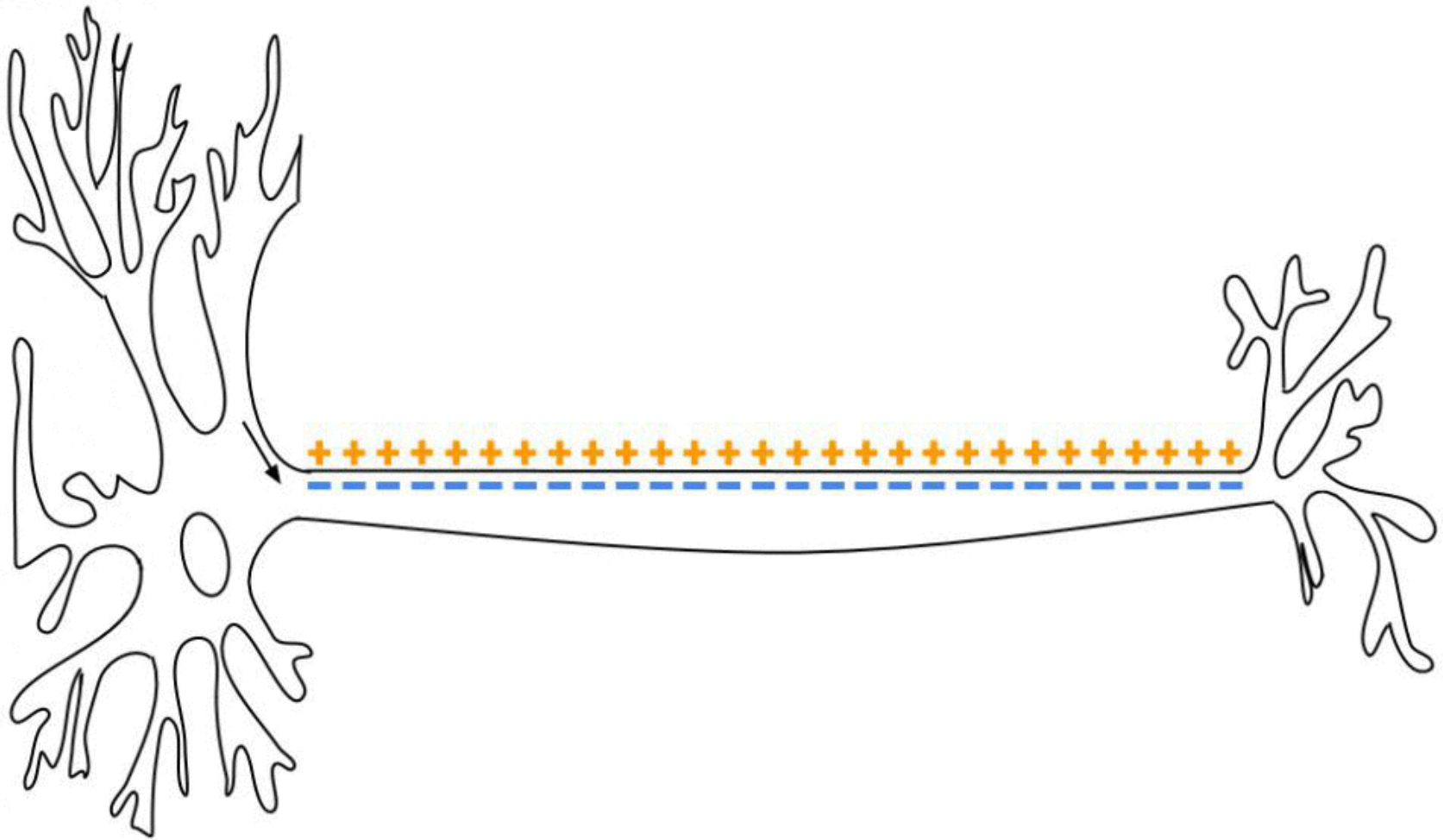
Computer Vision Research Laboratory  
Department of Computer Science and Engineering

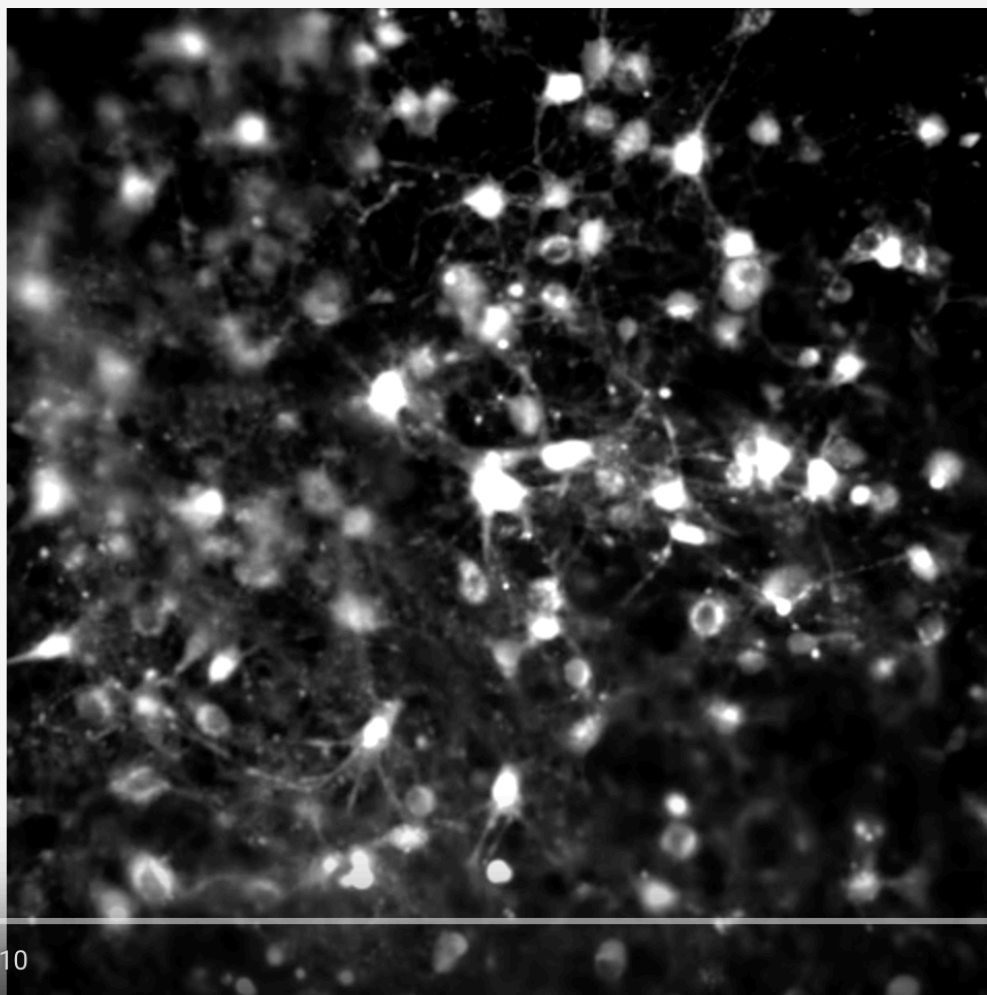


# A model of vision that works:



Chimp Brain in a jar © BY 2.0 Gaetan Lee





0:01 / 0:10

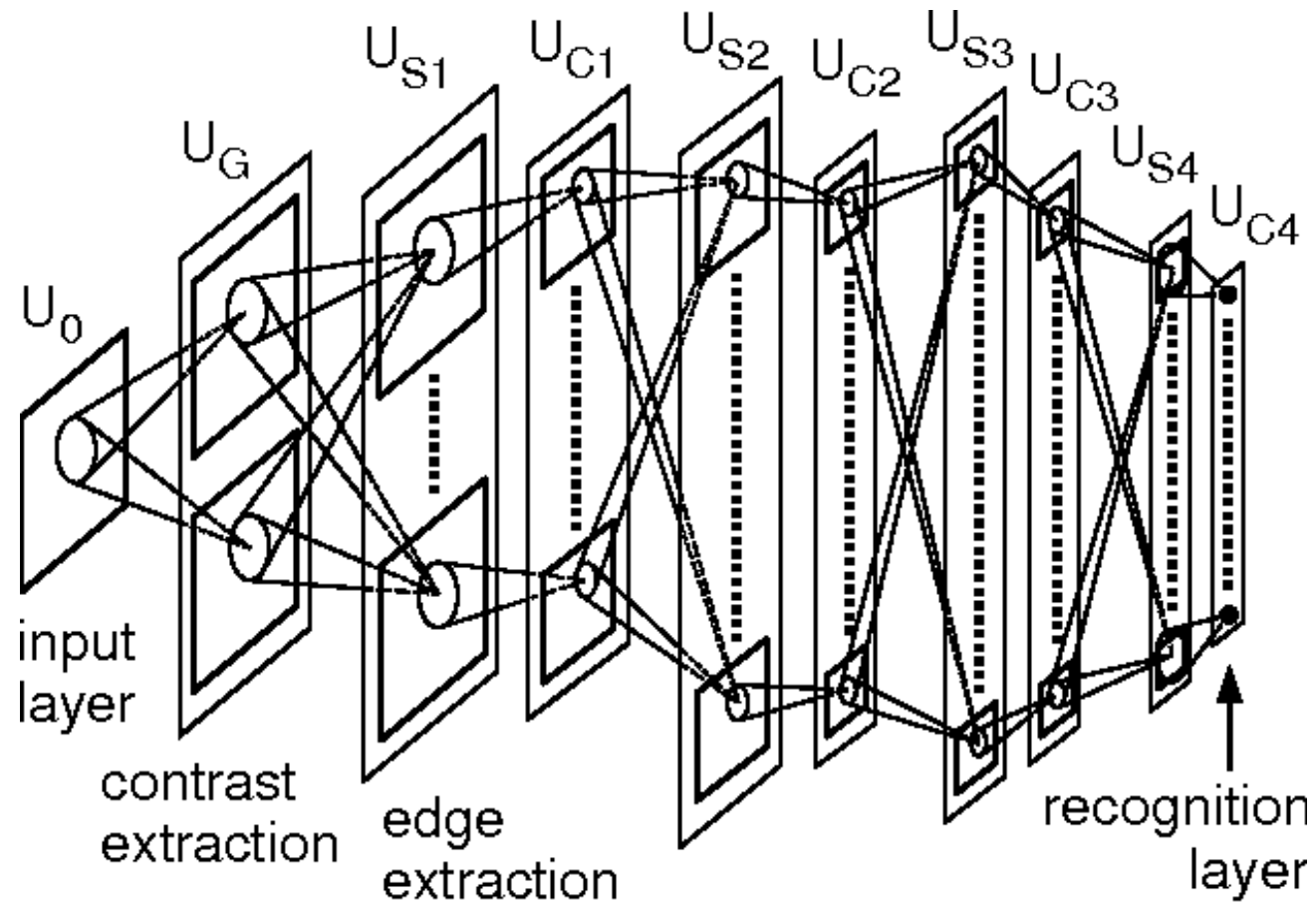


<https://www.youtube.com/watch?v=yy994HpFudc>



**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

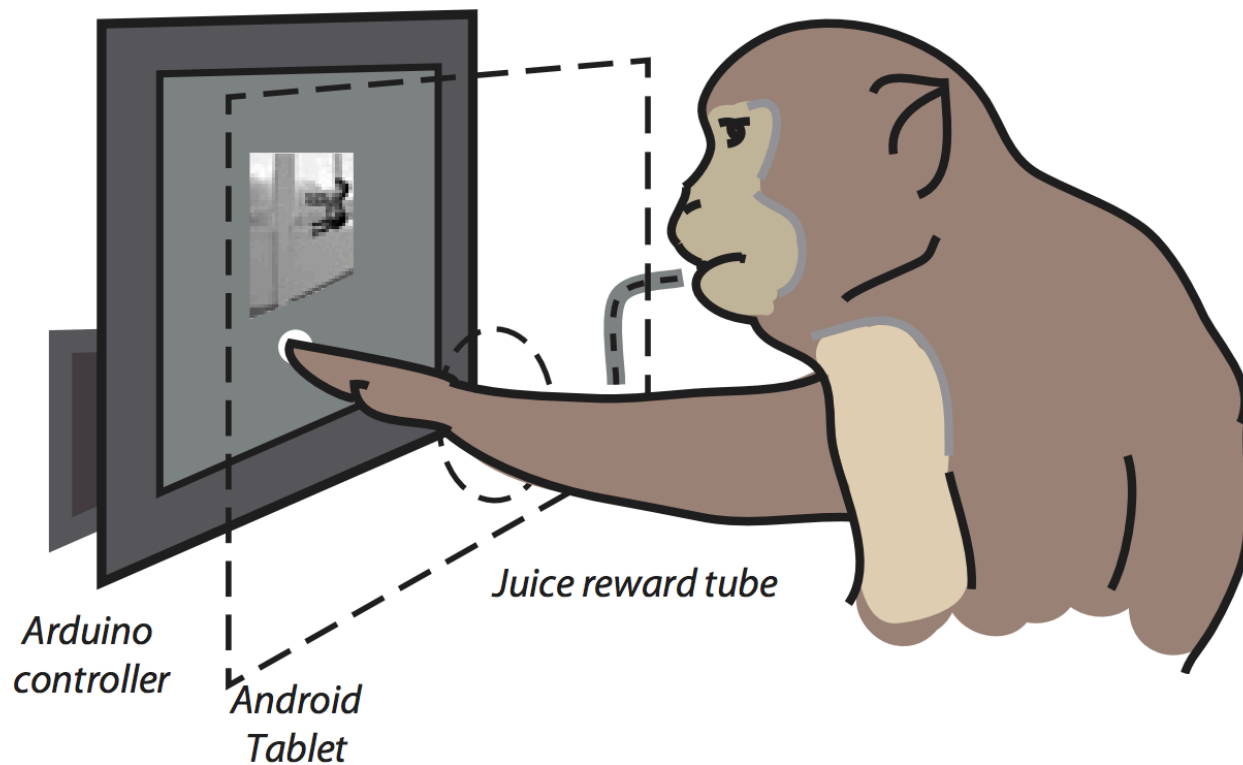
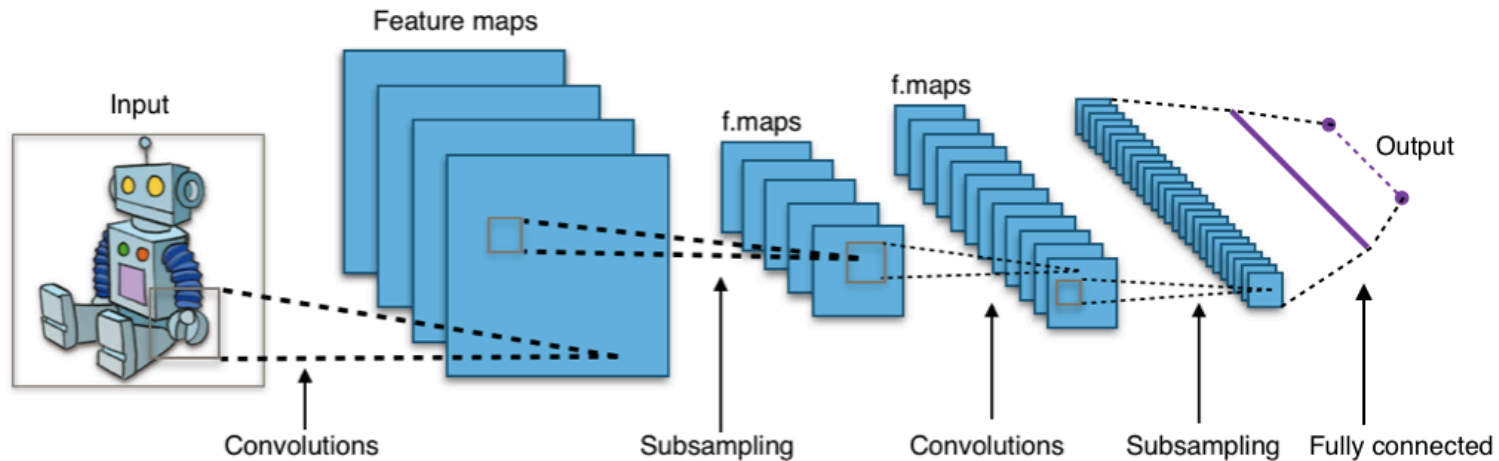


Image adapted from: Rajalingham et al. JNeurosci 2018

# CNN for Object Recognition

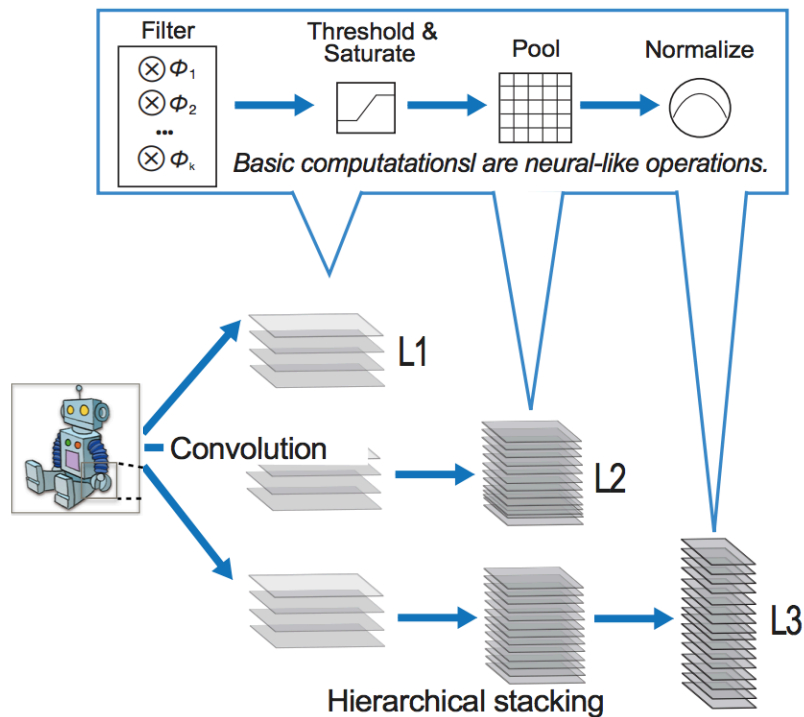


Typical CNN architecture © BY-SA 4.0 Aphex34

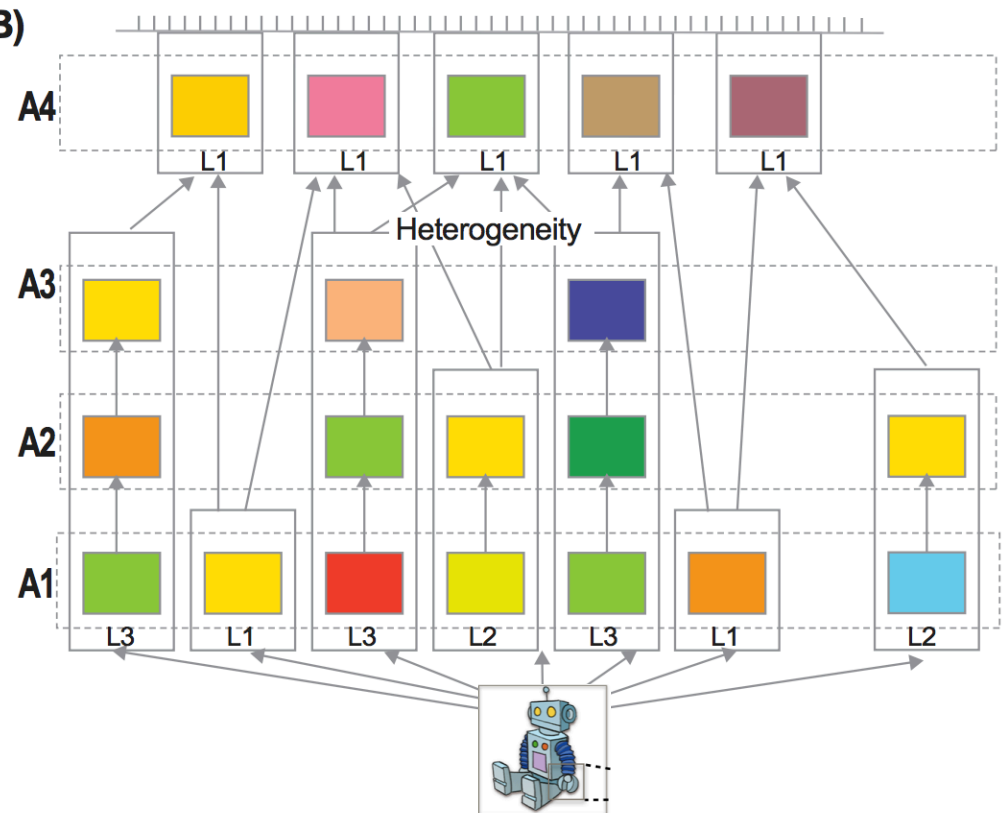


# Heterogeneous Hierarchical CNN

A) Basic operations



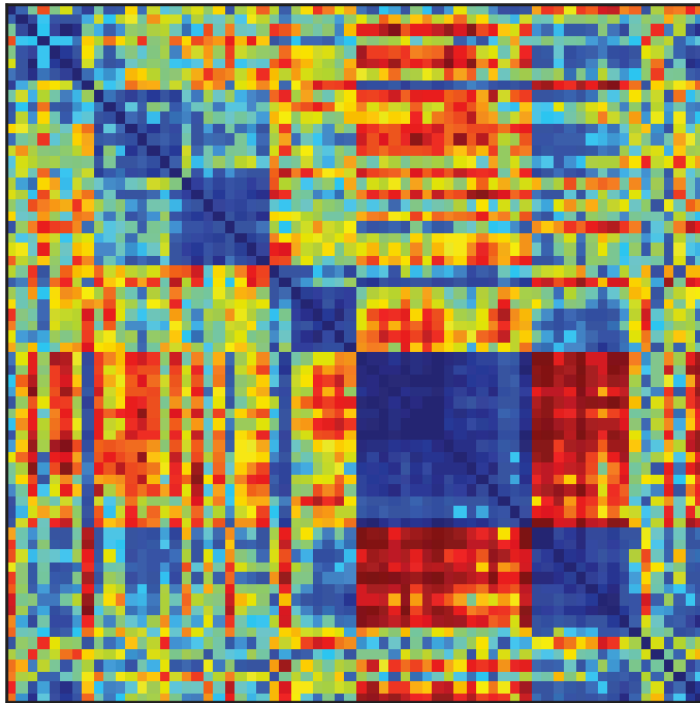
B)



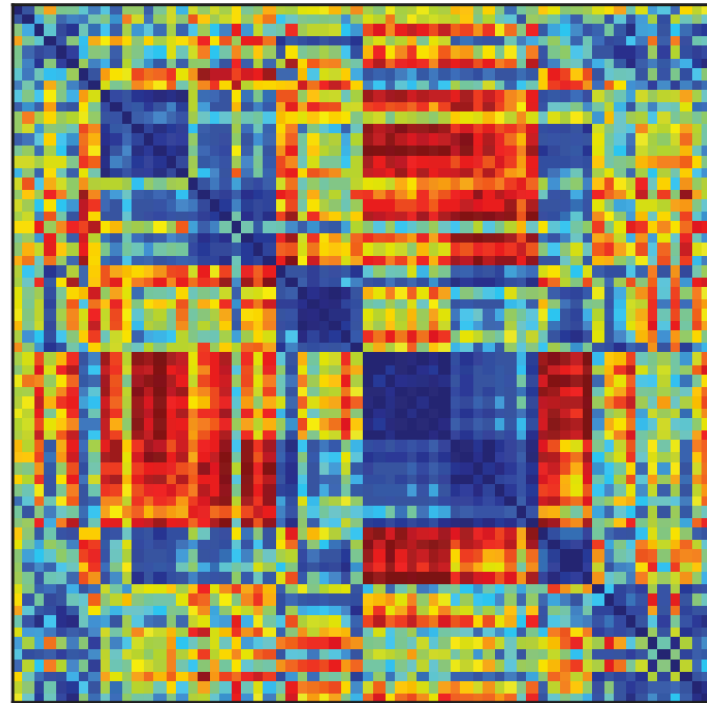
Yamins et al. NeurIPS 2013

# Population Responses: Model vs. Brain

HMO Model

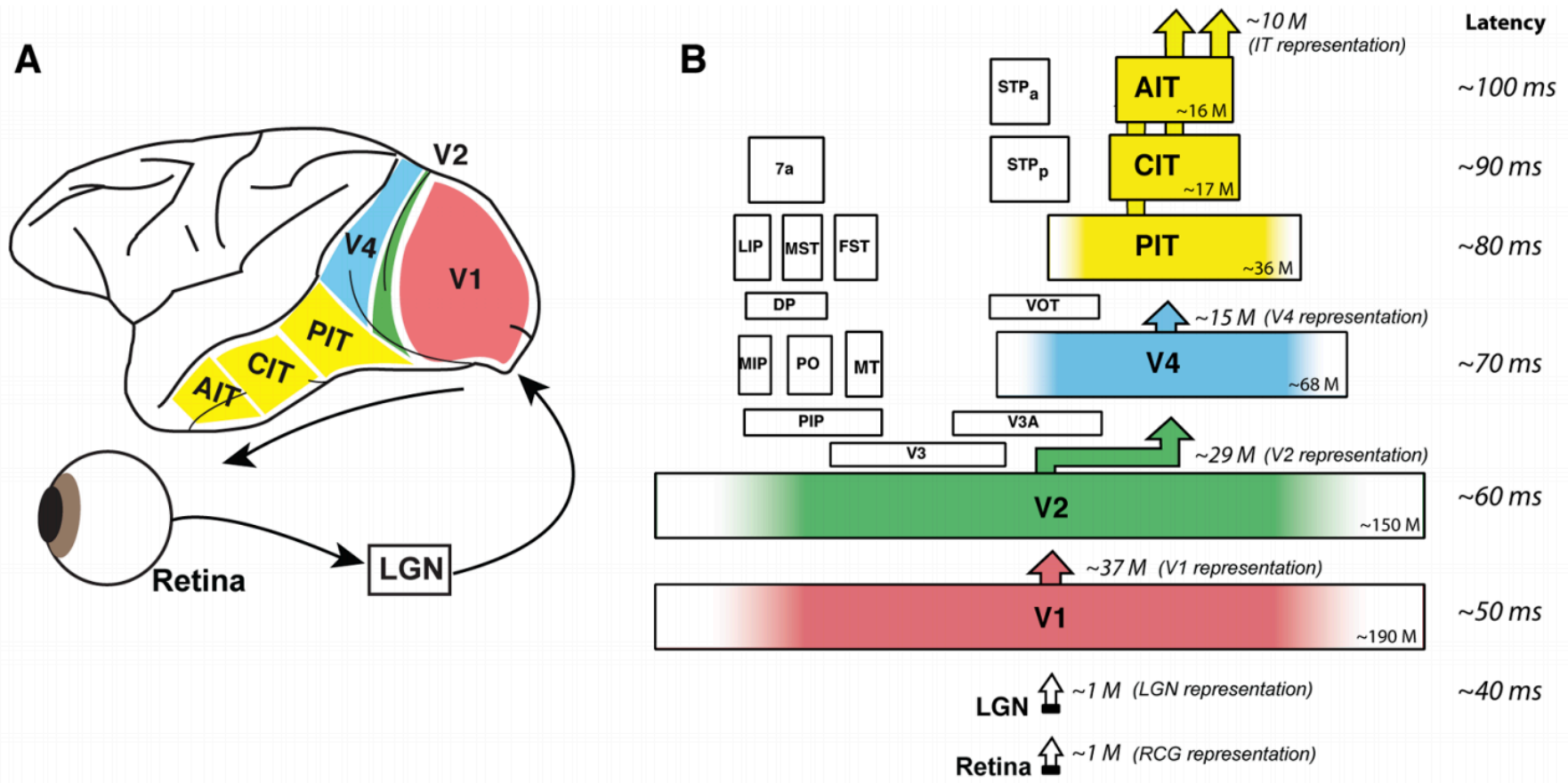


IT Neurons



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?

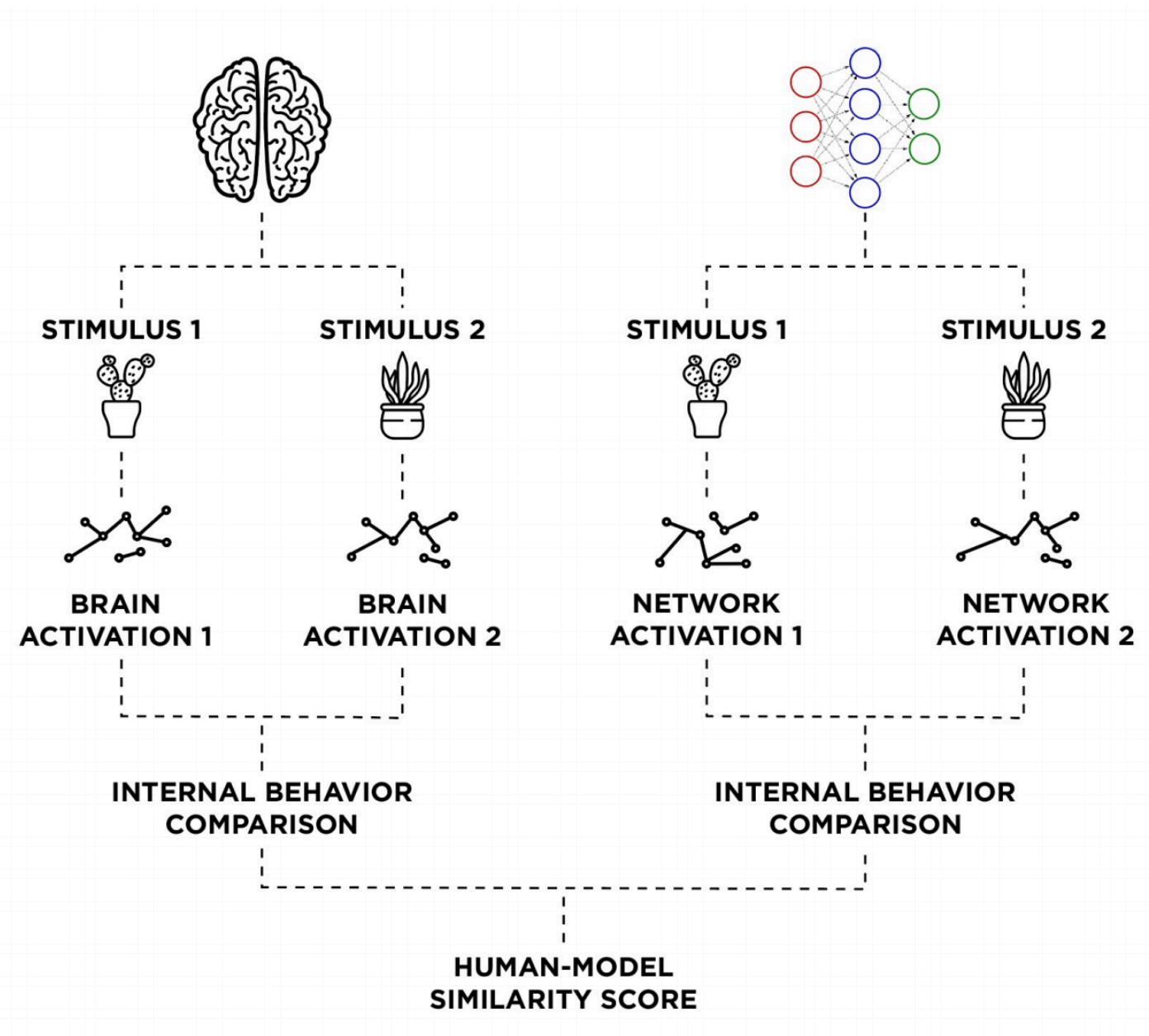


Nathaniel Blanchard

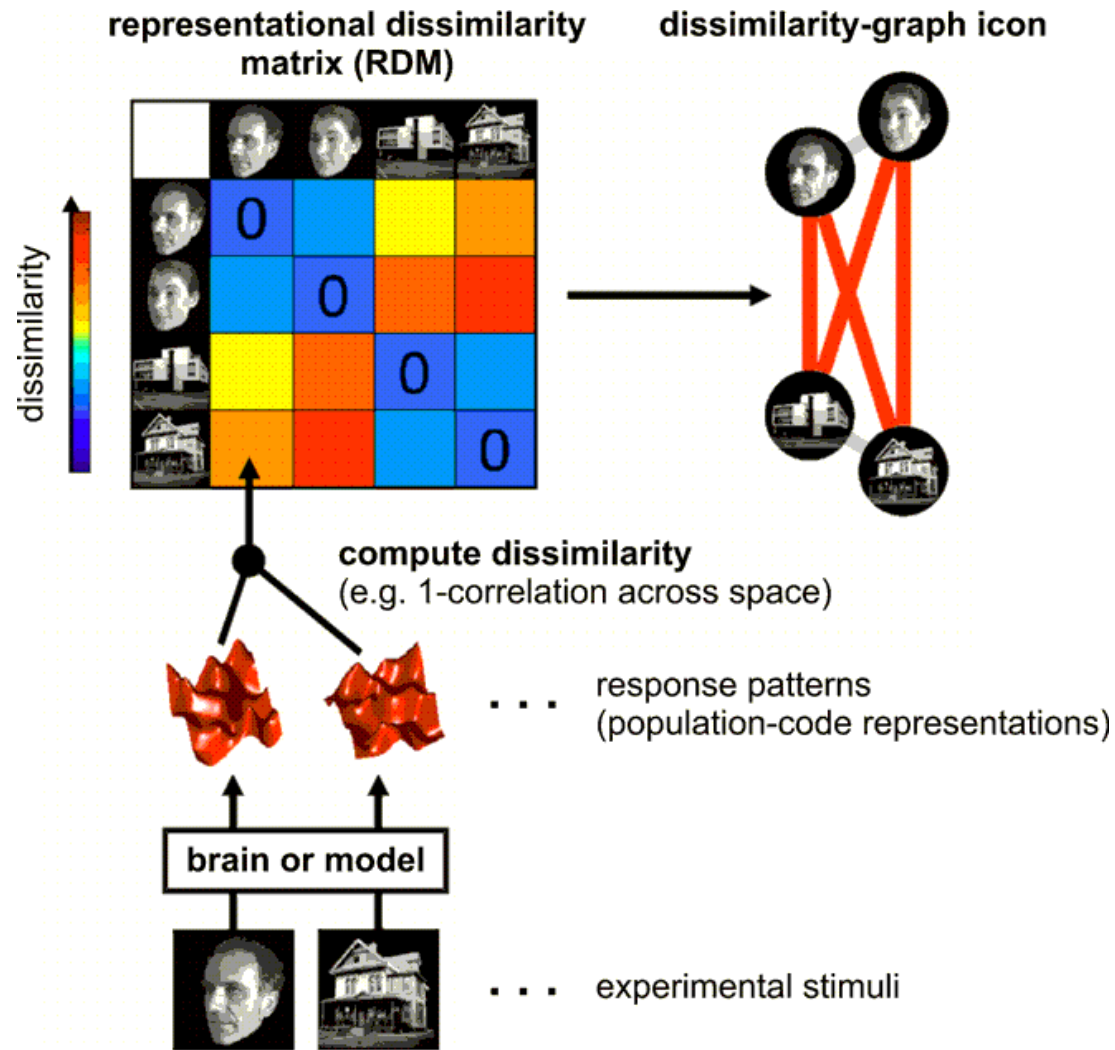
# A Neurobiological Evaluation Metric for Neural Network Model Search

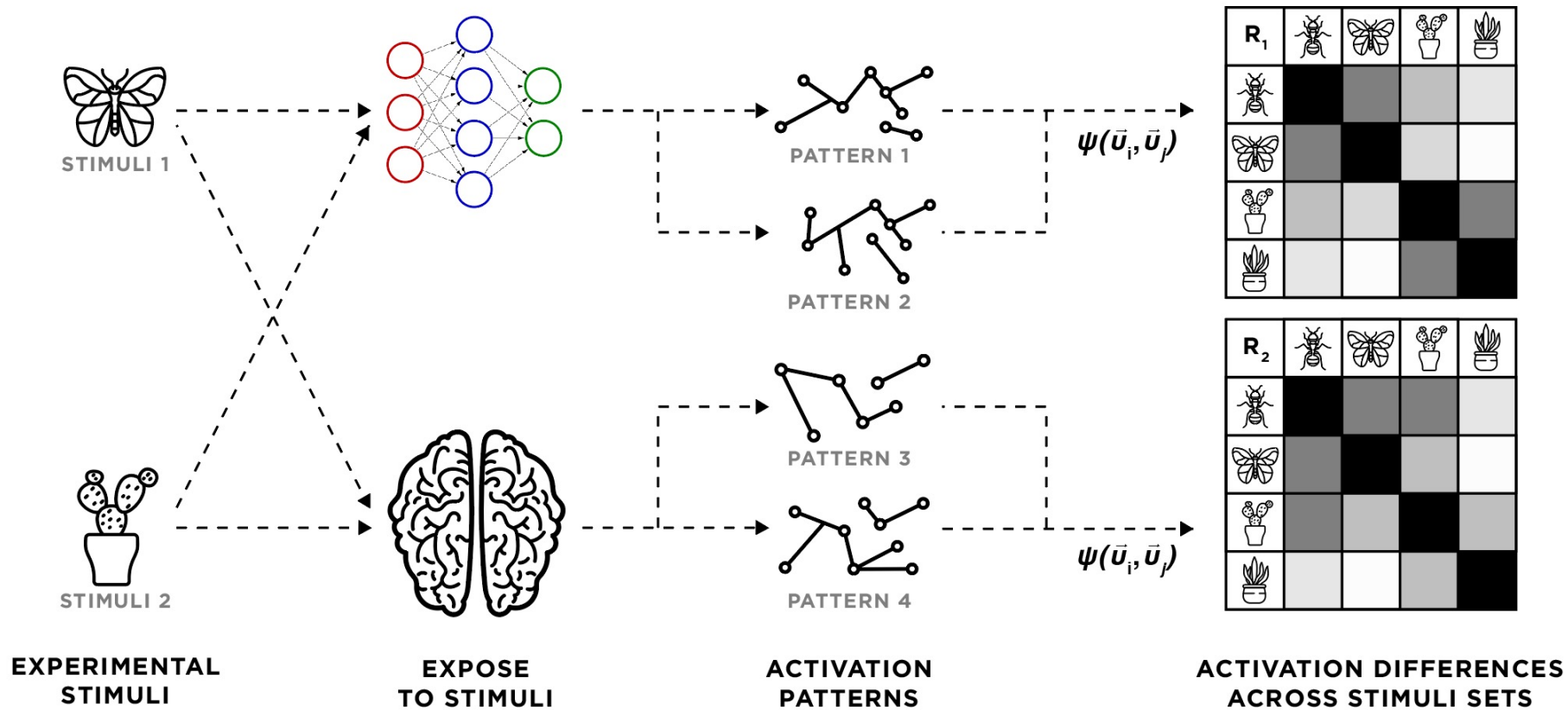
**IEEE/CVF CVPR, 2019**





# 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  $\vec{v}_j \in V$ :

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

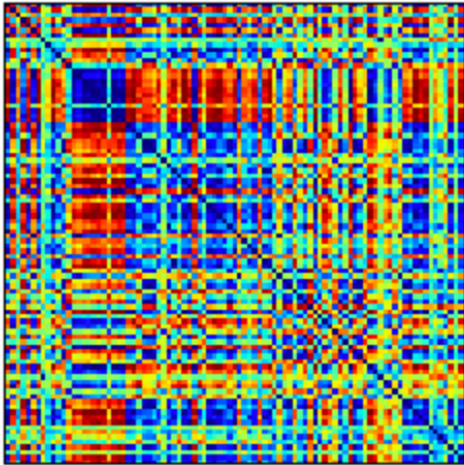
# RDM Step 3: Construct Matrix

An RDM  $R$  may then be constructed from  $S$ ,  $V$ , and  $\psi$  as:

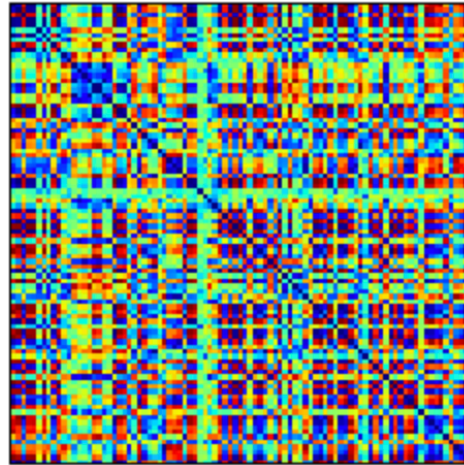
$$R = \begin{bmatrix} \psi(\vec{v}_1, \vec{v}_2) & \psi(\vec{v}_1, \vec{v}_3) & \dots & \psi(\vec{v}_1, \vec{v}_m) \\ & \psi(\vec{v}_2, \vec{v}_3) & \dots & \psi(\vec{v}_2, \vec{v}_m) \\ & & \ddots & \vdots \\ & & & \psi(\vec{v}_{m-1}, \vec{v}_m) \end{bmatrix}$$

# Works well for assessing biological fidelity:

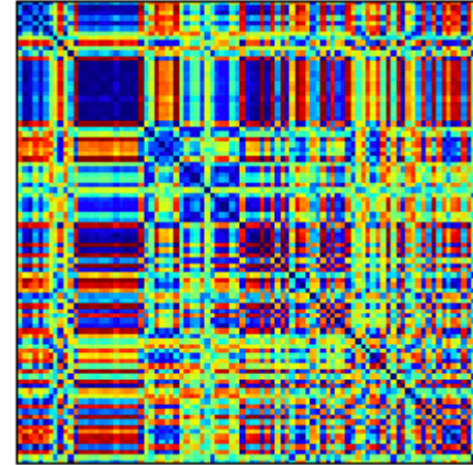
Pixels



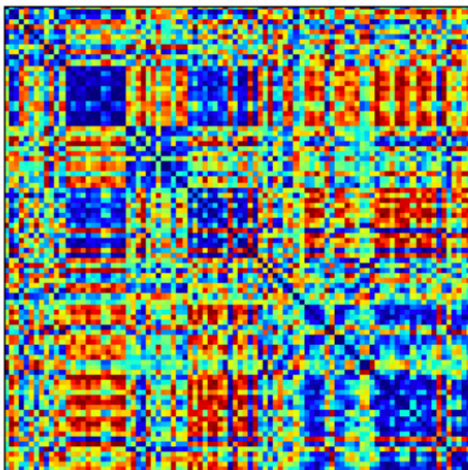
HMAX



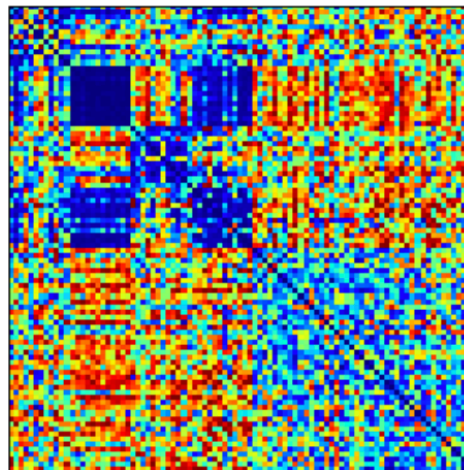
V1-like



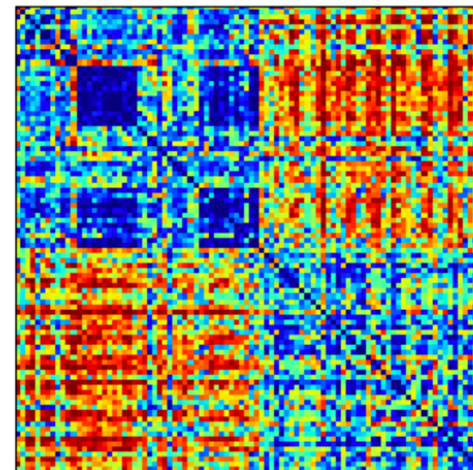
HMO



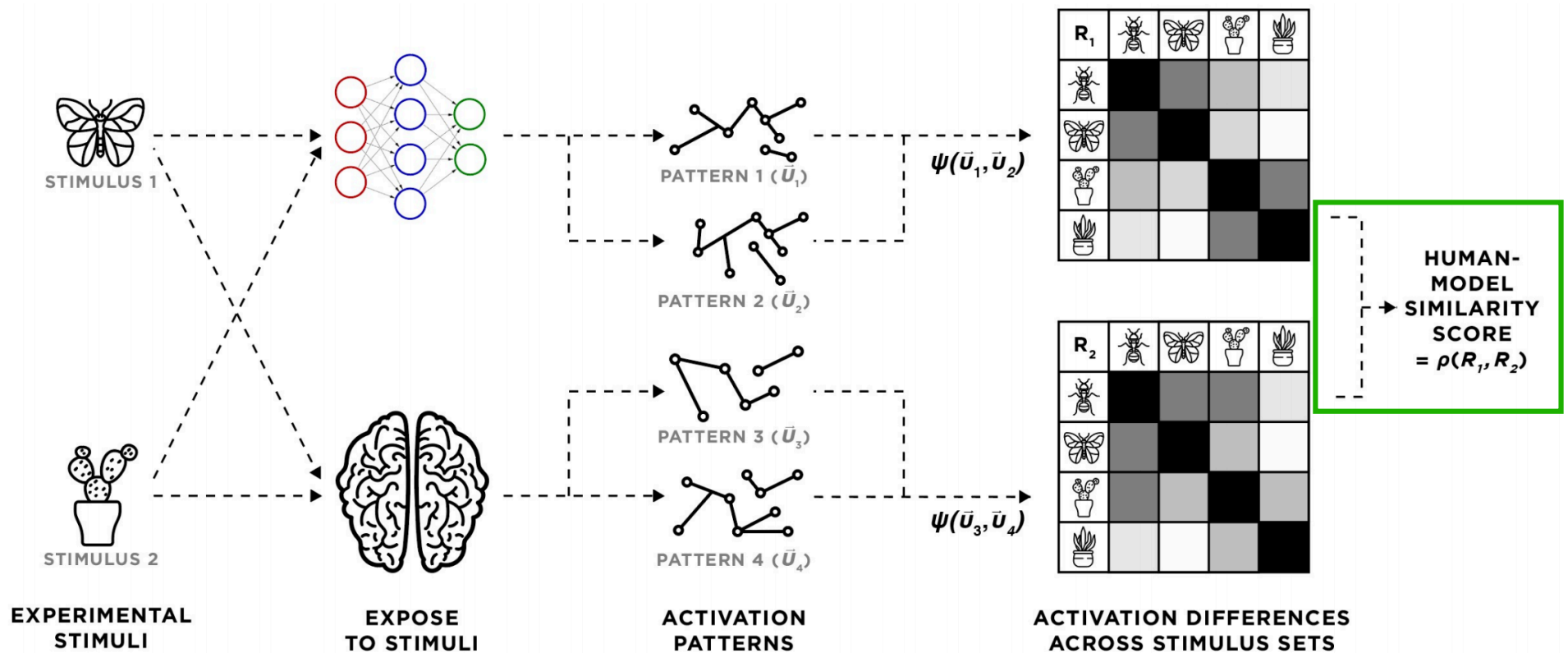
Monkey IT (Kriegeskorte, 2008)



Human (Kriegeskorte, 2008)



# Computing Human-Model Similarity



# Human Model Similarity Score

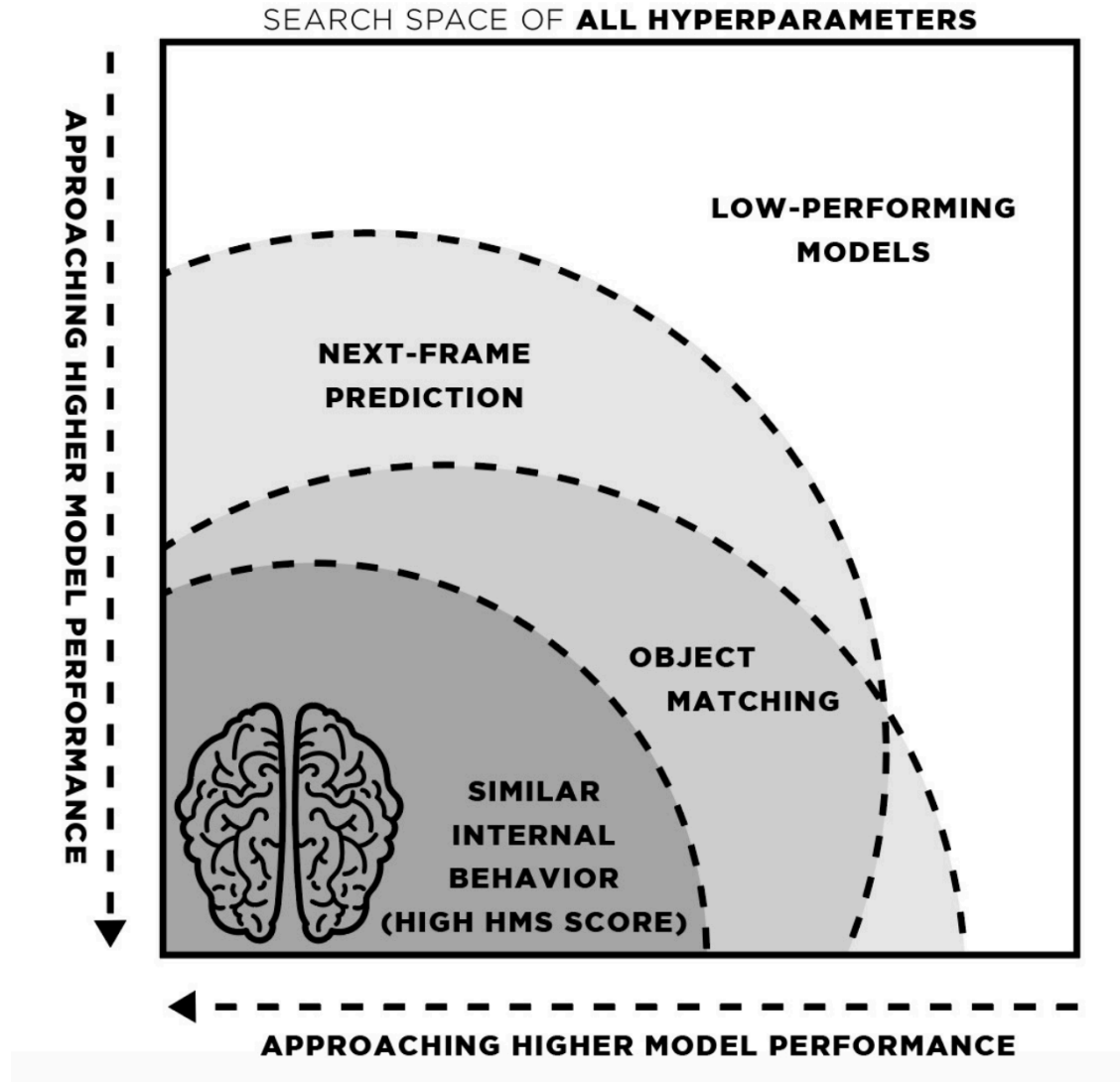
The diagram illustrates the calculation of the Human Model Similarity Score (HMS). At the top, 'RDM 1' and 'RDM 2' are labeled. Arrows point from each to the corresponding rank correlation terms in the equation below. The equation is  $HMS = \rho(\hat{R}_1, \hat{R}_2)$ . An arrow points from the text 'Spearman's Rank Correlation' to the  $\rho$  symbol in the equation.

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

Spearman's Rank Correlation



# AI as a search problem



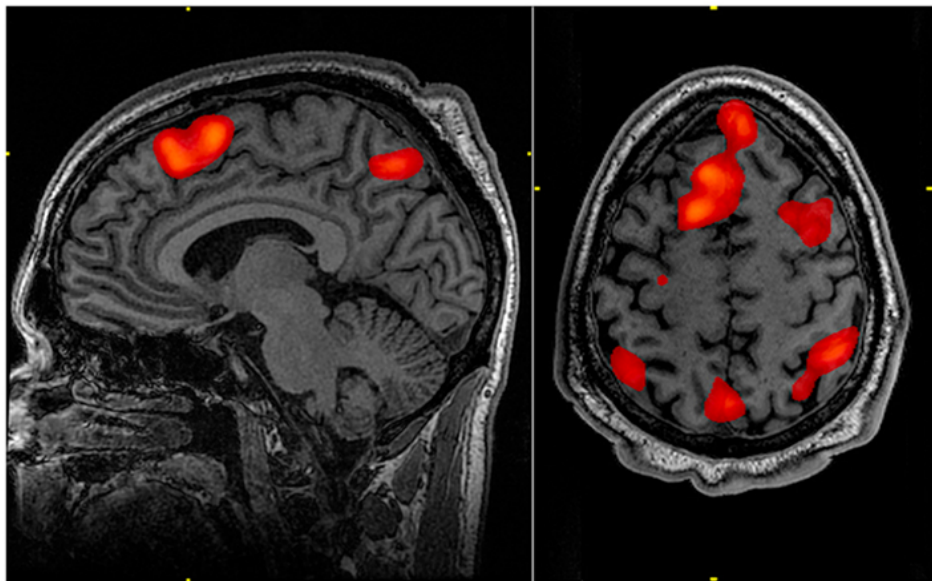
# Experiments

# 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

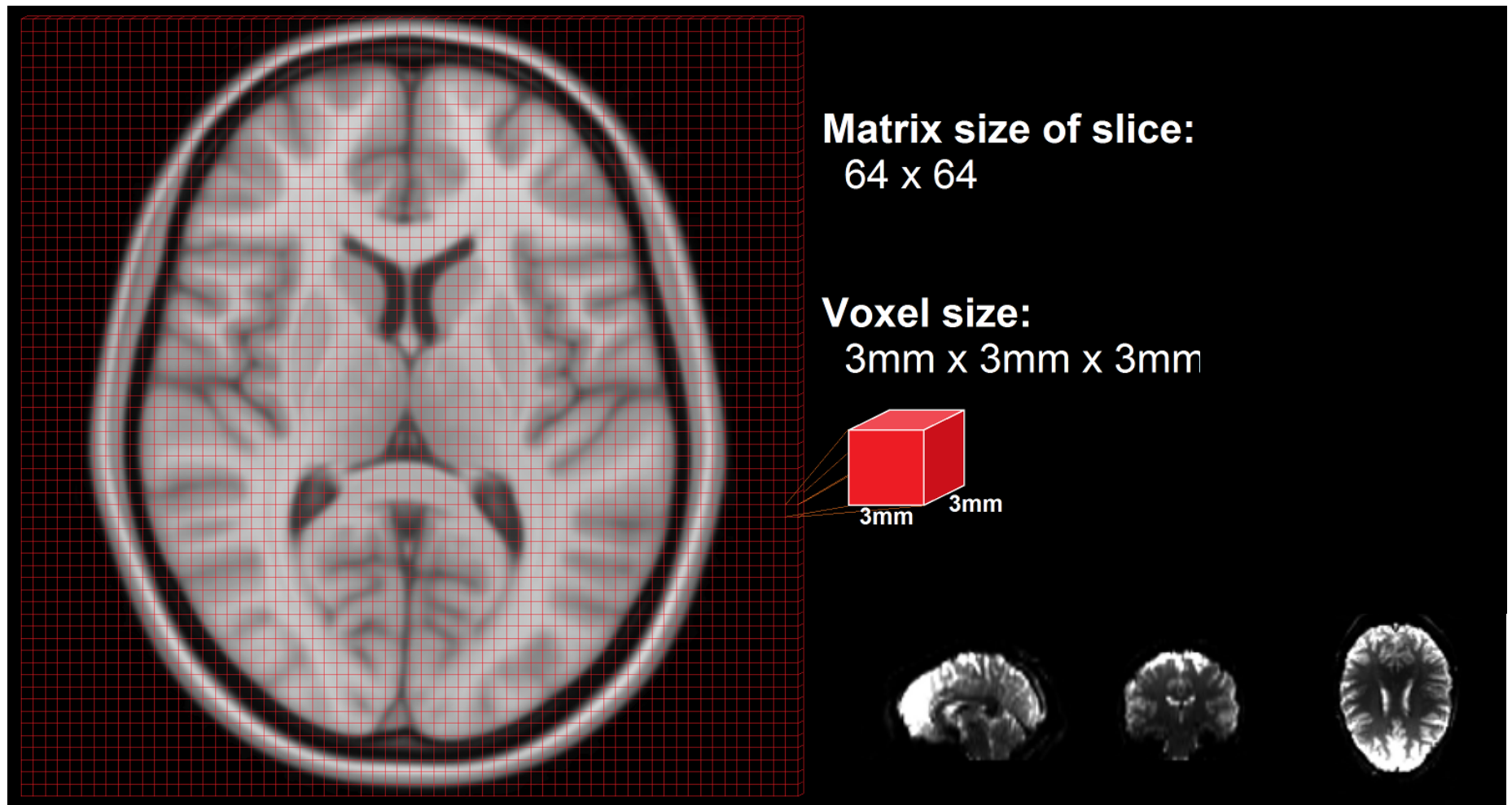


Image adapted from: <https://miykael.github.io/nipype-beginner-s-guide/neuroimaging.html>

# 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.

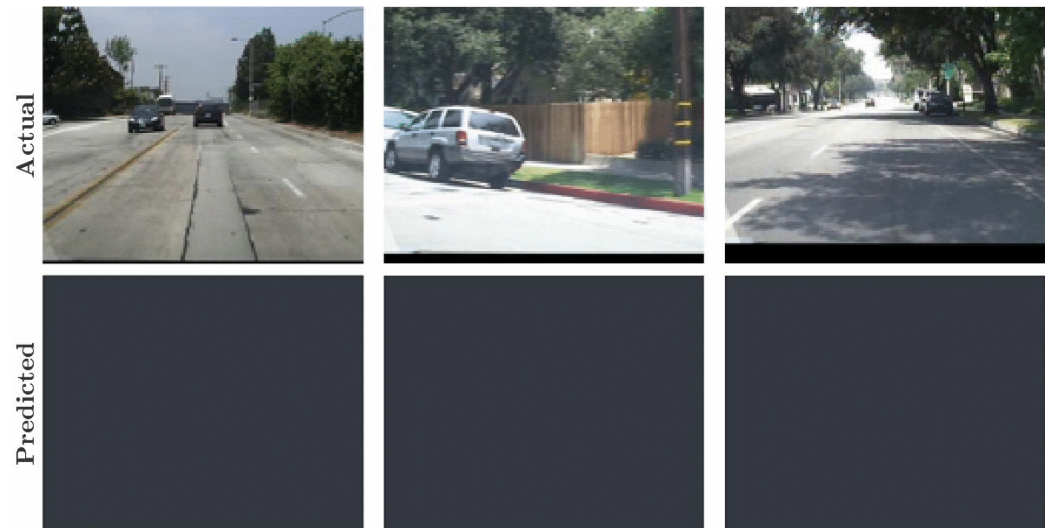
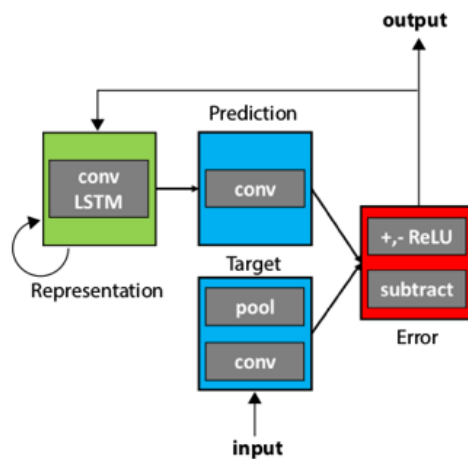
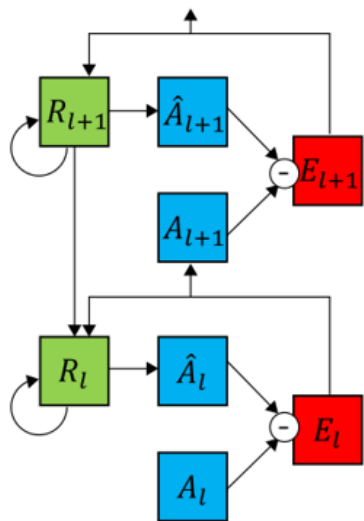
Subject RDMs were averaged together into a mean human brain RDM, which reduced noise.



# 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)

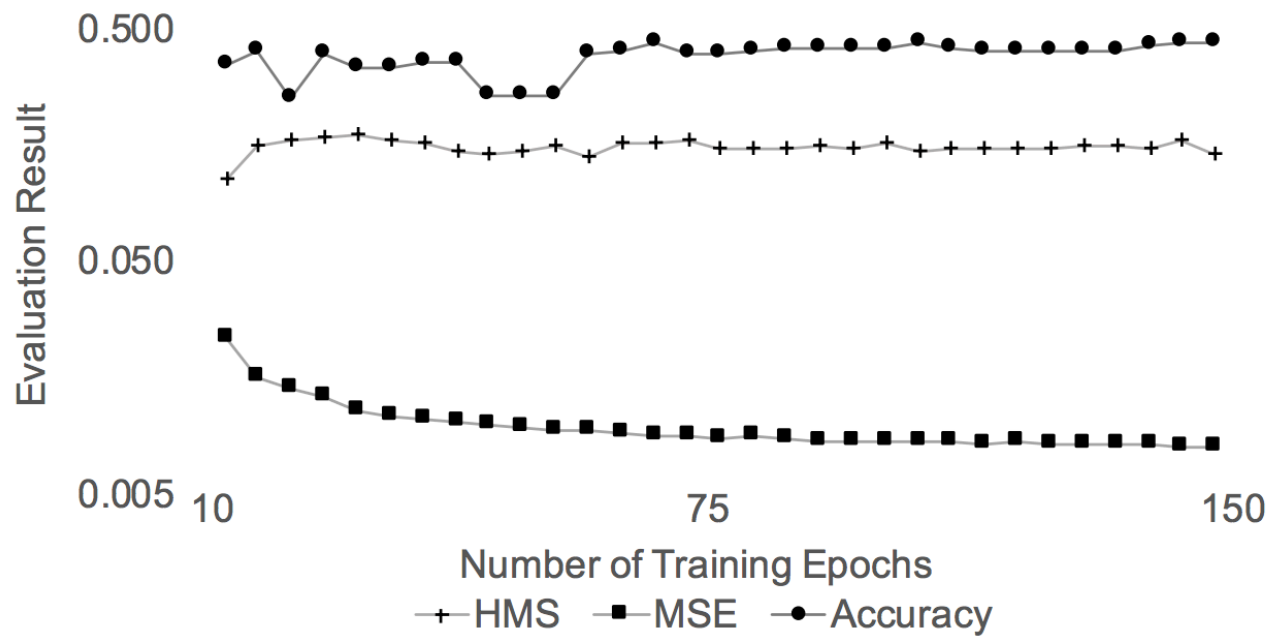


HMS is predictive of network performance  
on other metrics

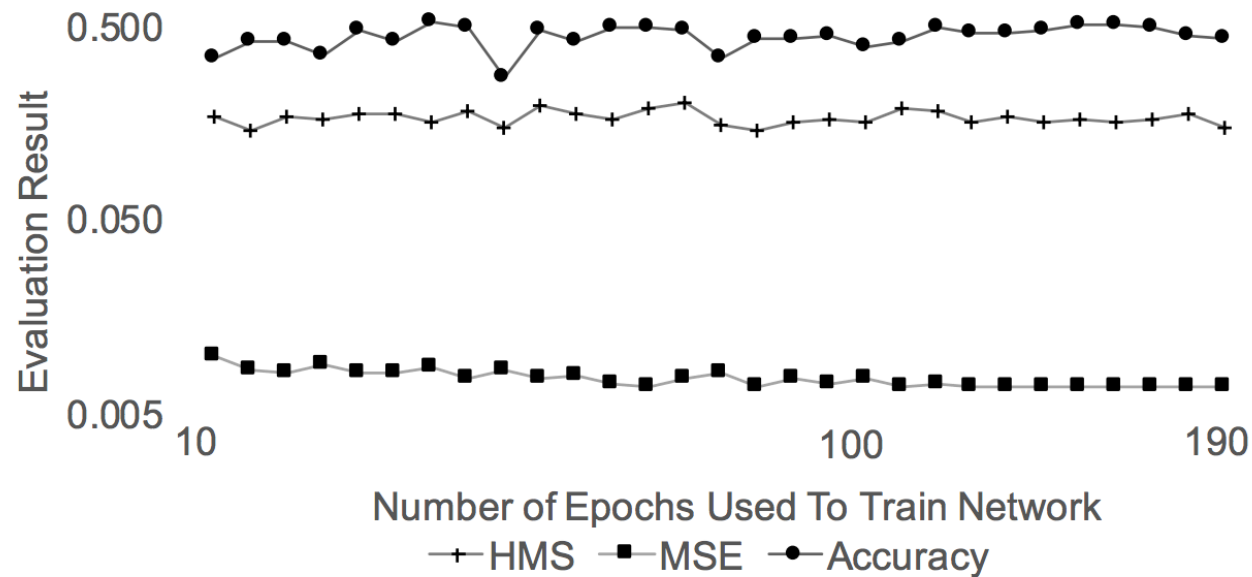
Variable	Accuracy	HMS	Learning Rate
Next Frame Prediction Error	-0.791**	<b>-0.646**</b>	0.635**
Object Matching Accuracy	.	<b>0.575**</b>	-0.517**
Human-Model Similarity	.	.	-0.452**

\*\* $p < 0.001$

# Within-Network Stability

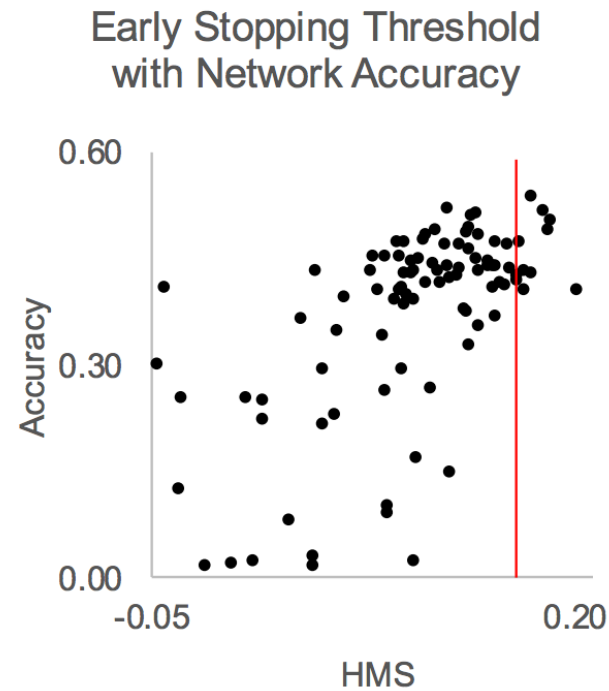
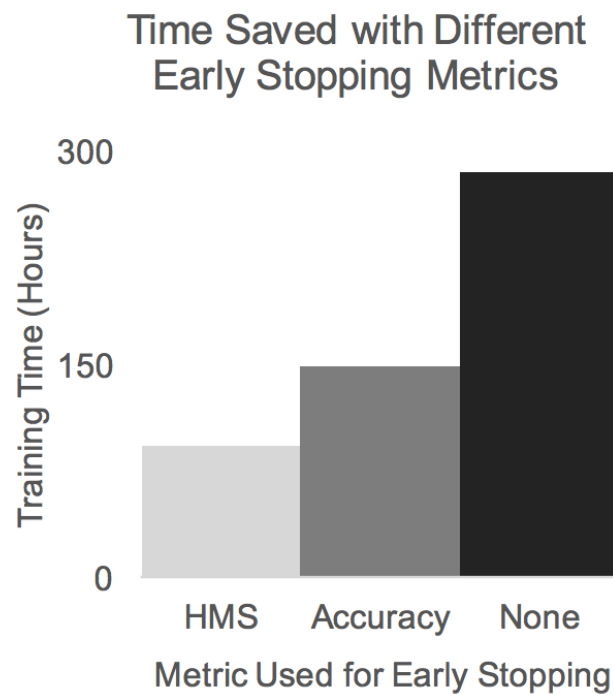


# Across-Network Stability



**66 Models, Mean**

# HMS-Driven Early Stopping

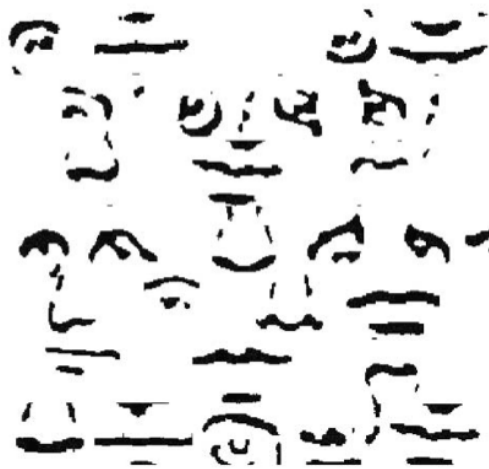


Is fMRI the best reference for this?

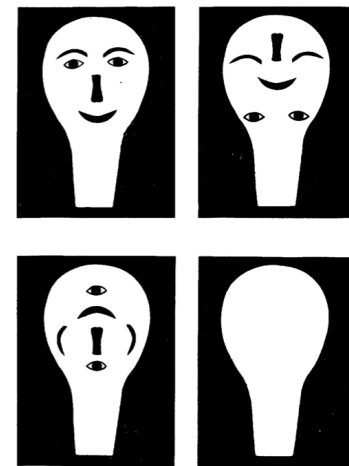
# Easier: Human Behavior

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!