## CSE 40171: Artificial Intelligence



Artificial Neural Networks with Functional Fidelity: Internal Behavior of Artificial and Biological Networks

### Homework #6 has been released It is due **tonight** at 11:59PM

# Project Updates are Due on 11/25 at 11:59PM

(See Course Website for Instructions)

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

# Monkey performing an object recognition task

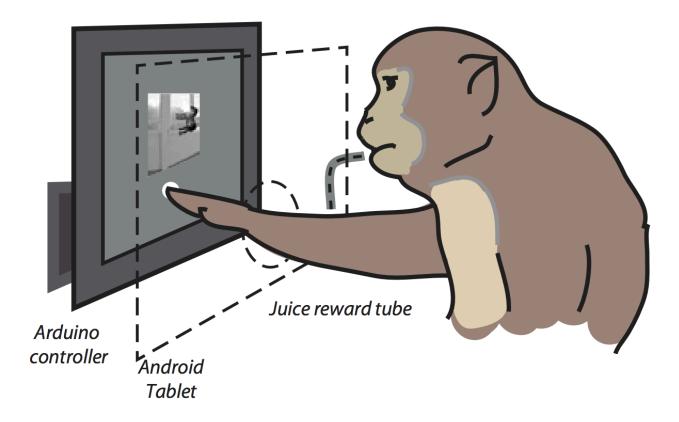
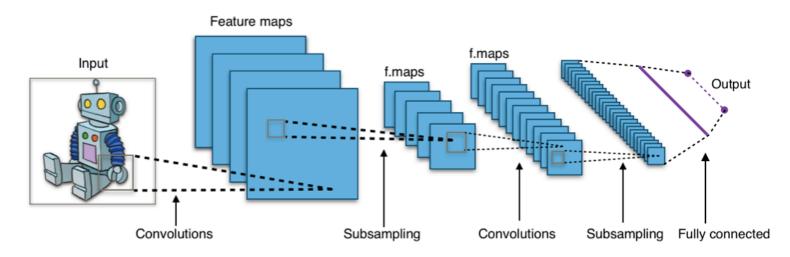


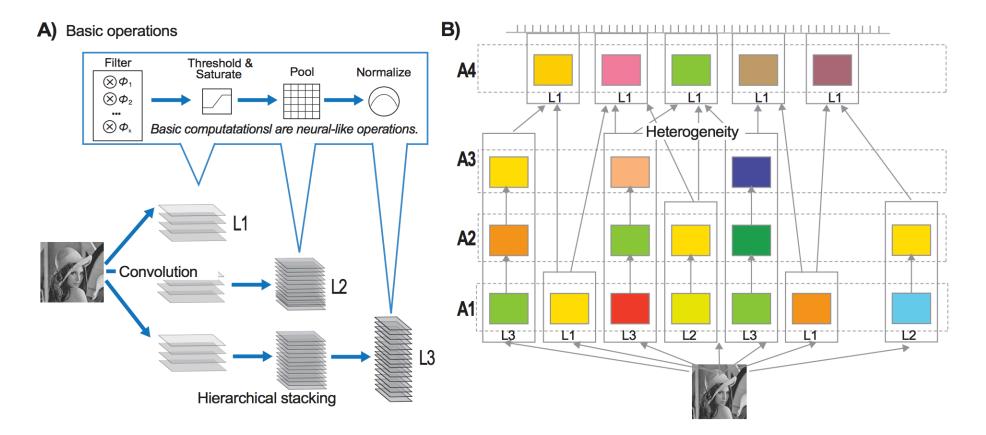
Image adapted from: Rajalingham et al. JNeurosci 2018

# CNN for Object Recognition



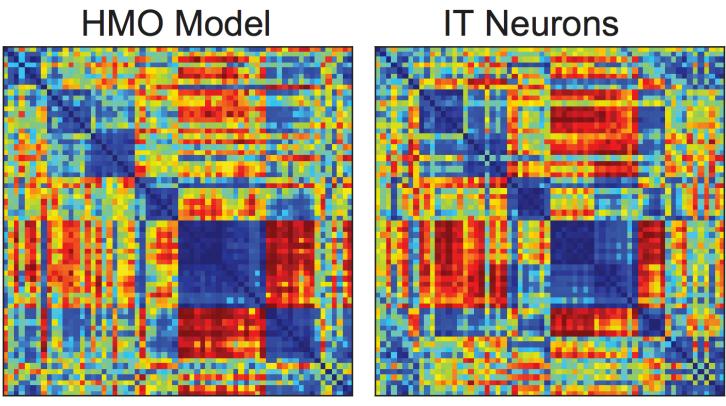
Typical CNN architecture 🞯 BY-SA 4.0 Aphex34

## Heterogeneous Hierarchical CNN



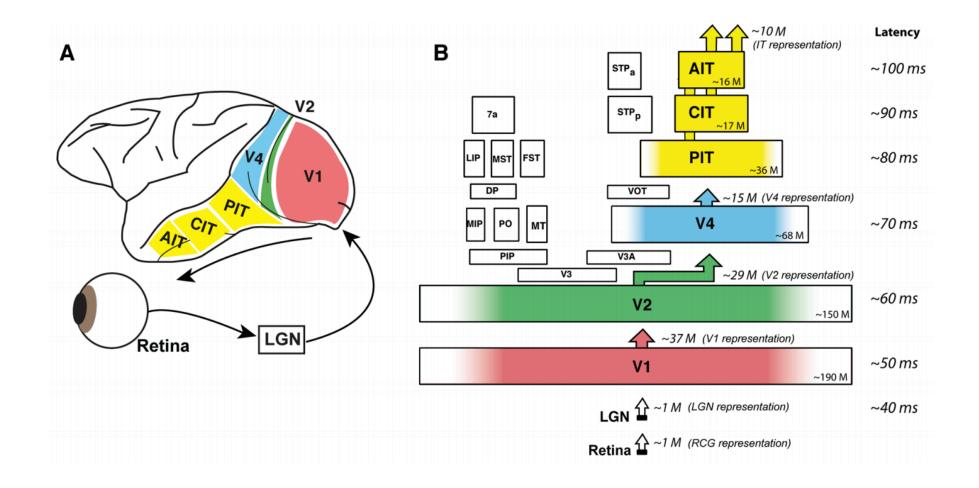
Yamins et al. NeurIPS 2013

# Population Responses: Model vs. Brain

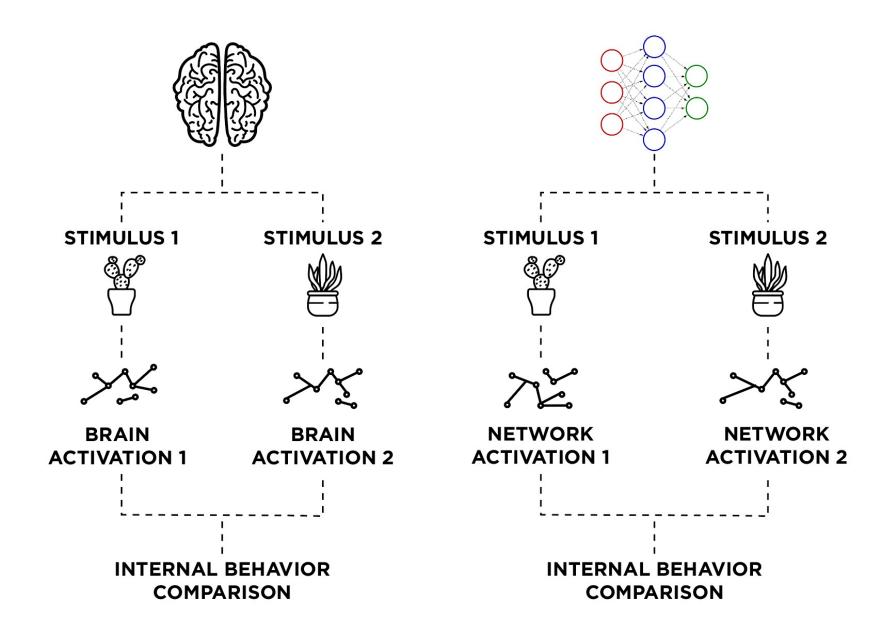


Yamins et al. NeurIPS 2013

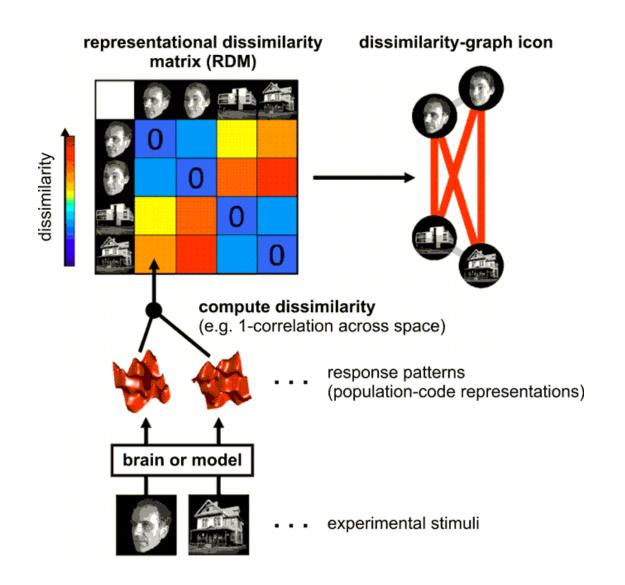
# Where in the brain is area IT?



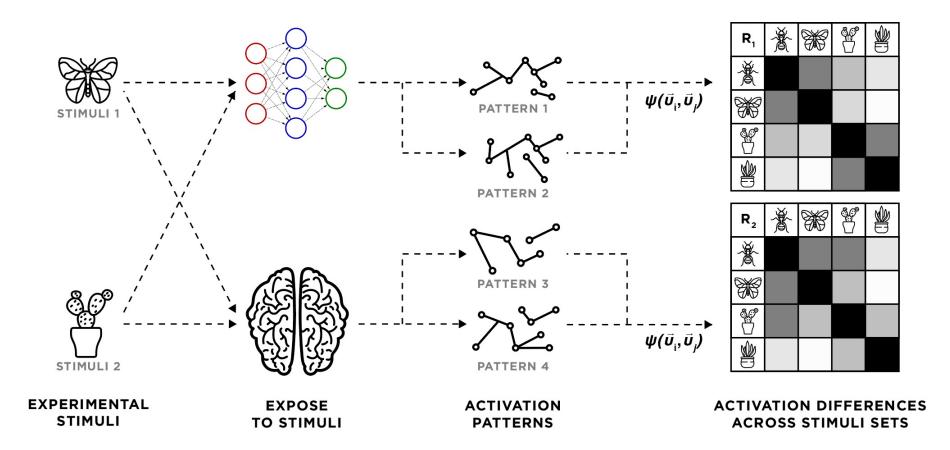
How do we compare the activity in brains with the activity in artificial neural networks?



### Representational Similarity Analysis



N. Kriegeskorte, M. Mur and P. A. Bandettini, "Representational Similarity Analysis – Connecting the Branches of Systems Neuroscience," Frontiers in Systems Neuroscience, 2008



### 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,  $f_1, f_2, \ldots, f_n$ , in response to *s*.

### **RDM Step 1: Data Representation**

If one expands the representation of *s* to a set of *m* stimuli  $S = s_1, s_2, ..., s_m$ , the natural extension of  $\overrightarrow{v}$  is the set of feature value collections  $V = \overrightarrow{v_1}, \overrightarrow{v_2}, ..., \overrightarrow{v_m}$ , in which  $s_i \in S$  is paired with  $\overrightarrow{v_i} \in V$  for each i = 1, 2, ..., m.

### RDM Step 2: Dissimilarity

Define the dissimilarity score between any two  $\overrightarrow{v_i} \in V$  and  $\overrightarrow{v_j} \in V$ :

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

### RDM Step 3: Construct Matrix

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

$$R = \begin{bmatrix} \psi(\vec{v}_{1}, \vec{v}_{2}) & \psi(\vec{v}_{1}, \vec{v}_{3}) & \dots & \psi(\vec{v}_{1}, \vec{v}_{m}) \\ \psi(\vec{v}_{2}, \vec{v}_{3}) & \dots & \psi(\vec{v}_{2}, \vec{v}_{m}) \\ & \ddots & \vdots \\ \psi(\vec{v}_{m-1}, \vec{v}_{m}) \end{bmatrix}$$

# 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  $1.95 \times 1.95 \times 2$ mm<sup>3</sup> 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.

H. Nili, C. Wingfield, A. Walther, L. Su, W. Marslen-Wilson, and N. Kriegeskorte. A toolbox for representational similarity analysis. PLoS Computational Biology, 10(4):e1003553, 2014.



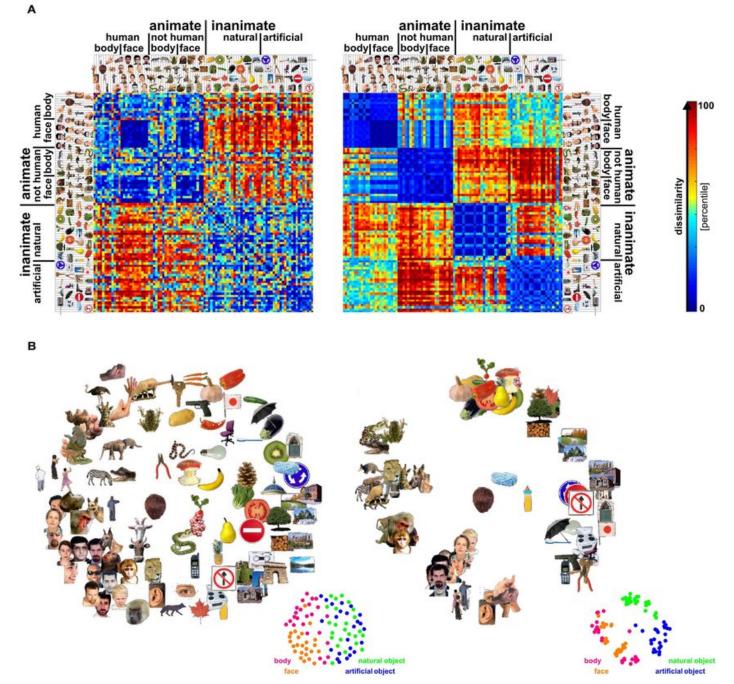
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## fMRI Stimuli Set

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#### hIT activity patterns

#### human similarity judgments



Mur et al. Frontiers in Psychology 2013

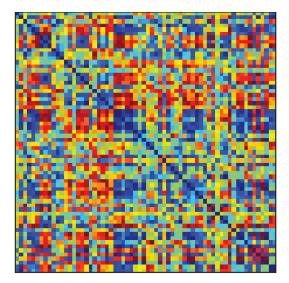
# Back to the brain vs. model experiment...

### Electrophysiology Data: Yamins et al. 2013

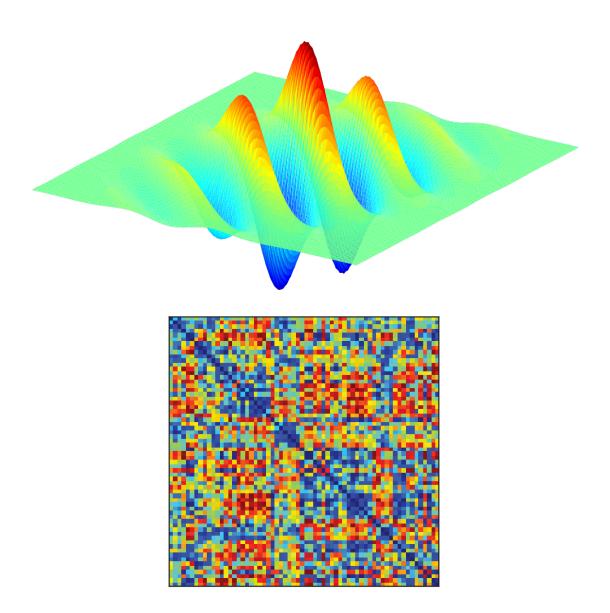
- Large-scale parallel array electrophysiology recordings in the visual cortex of awake behaving macaques
- Passively viewing animals shown random stimulus sequences with durations comparable to those in natural primate fixations (200 ms)
- Electrode arrays were surgically implanted in V4 and IT, and recordings took place daily over a period of months
- A total of 296 multi-unit responses were recorded from two animals
- For each testing stimulus and neuron, final neuron output responses were obtained by averaging data from between 25 and 50 repeated trials

## Control Model: Pixels





## Control Model: V1-Like



# Control Model: SIFT

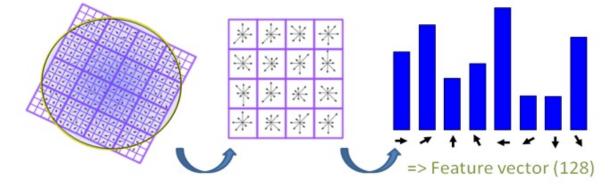
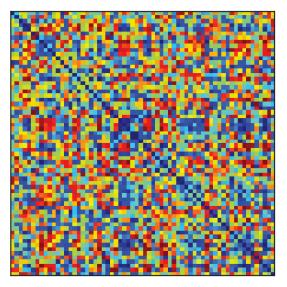
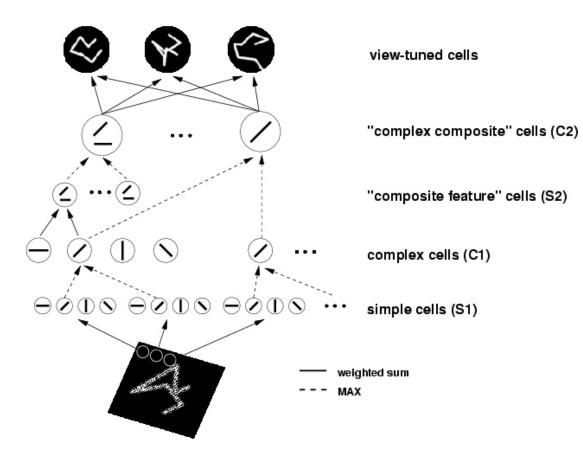
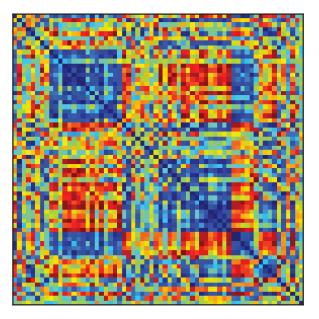


Image Credit: https://gilscvblog.com/2013/08/18/a-short-introduction-to-descriptors/



# Control Model: HMAX

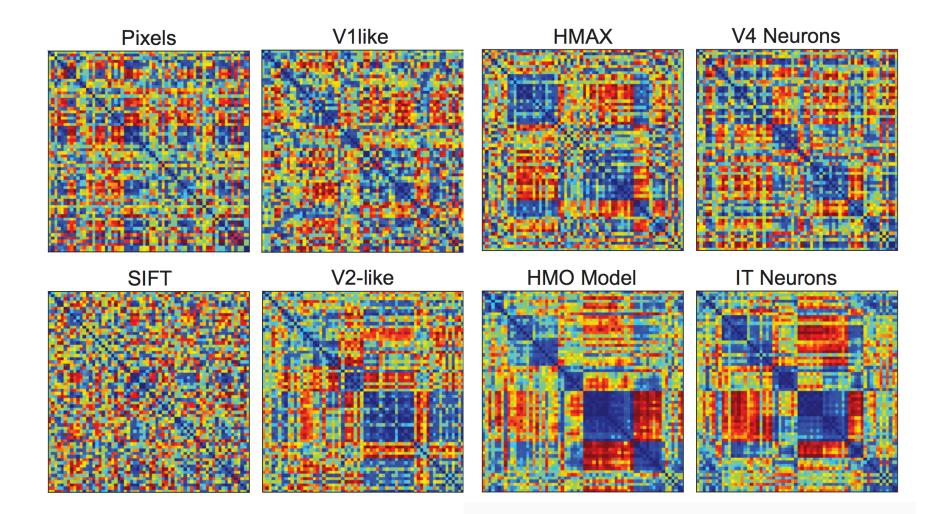




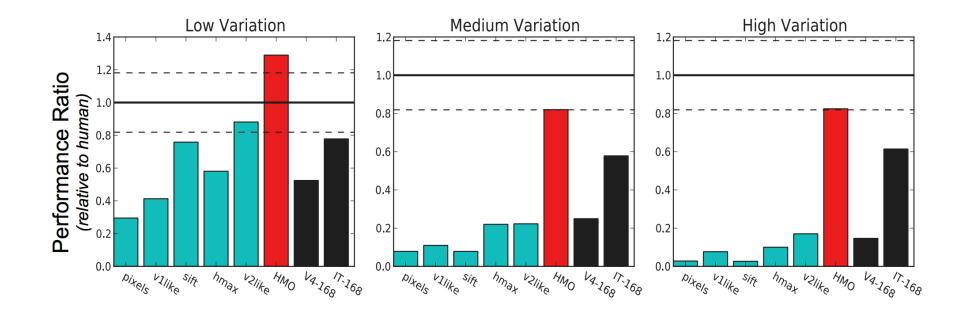
Riesenhuber and Poggio Nat. NeuroSci. 1999

Yamins et al. NeurIPS 2013

### Comparison to Monkey Recordings

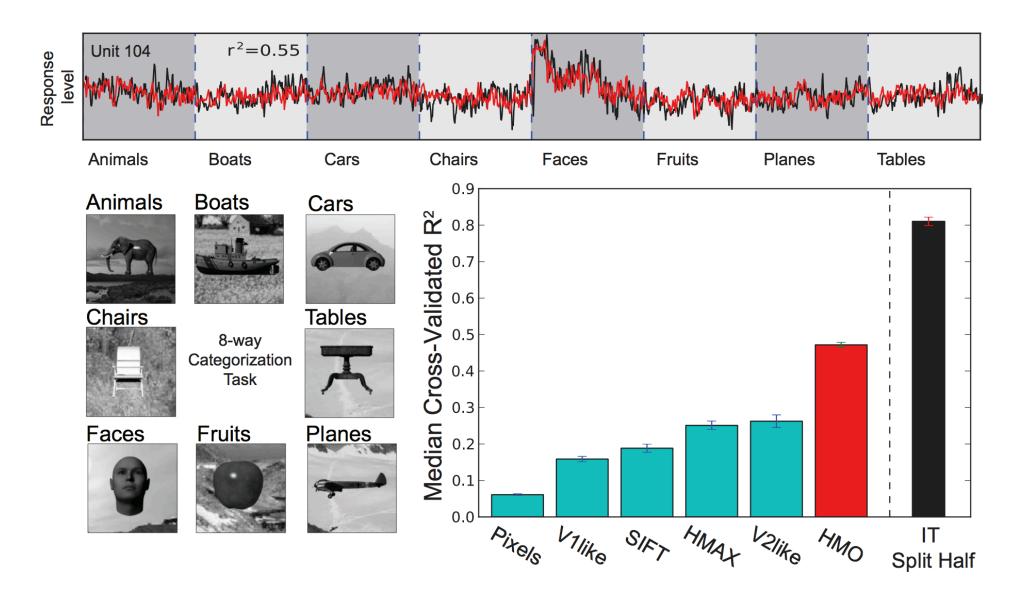


### 8-way Categorization Performances

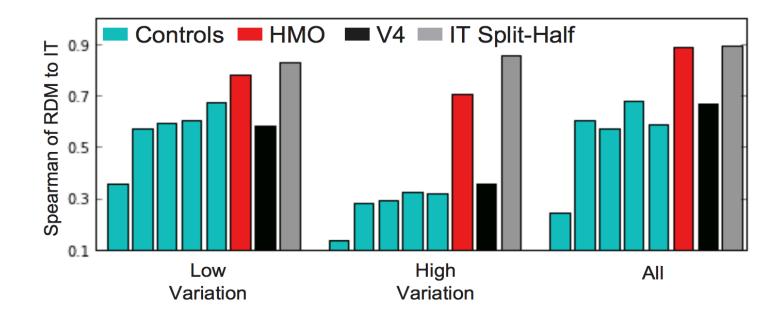


#### Black curve: actual neural response

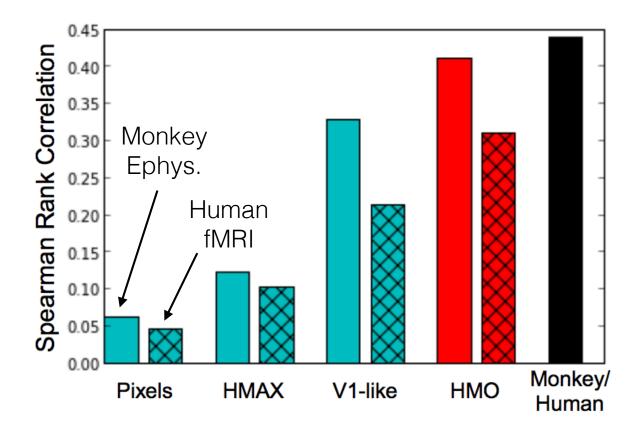
### Red curve: prediction for a single sample IT neuron



# Comparison of RDMs

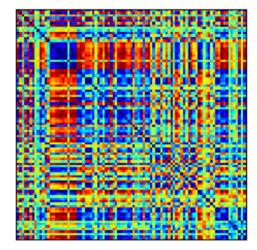


# Model comparison to monkey IT and human ventral stream

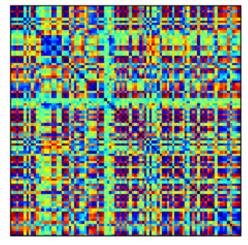


## **RDM** Comparison

#### **Pixels**

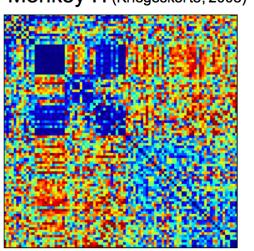


HMAX

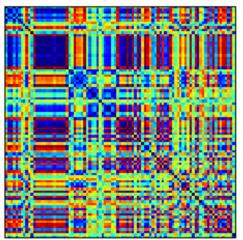


Monkey IT (Kriegeskorte, 2008)

HMO



V1-like



Human (Kriegeskorte, 2008)

