

# CS375 / Psych 249:

## Large-Scale Neural Network Models for Neuroscience

---

### Lecture 4: Model-Brain Mapping Methods

2025.01.14-16

Atlas Kazemian

Departments of Psychology  
Stanford Neuroscience and Artificial Intelligence Laboratory  
Stanford University



# Outline

- Comparison methods
  1. The early days
    - Examples: subjective comparisons, sparsity, response properties.
  2. Using stimulus-by-stimulus similarity matrices.
    - Examples: RSA, CKA
  3. Learning a mapping from models to neural data.
    - Examples: One to one matching, linear regression, procrustes, soft matching, nonlinear mapping
- Selecting the right method:
  - Bidirectionally vs symmetry.
  - Using IATC for choosing the correct metric
- Noise ceiling estimates

# Why do we compare neural networks to the brain?

As **scientists** we care about understanding the brain:

- Does the model encode similar features as neural populations?
- Is the model solving the task using similar transformations?
- Which architectural or learning constraints allow us to better explain neural responses.

As **engineers** we care about building a good model of the brain:

- Models allow rapid, large-scale testing of hypotheses that would be infeasible in humans or animals (ablation studies)
- Models can inform brain–computer interfaces and personalized treatments.

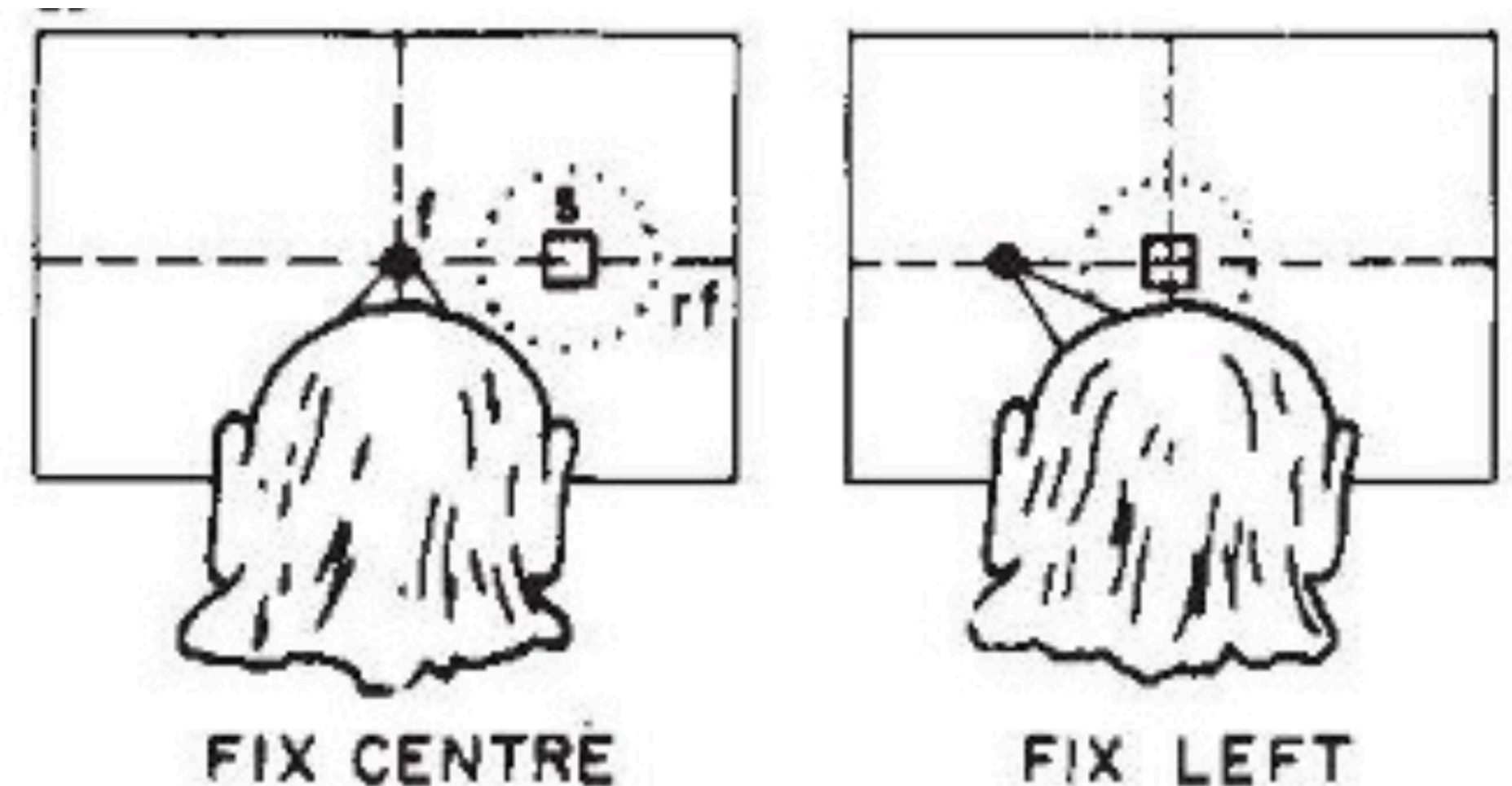
# Early days: subjective comparisons

**Idea:** subjectively compare properties in models and neural data

**Zipser & Andersen (1988):** Study how the brain encodes retinal location and eye position together to represent object location in posterior parietal cortex.

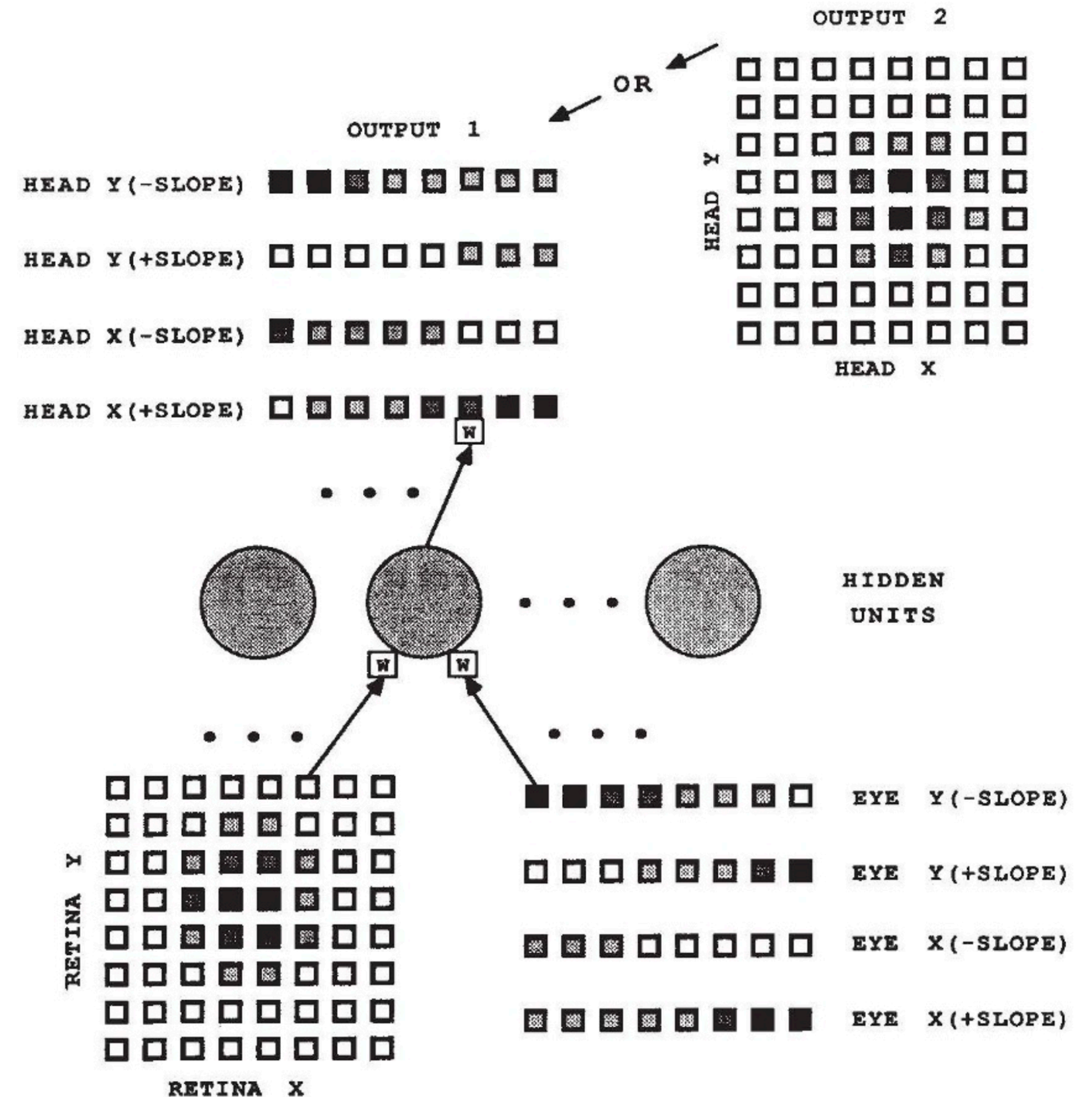
## Neural data

- Single-unit recordings from area 7a in awake monkeys
- Visual stimulus is flashed at many retinal (x, y) locations during fixation
- Firing rate measured for each location



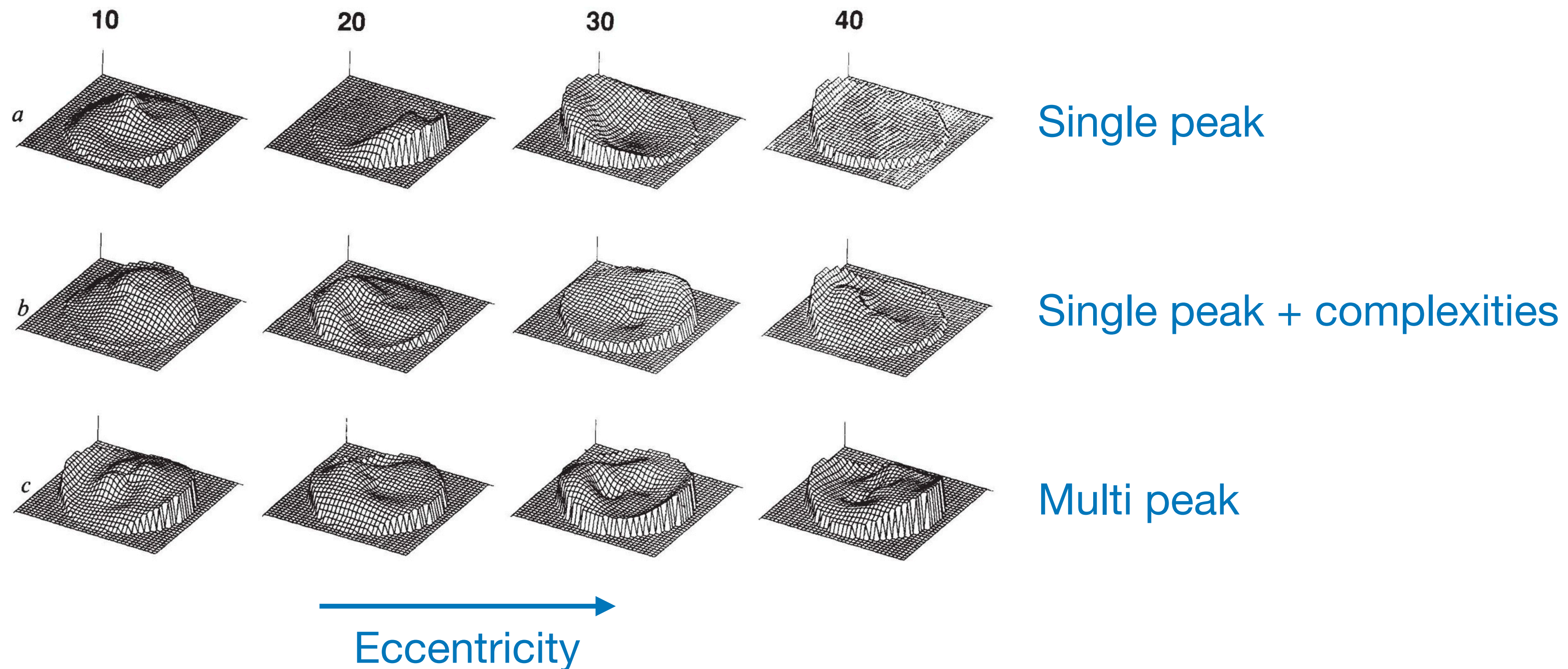
# Neural Network Model

- 3-layer feedforward trained with backpropagation
- **Inputs:**
  1. Retinal position
  2. Eye position
- **Task:** Learn head-centered target locations



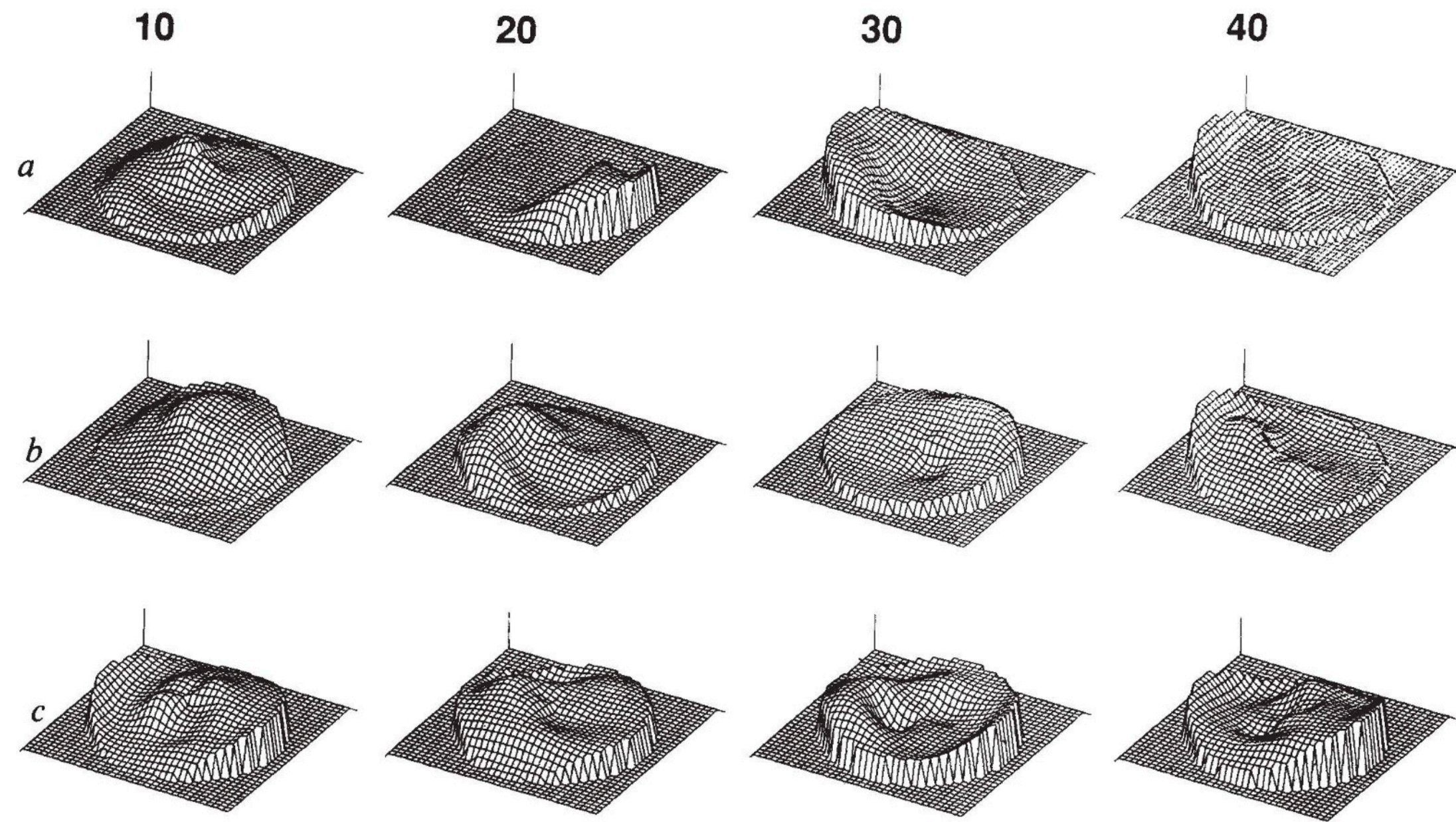
# Comparing neural data with the model

## Monkey receptive fields

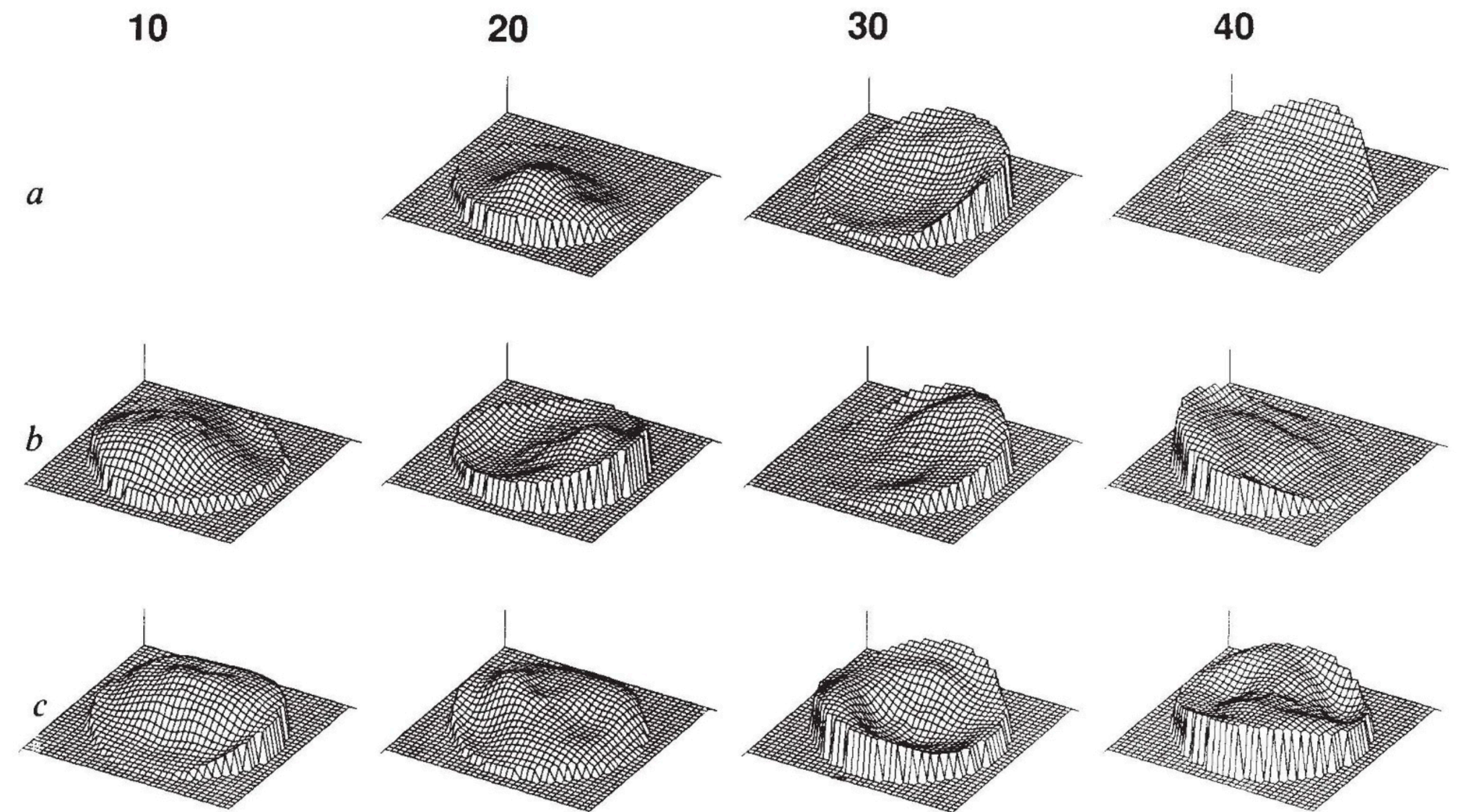


# Comparing neural data with the model

**Monkey receptive fields**



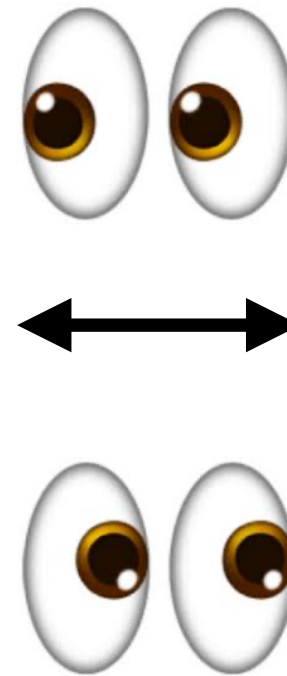
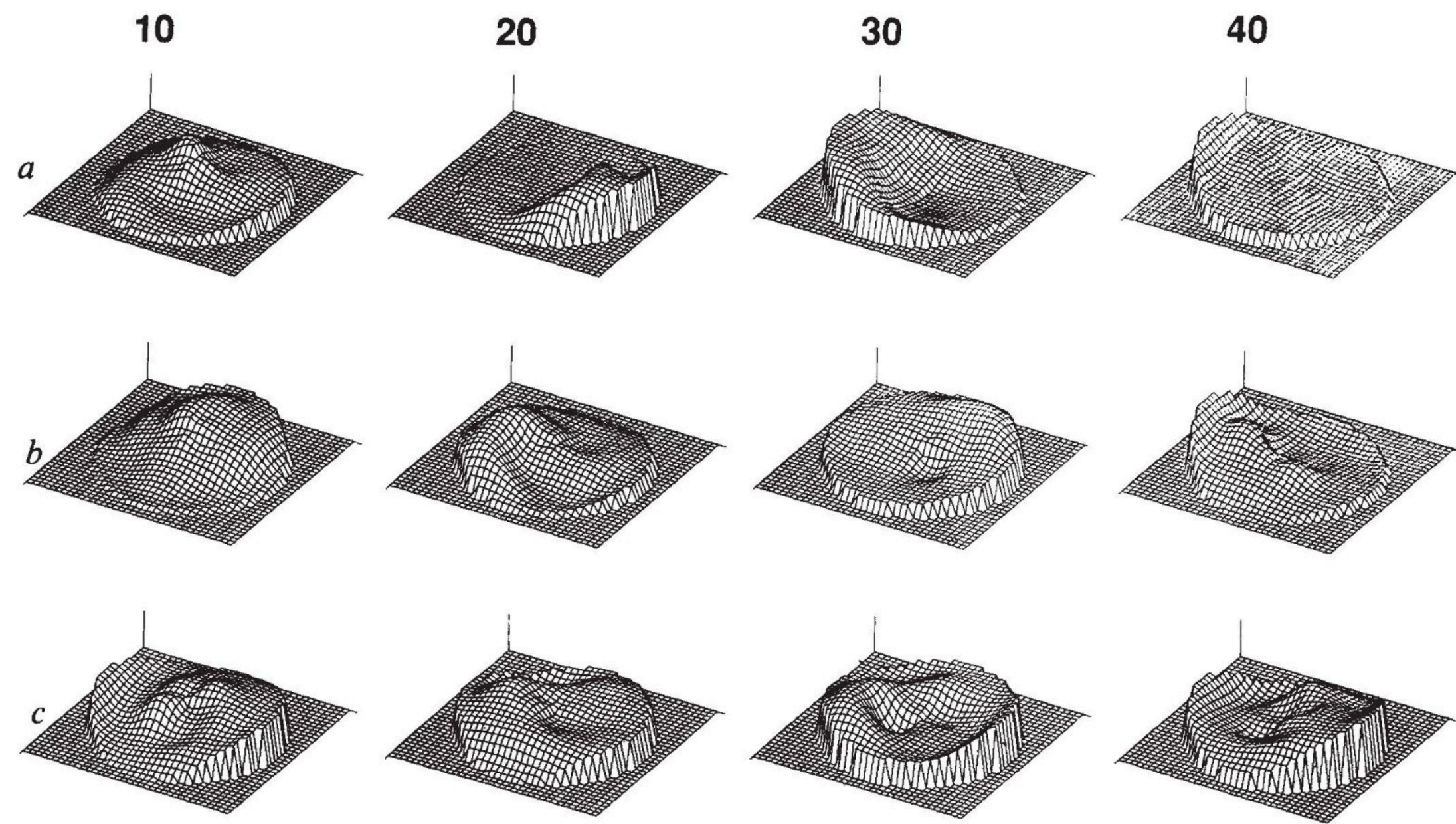
**Model receptive fields**



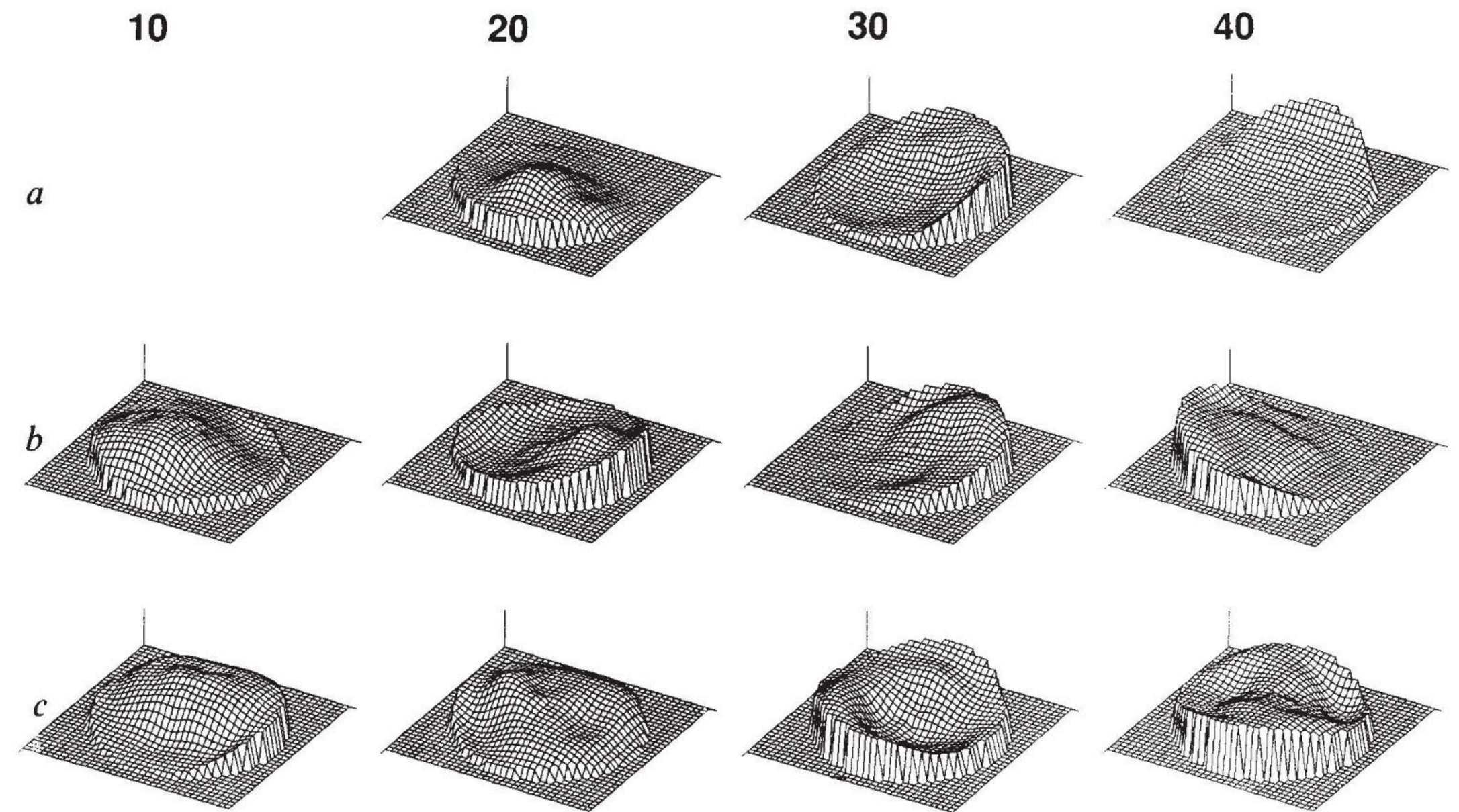
# Comparing neural data with the model

*...“The comparison process contains an element of subjectivity, but it demonstrates that the trained model generates retinal receptive fields remarkably similar to the experimentally observed fields.”...*

**Monkey receptive fields**



**Model receptive fields**



# Early days: comparing sparseness

**Idea:** Move beyond subjective comparison by comparing sparseness and population statistics

**Rolls & Tovee (1995):** Are object representation in IT encoded using a dense, localist, or sparse distributed code?

## Neural data

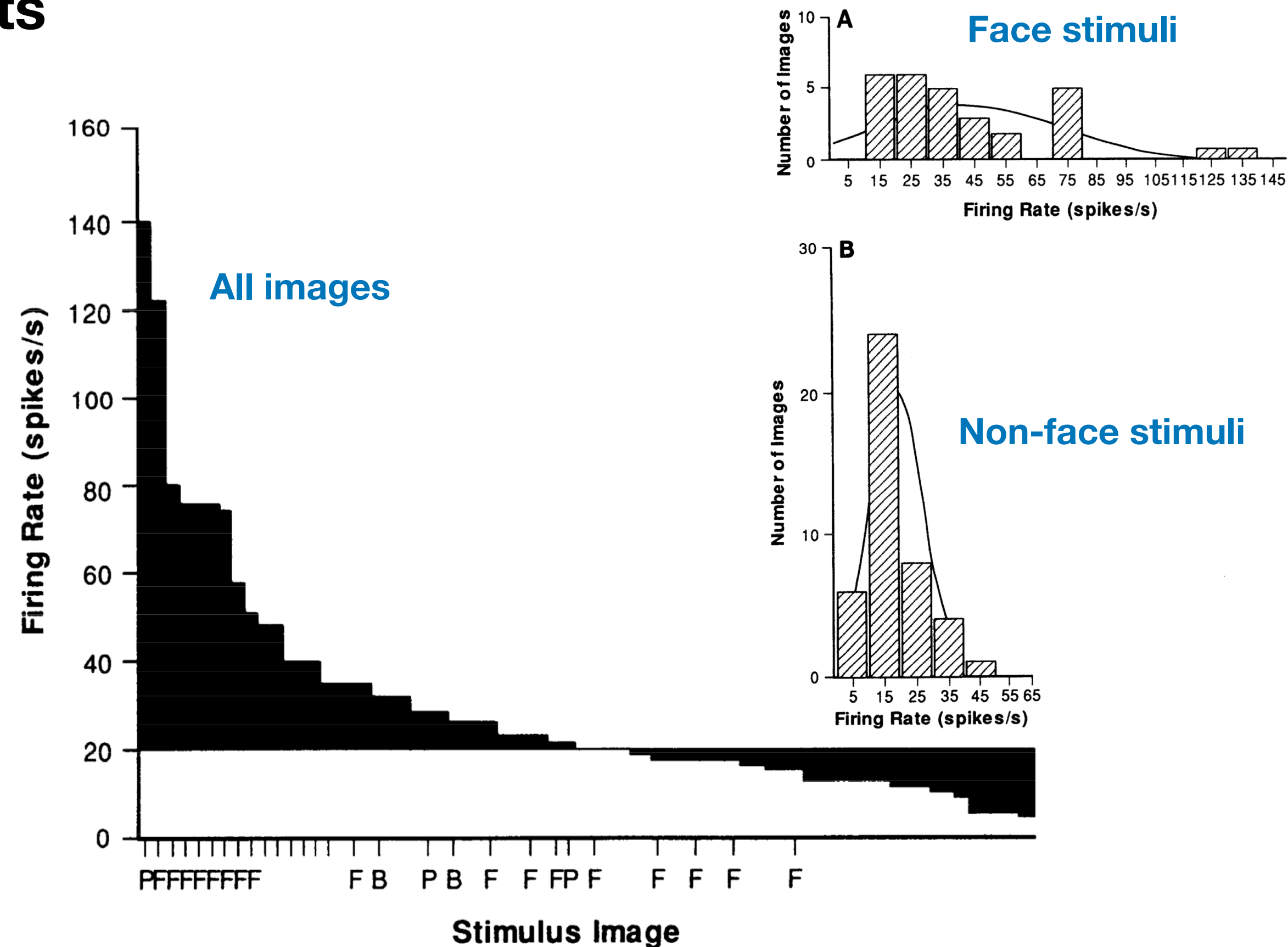
- Single-unit recordings from macaque IT
- Monkeys viewed large “diverse” sets of complex visual stimuli (objects, faces, scenes)



## Representational theories

1. **Dense distributed coding:** Many neurons active for most stimuli
2. **Localist (grandmother-cell) coding:** One neuron per object
3. **Sparse distributed coding:** Few neurons active per stimulus

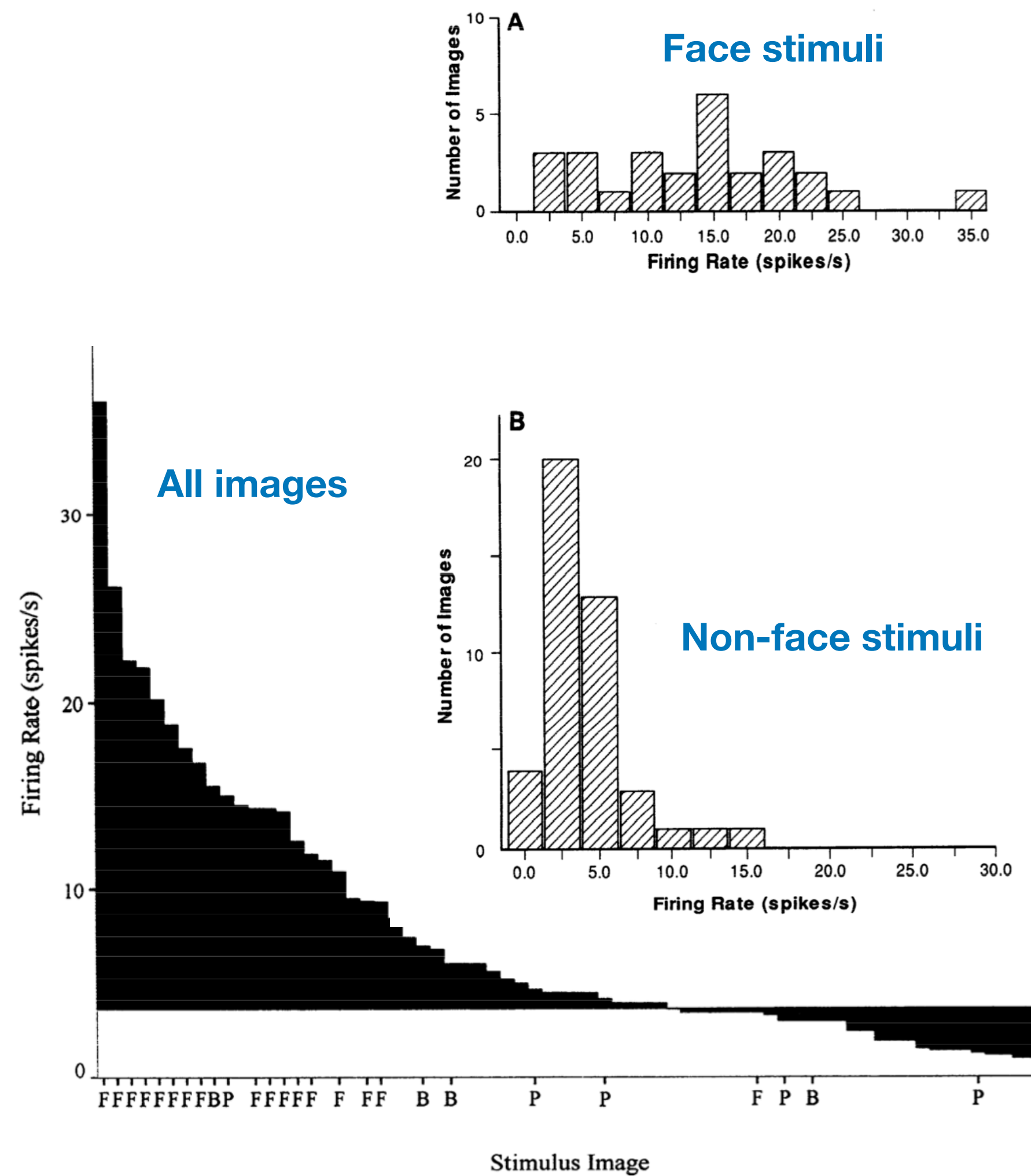
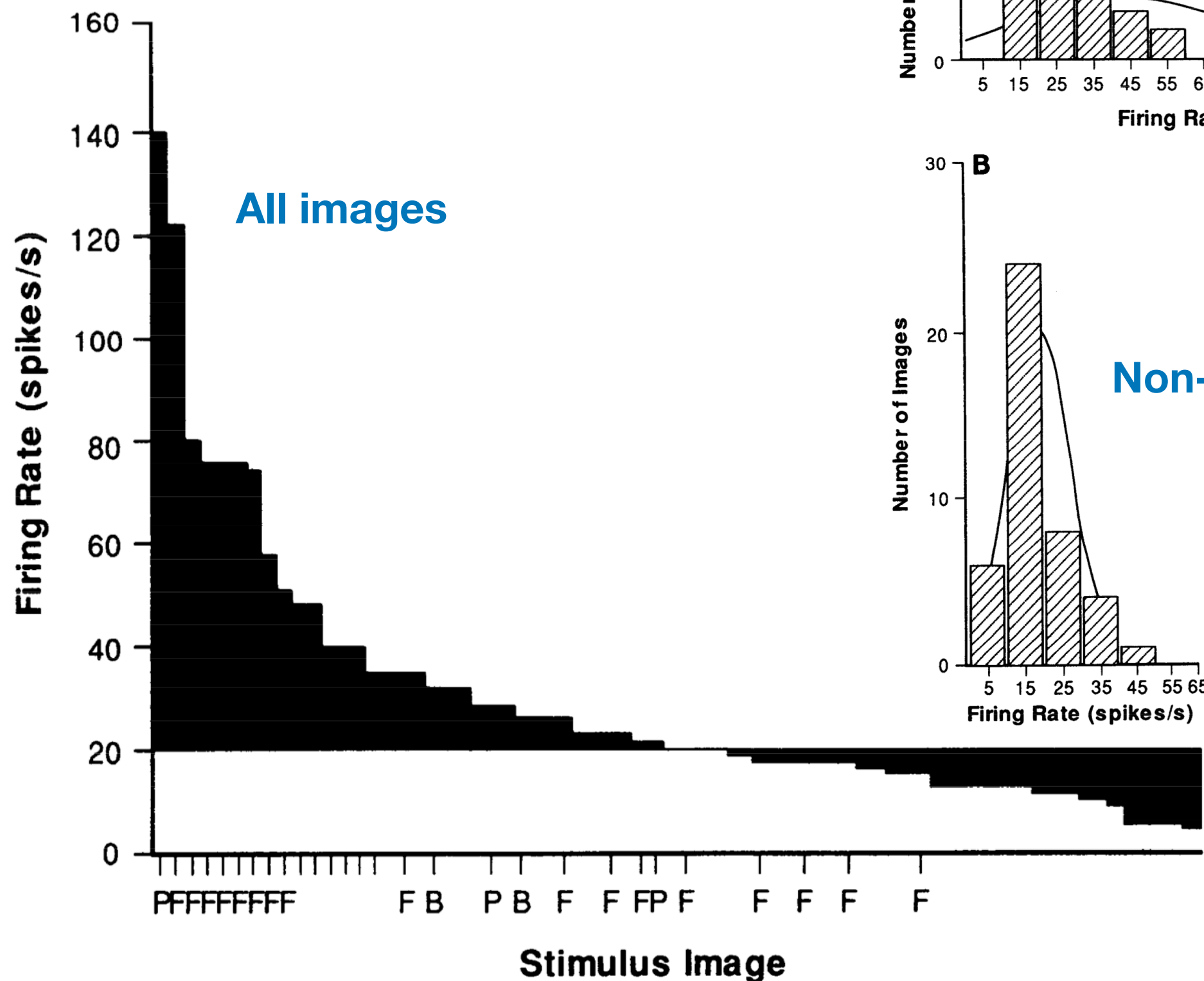
## Results



# Representational theories

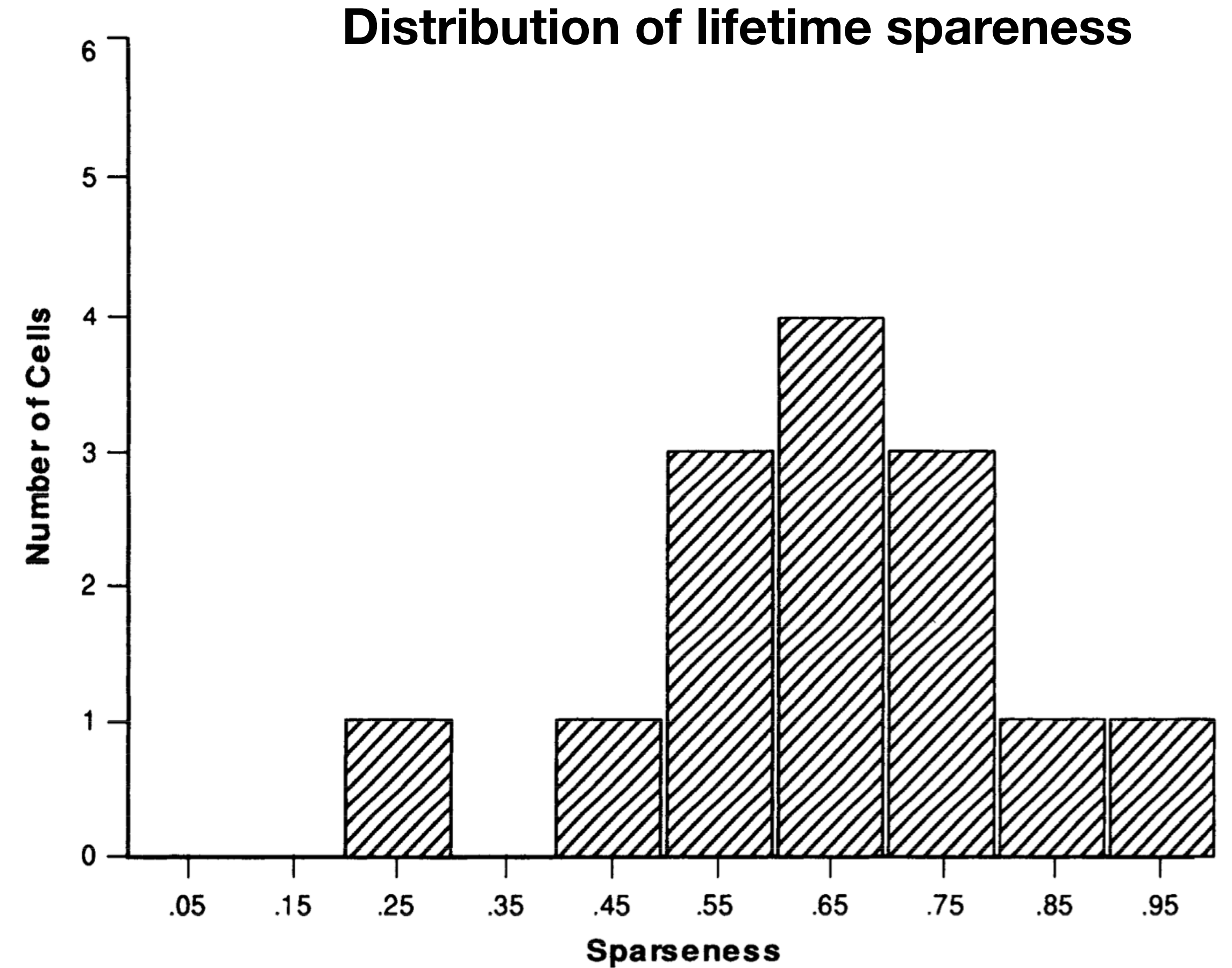
- 1. **Dense distributed coding:** Many neurons active for most stimuli
- 2. **Localist (grandmother-cell) coding:** One neuron per object
- 3. **Sparse distributed coding:** Few neurons active per stimulus

## Results



# Comparing neural data with theory

- **Lifetime sparseness:** how selectively a *neuron* responds across a large set of different visual stimuli over its lifetime.
- “Which class of representations could plausibly generate these neural response statistics?”



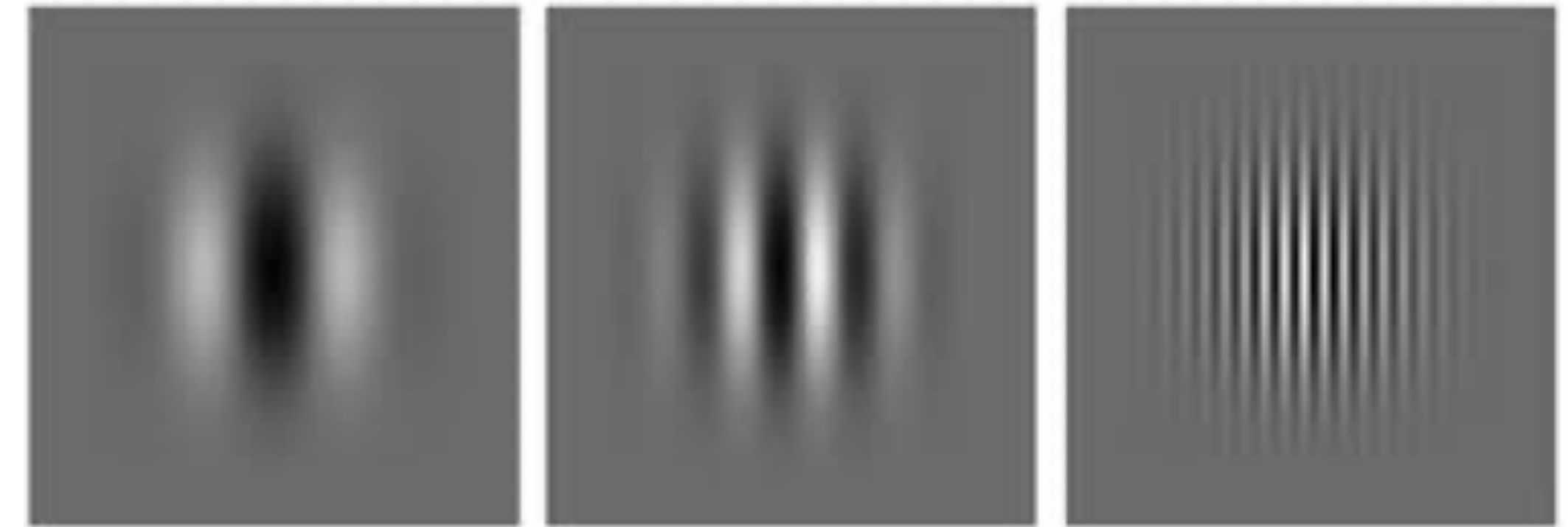
*"The mean response sparseness of **0.60** of this population of face-selective neurons indicates that, **within the class faces**, these neurons implement **distributed encoding**"*

# Early day: Comparing response properties

**Idea:** compare tuning properties of cells with those of networks

**De Valois et al. (1982):**

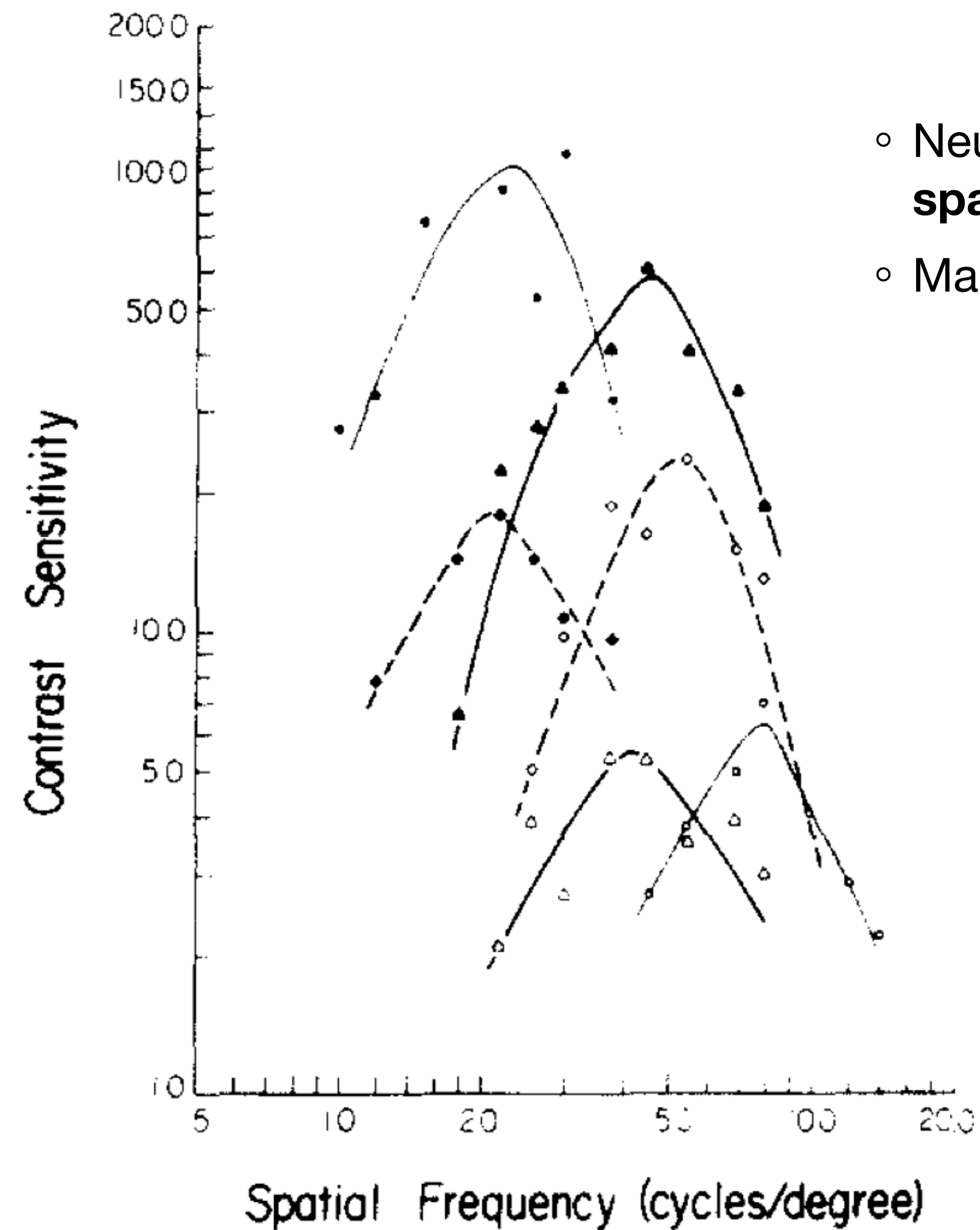
- How selectively do V1 neurons respond to different spatial frequencies in sine gratings? (Are V1 neurons bandpass filters?)



## Neural data

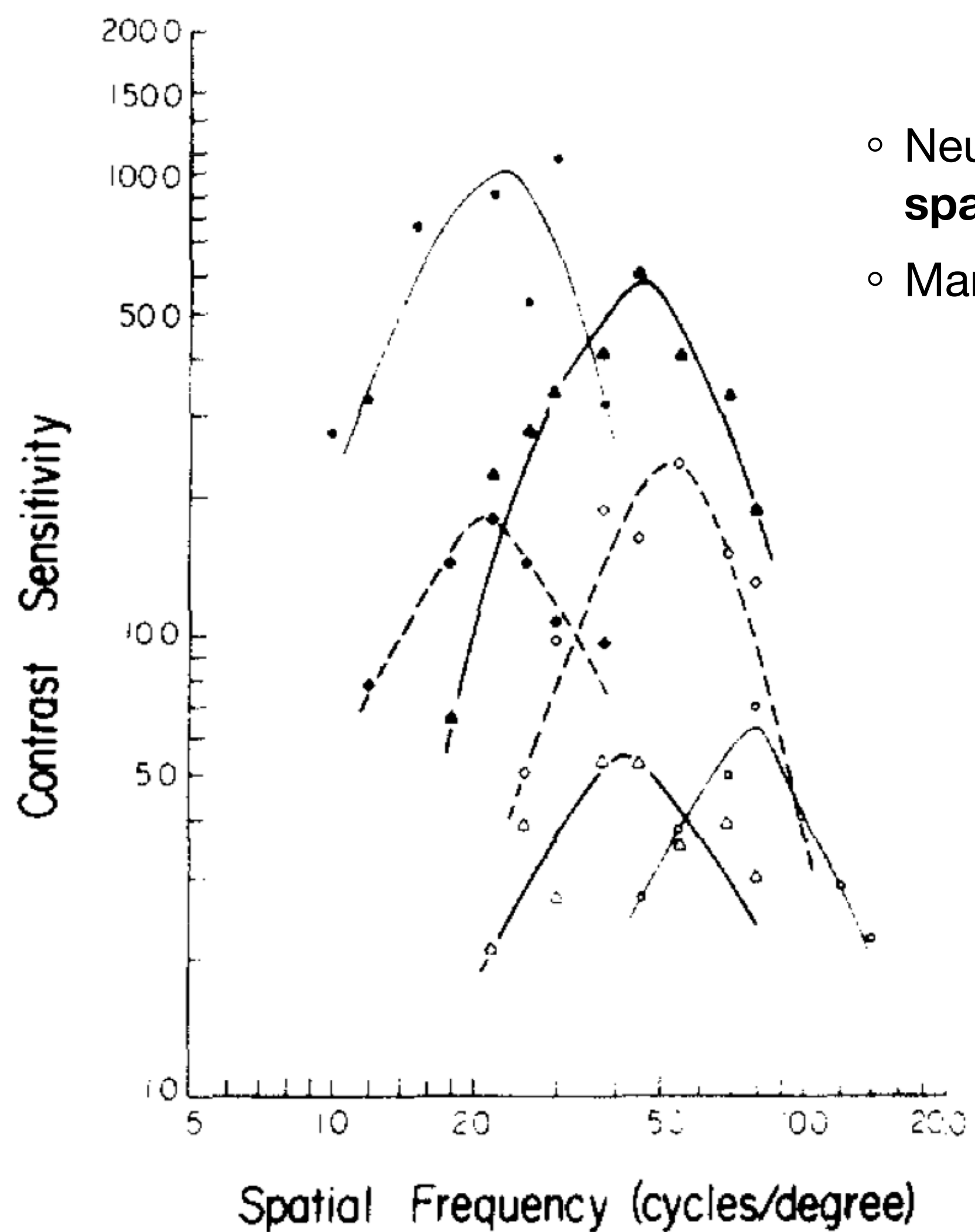
- single-unit recordings from macaque V1
- Present sinusoidal gratings at many orientations and spatial frequencies
- Spatial frequency = Number of cycles (dark-light)/ visual angle (degrees)

# Comparing results to neural networks

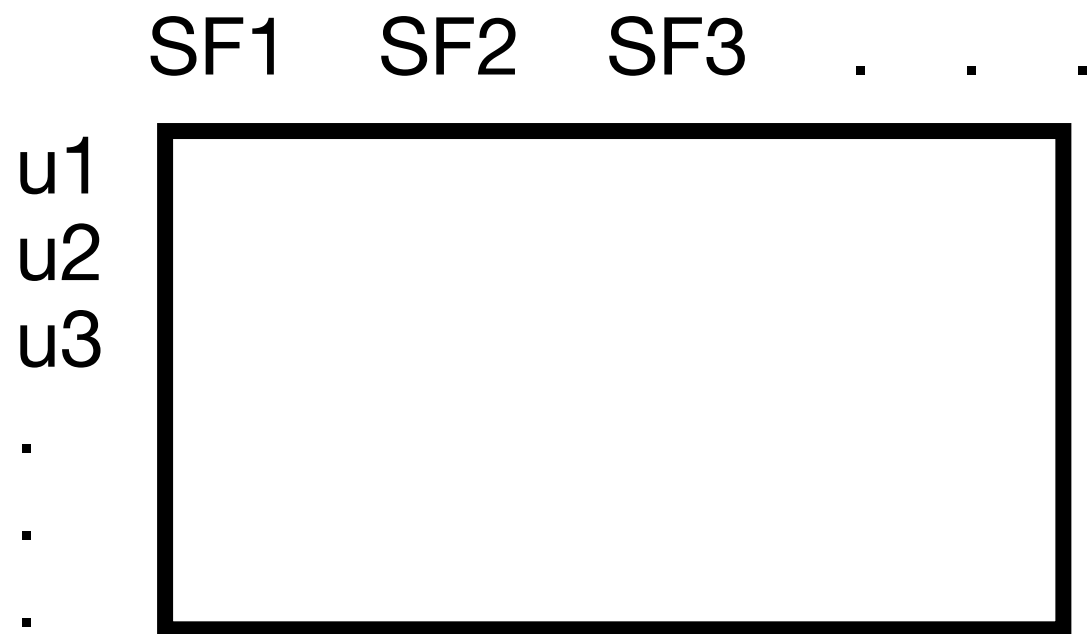


- Neurons span a **wide range of preferred spatial frequencies**
- Many neurons are **narrowly tuned**

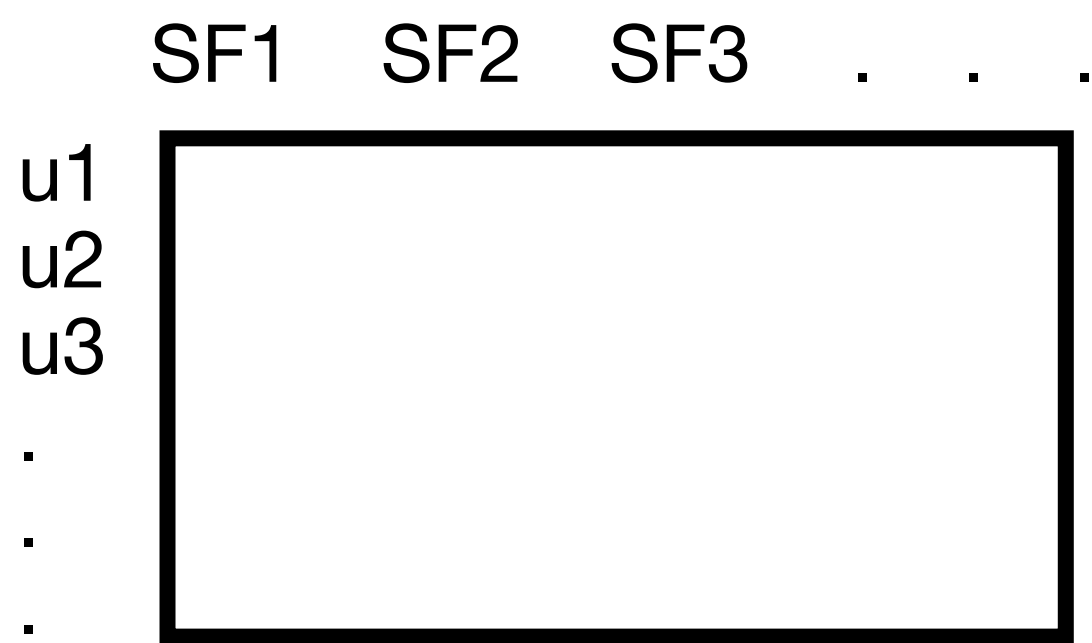
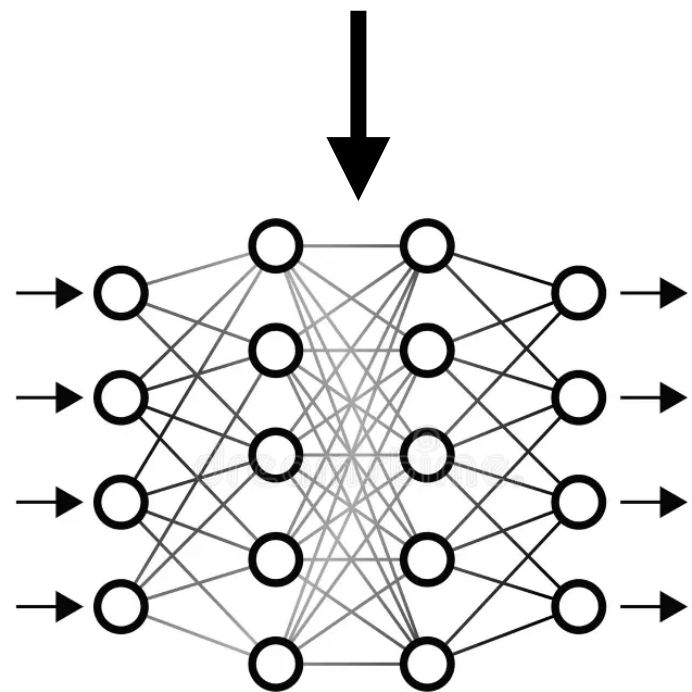
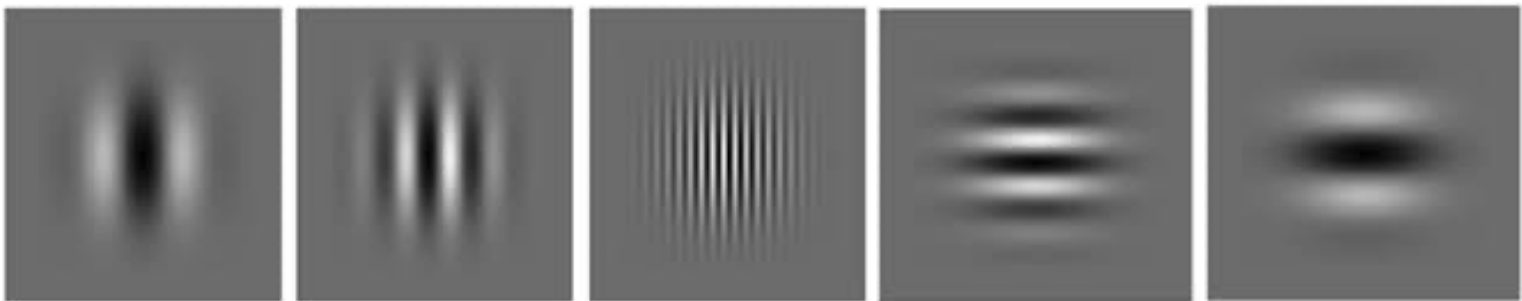
# Comparing results to neural networks



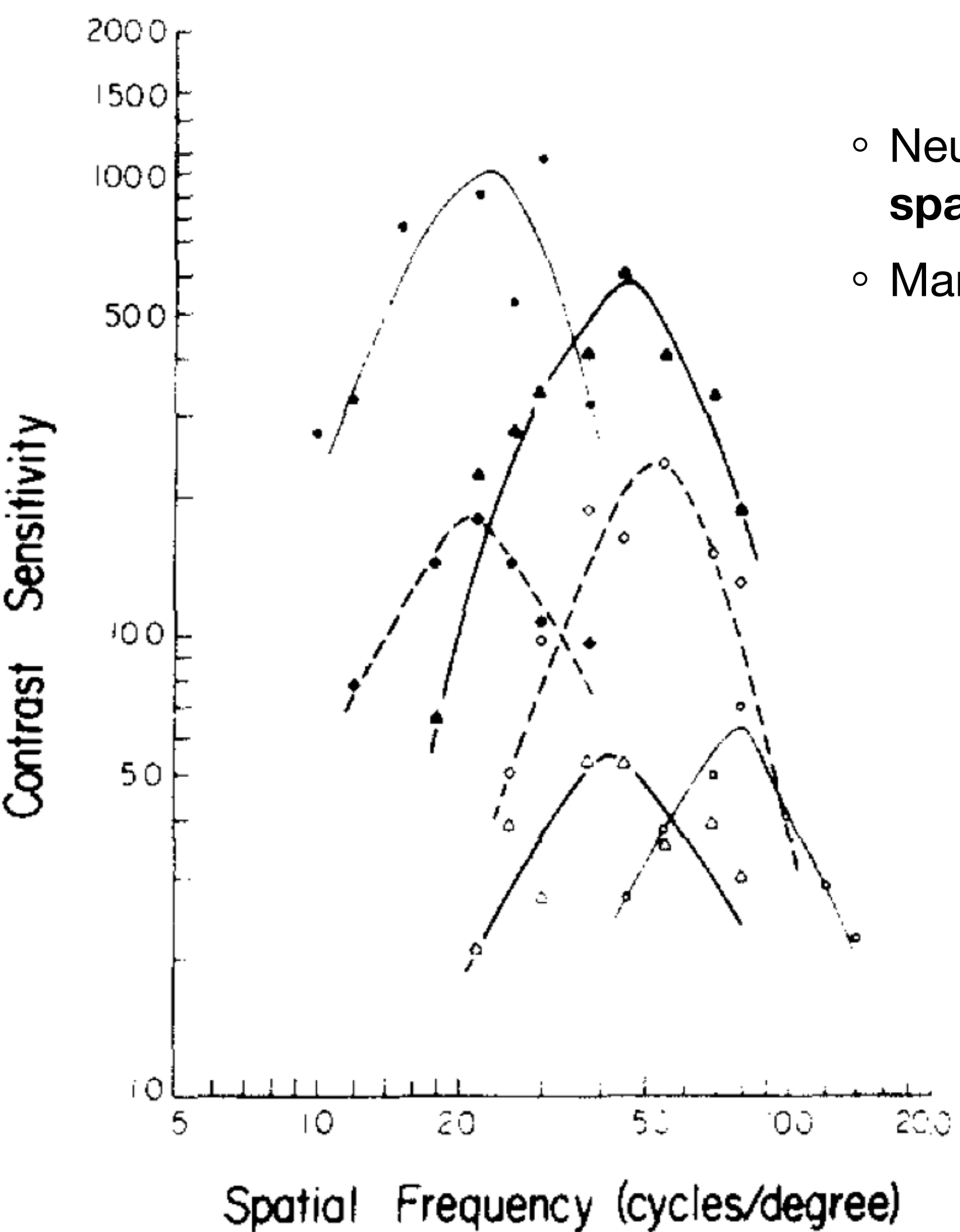
- Neurons span a **wide range of preferred spatial frequencies**
- Many neurons are **narrowly tuned**



Input the same grating stimuli to a trained model

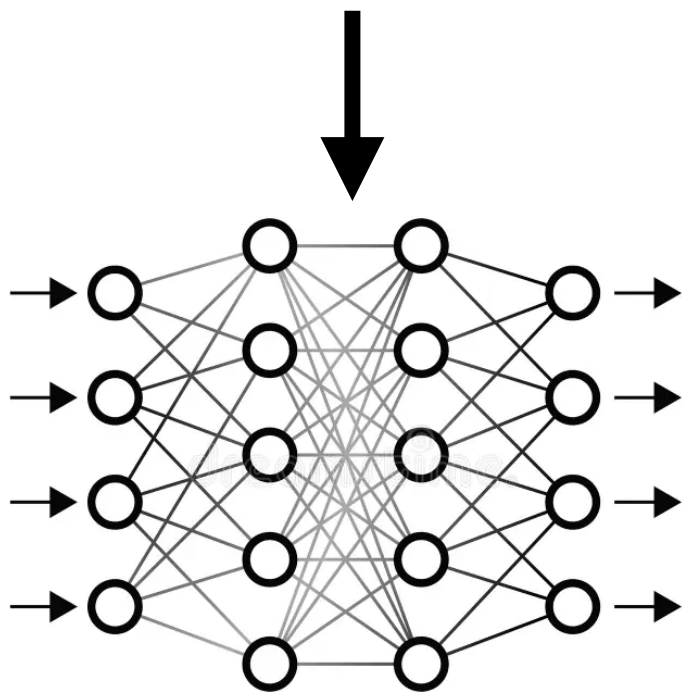
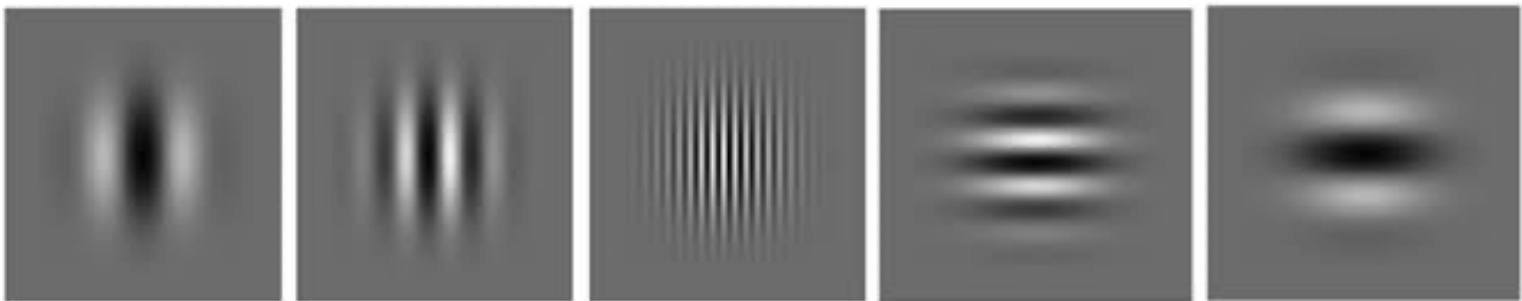


# Comparing results to neural networks

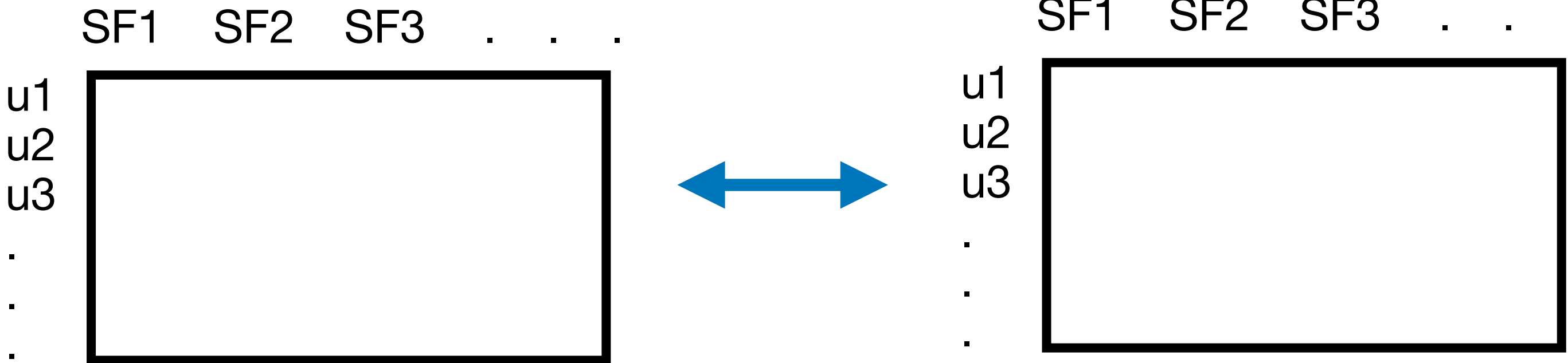


- Neurons span a **wide range of preferred spatial frequencies**
- Many neurons are **narrowly tuned**

Input the same grating stimuli to a trained model



Compare distributions of peak SF in model and neural data



# Brain score platform

## How to use

```
from brainscore_vision import load_benchmark
benchmark = load_benchmark("Marques2020_DeValois1982-peak_sf")
score = benchmark(my_model)
```

[Benchmark API](#)

[Code examples](#)

Data:  
Marques2020\_DeValois1982

[Find on GitHub](#)

## Model scores

### Score Legend

Min Alignment

Max Alignment

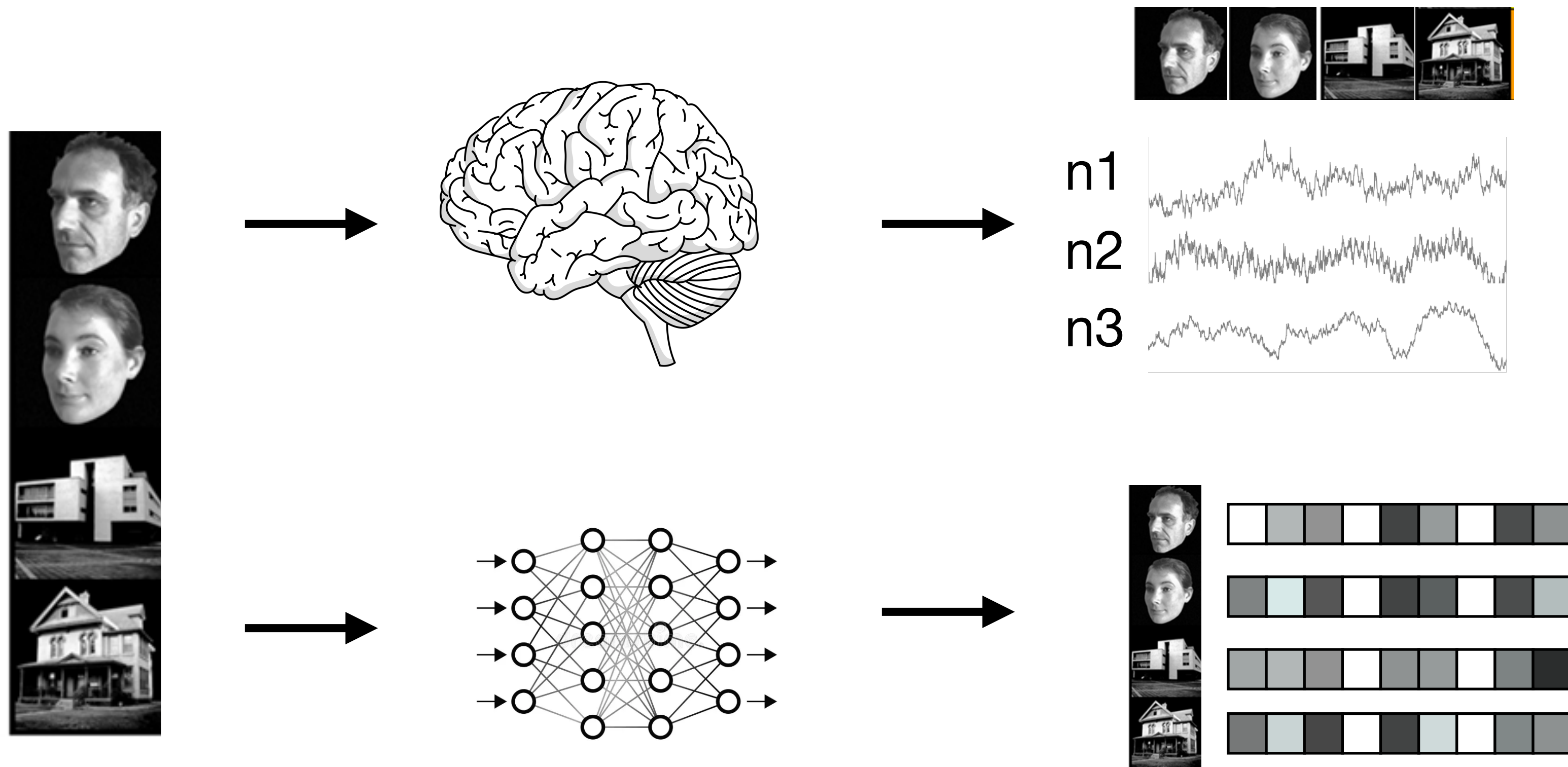
Metric: peak\_sf

[Find on GitHub](#)

Rank	Model	Score
1	<a href="#">resnet-18-LC_w_sh_100_iter_m</a>	.964
2	<a href="#">resnet50_imagenet_10_seed-0</a>	.950
3	<a href="#">alexnet_training_seed_01</a>	.943
4	<a href="#">resnet-18-LC_m</a>	.941
5	<a href="#">resnet50_linf_4_robust</a>	.935
6	<a href="#">alexnet_training_seed_08</a>	.933

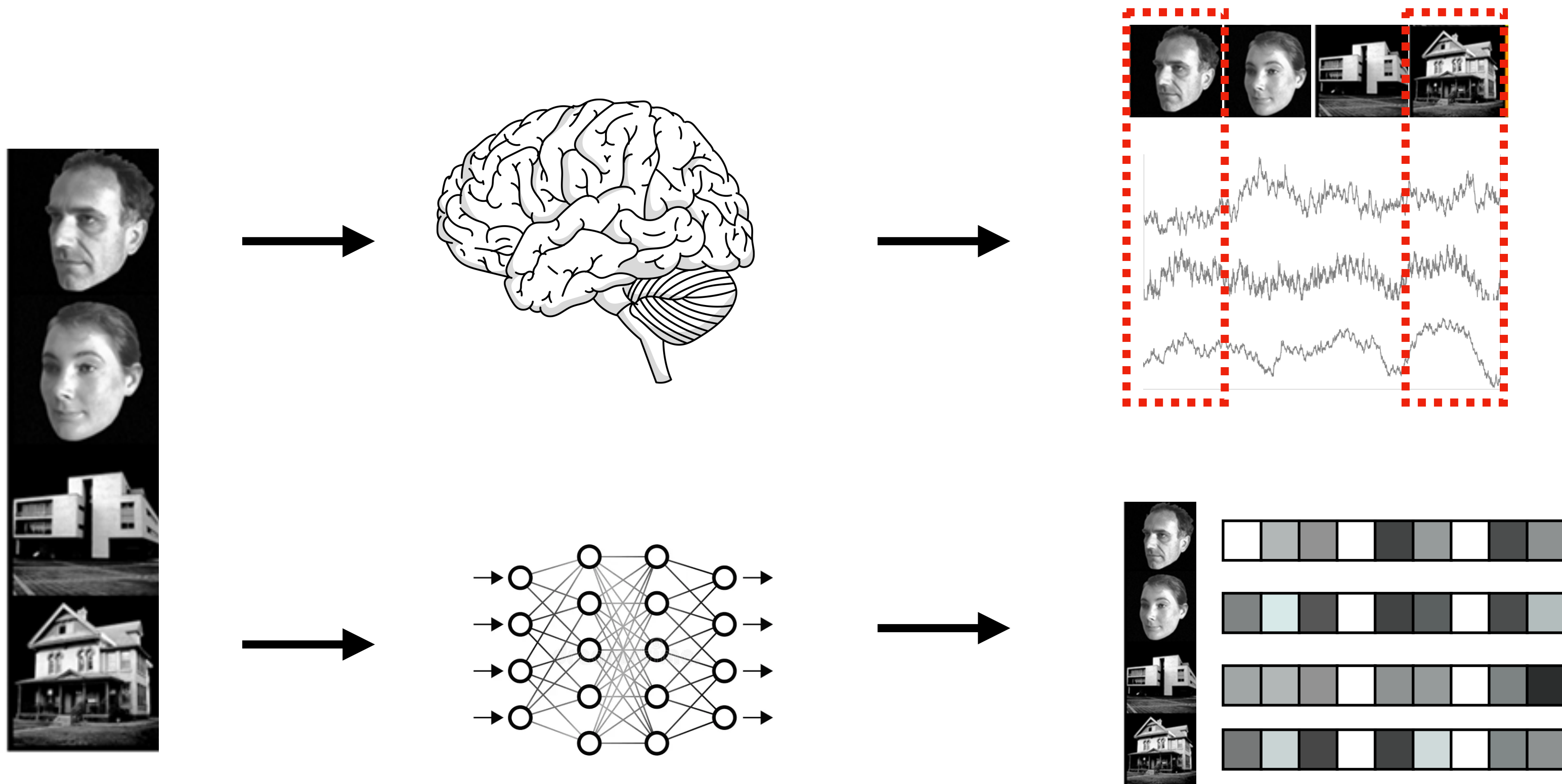
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



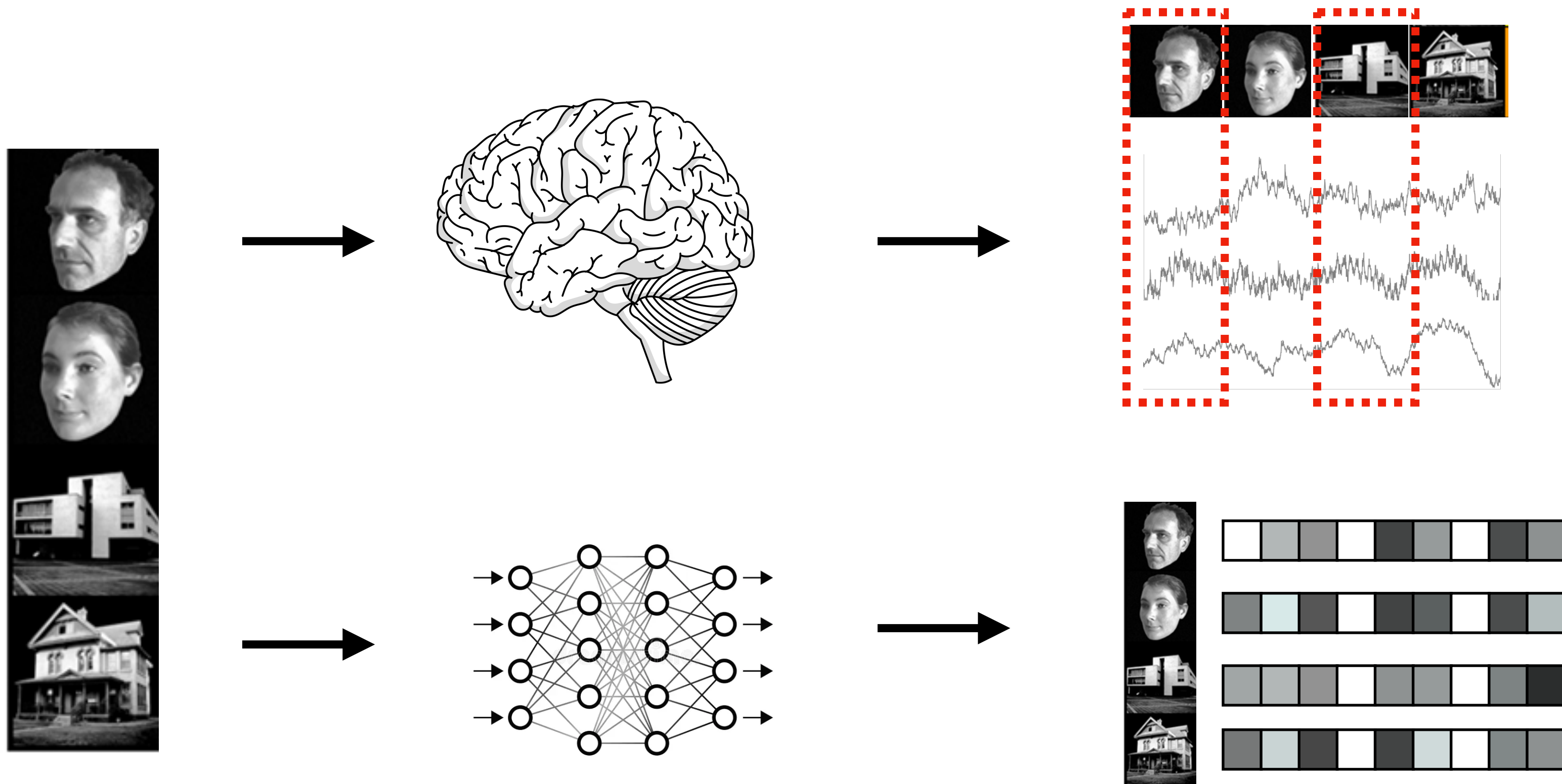
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



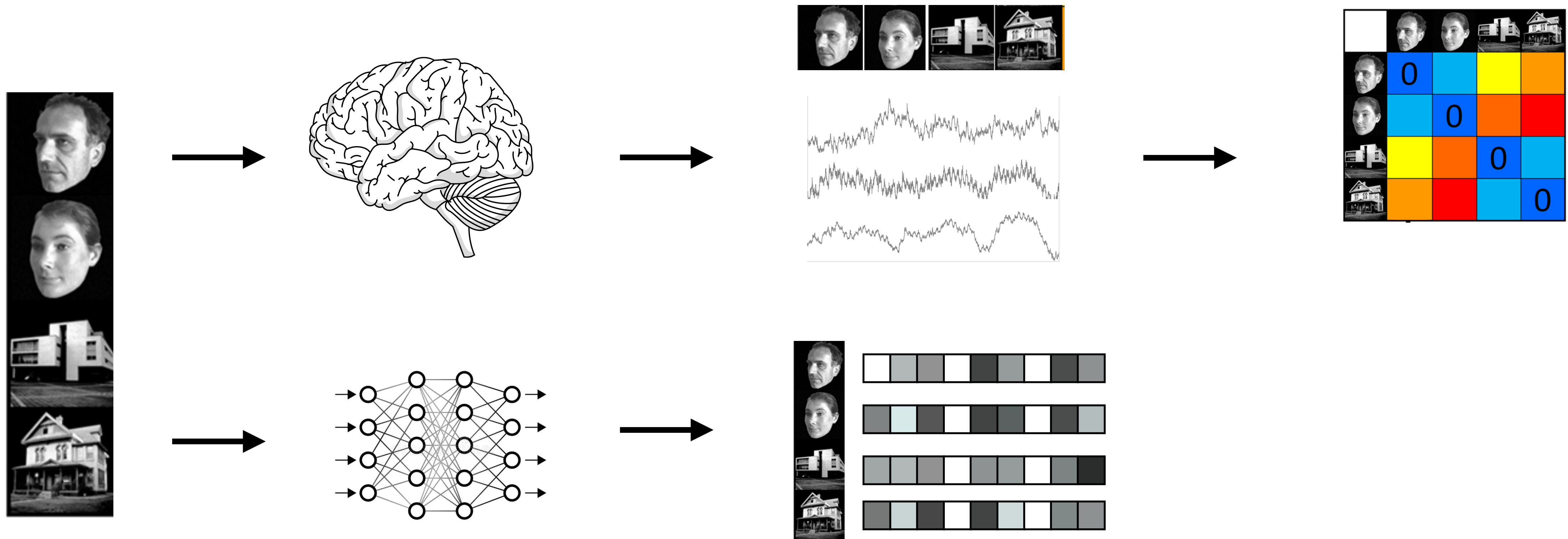
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



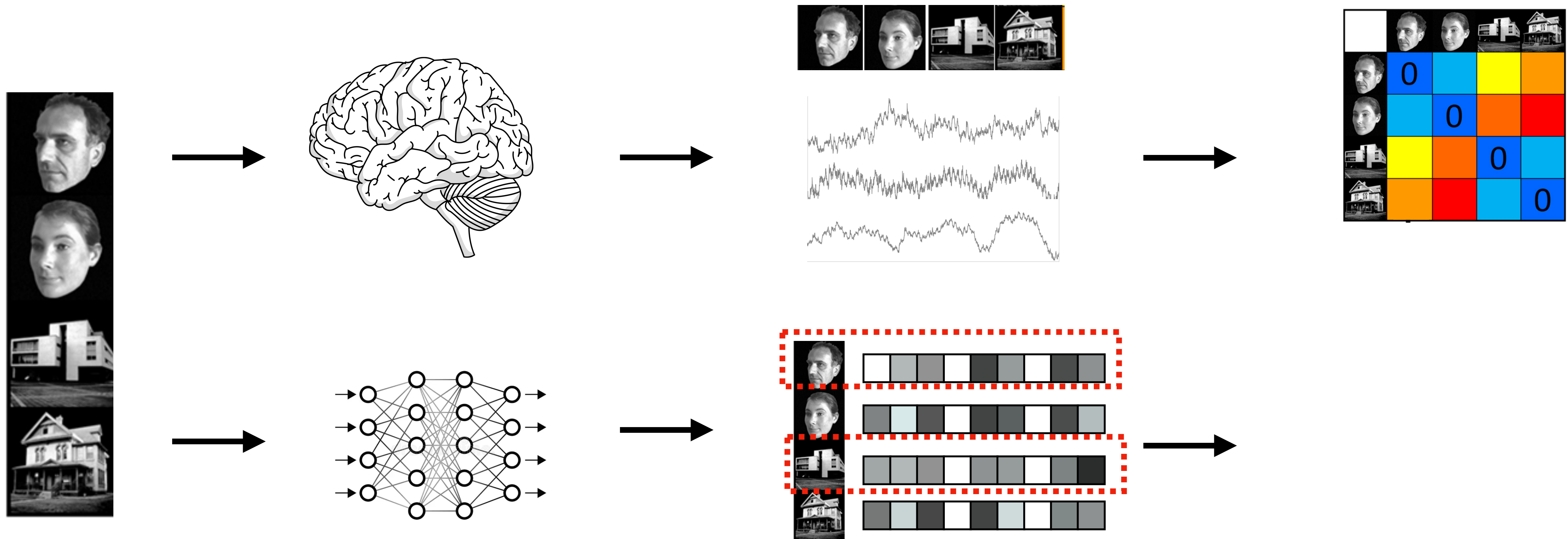
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



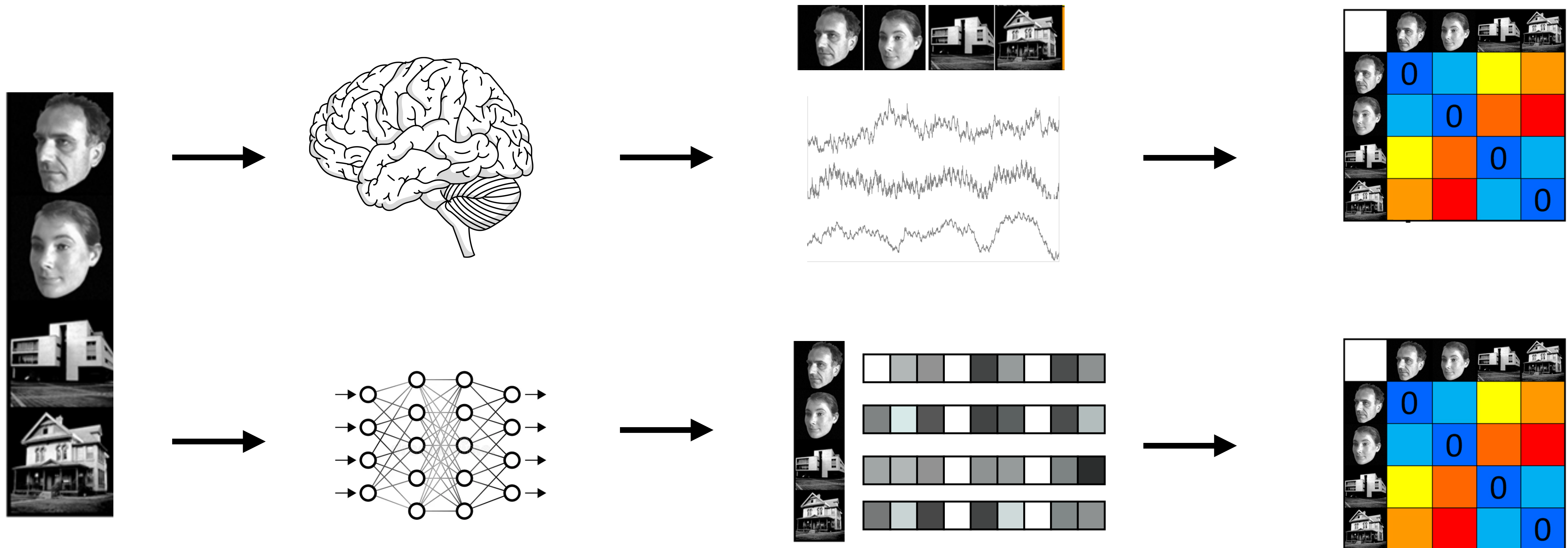
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



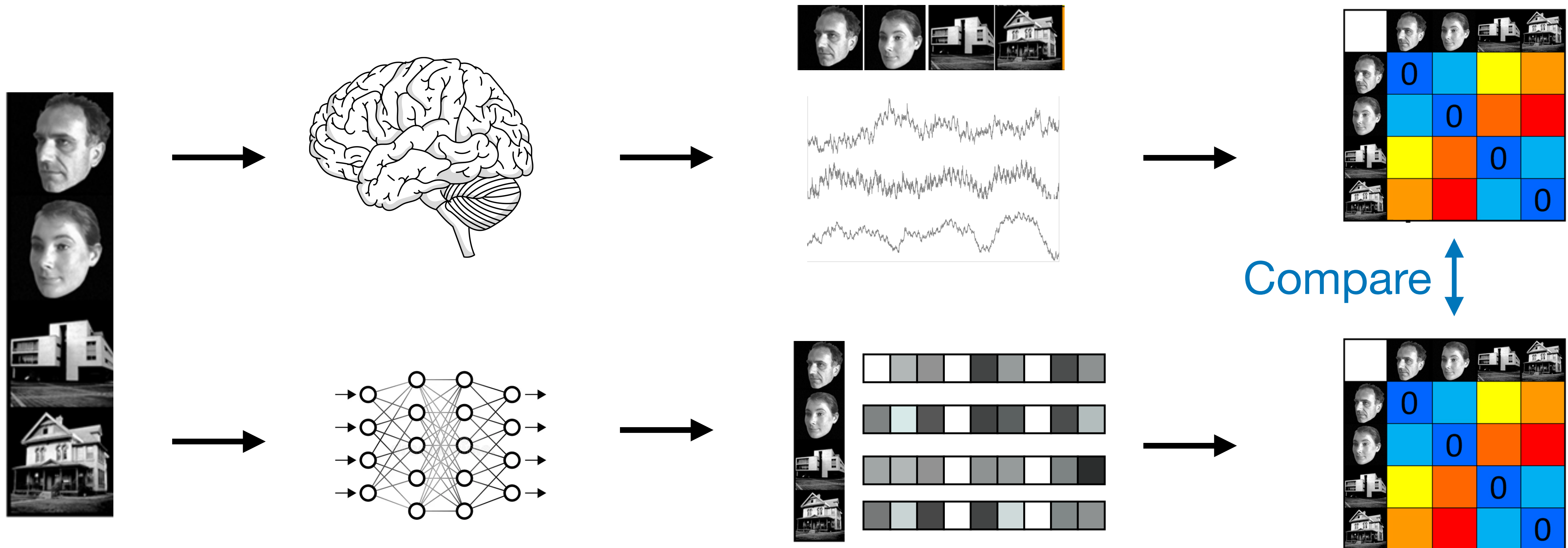
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



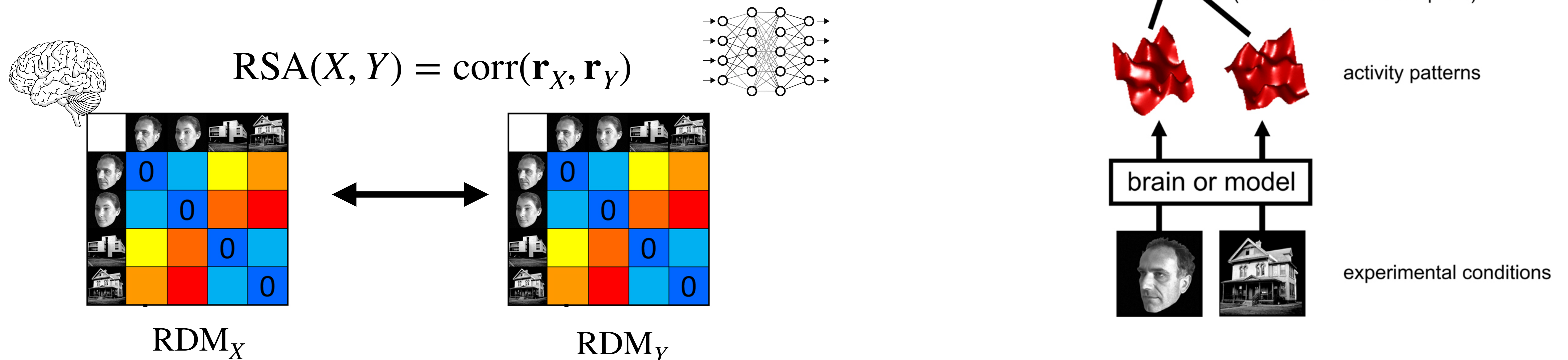
## 2. Using stimulus-by-stimulus similarity matrices

- ❖ Compare representations via stimulus–stimulus relationships
- ❖ Ignore neuron-to-neuron correspondence entirely



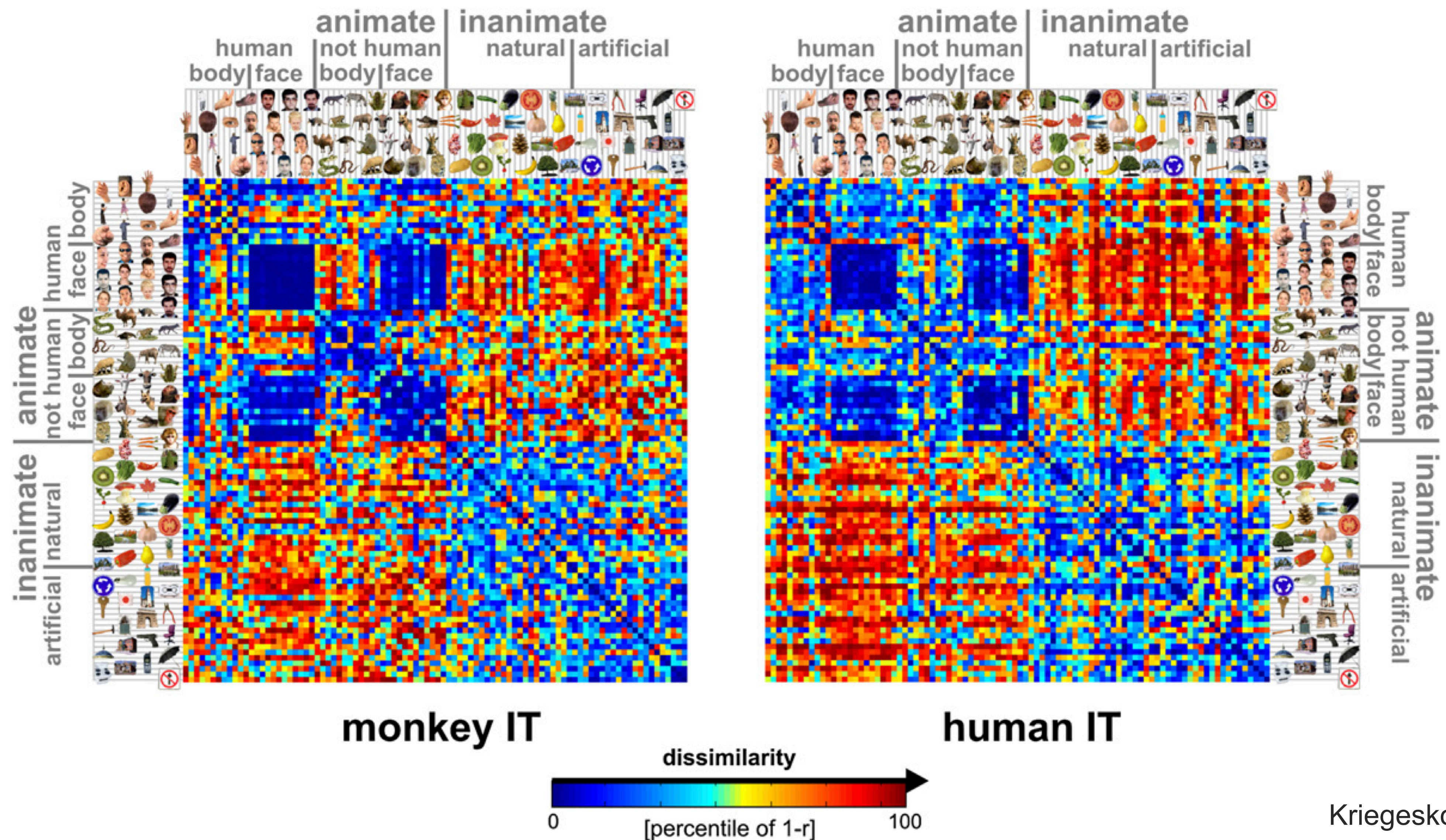
# Example: Representational Similarity Analysis (RSA)

- ❖ “Do two systems organize stimuli in the same geometric way?”
- ❖ **Pros:** For the first time allowed comparison between any systems as long as the stimuli was the same. Also doesn't need any training params.
- ❖ **Cons:** Very similar systems (up to a linear transform) can look very different under RSA



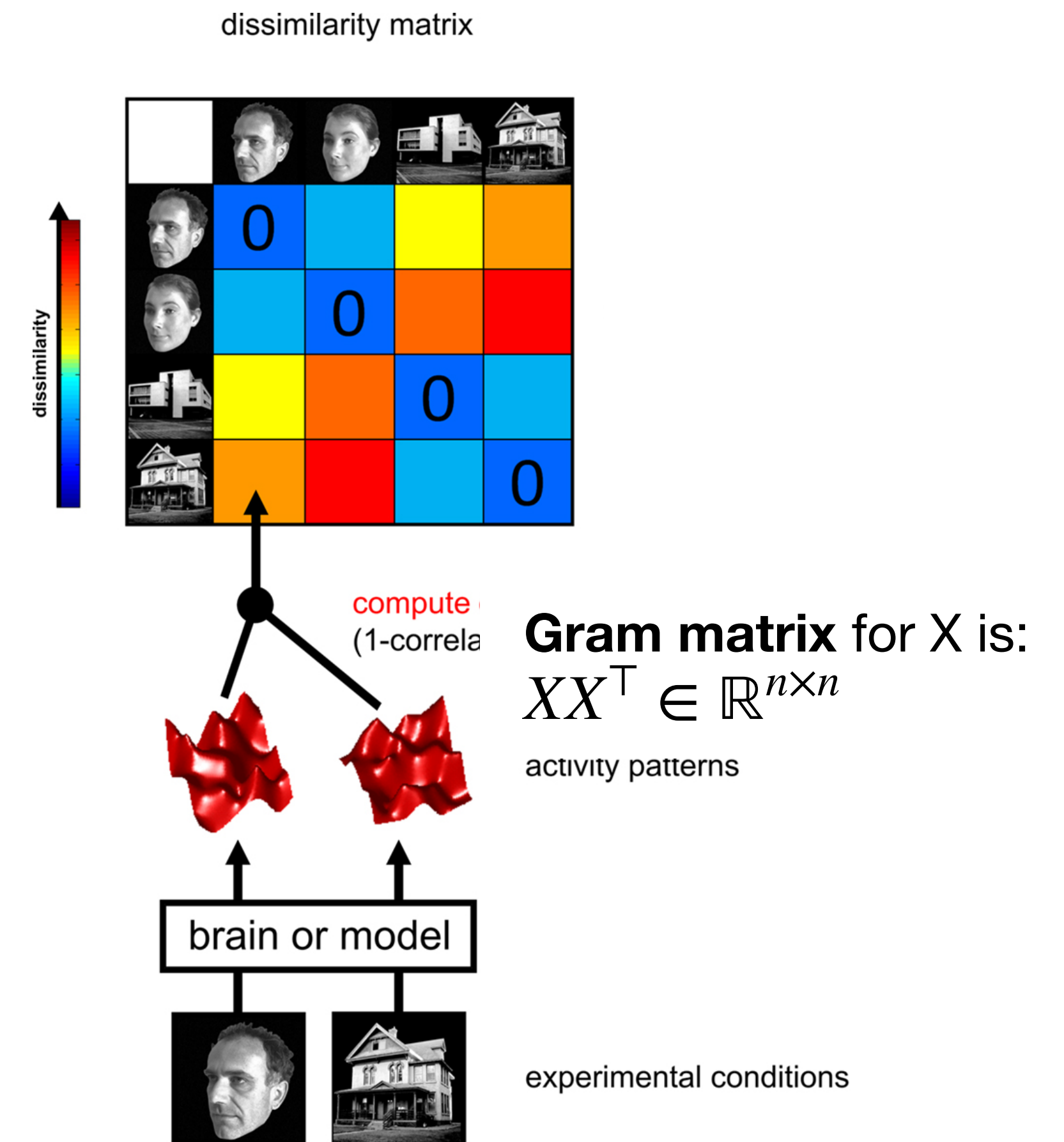
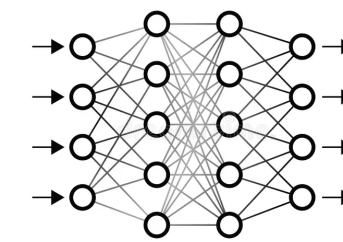
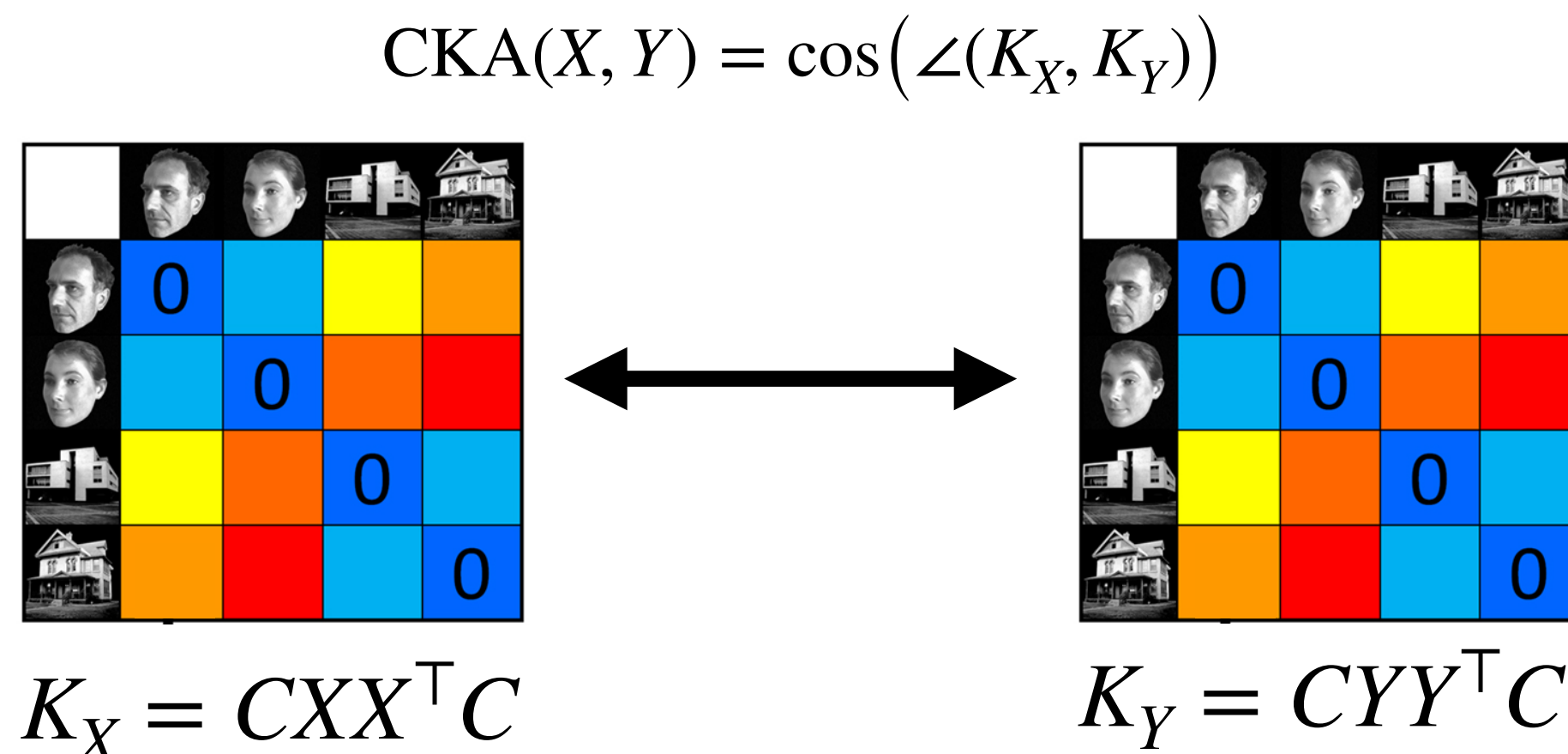
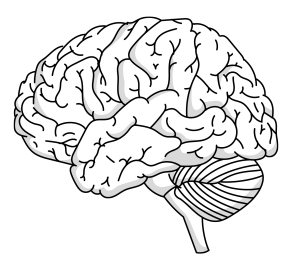
# Example: RSA

$r=0.49$



# Example: Centered Kernel Alignment (CKA)

- ❖ Similar method to RSA, but operates on **similarities** rather than **distance**
- ❖ Computes cosine similarity between centered gram matrices
- ❖ **Pros:** more flexible than RSA, invariant to rotation, scaling
- ❖ **Cons:** Similar systems (up to a linear transform) can look different under CKA.

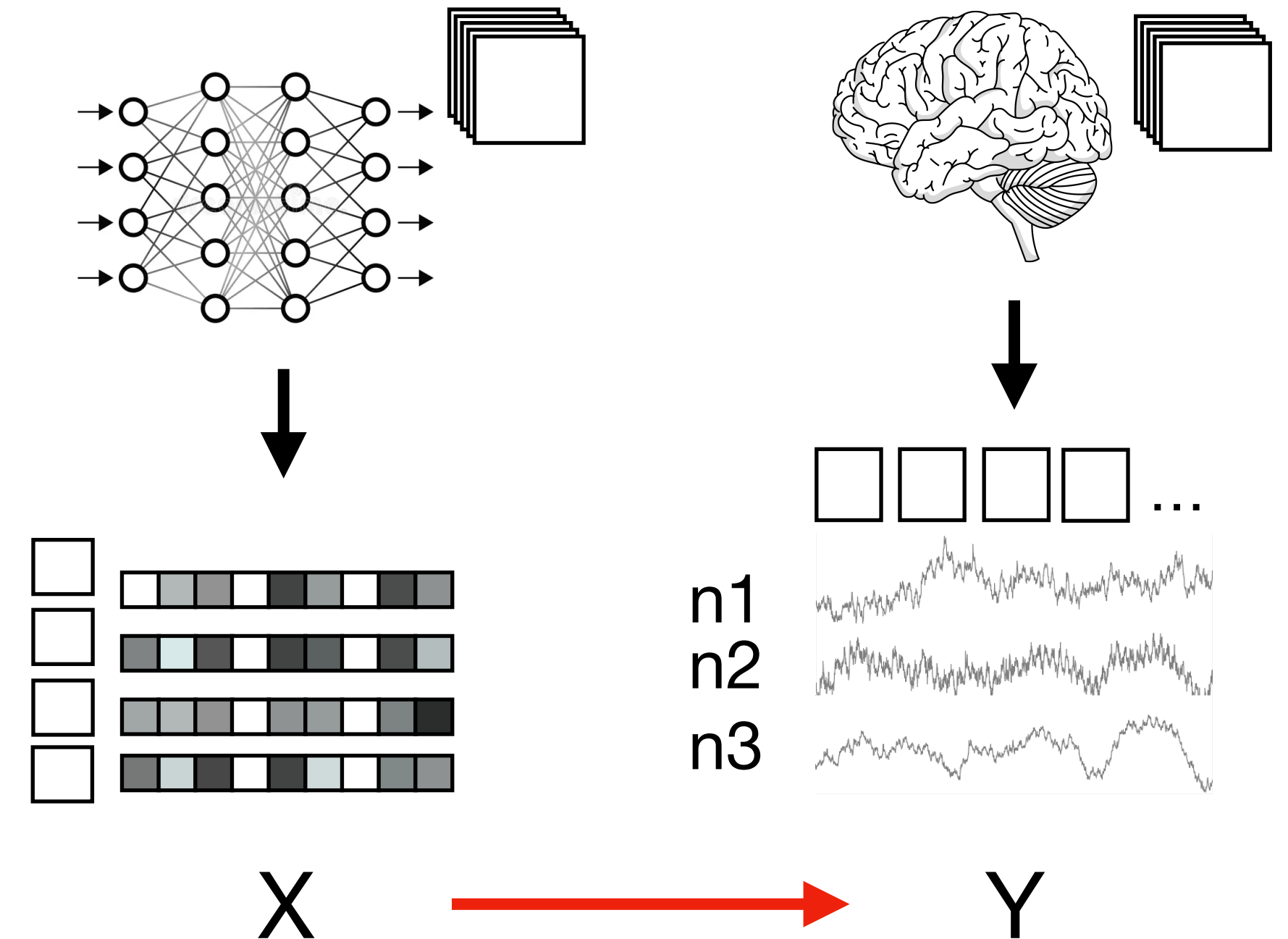


# 3. Learning mappings from the model to the brain

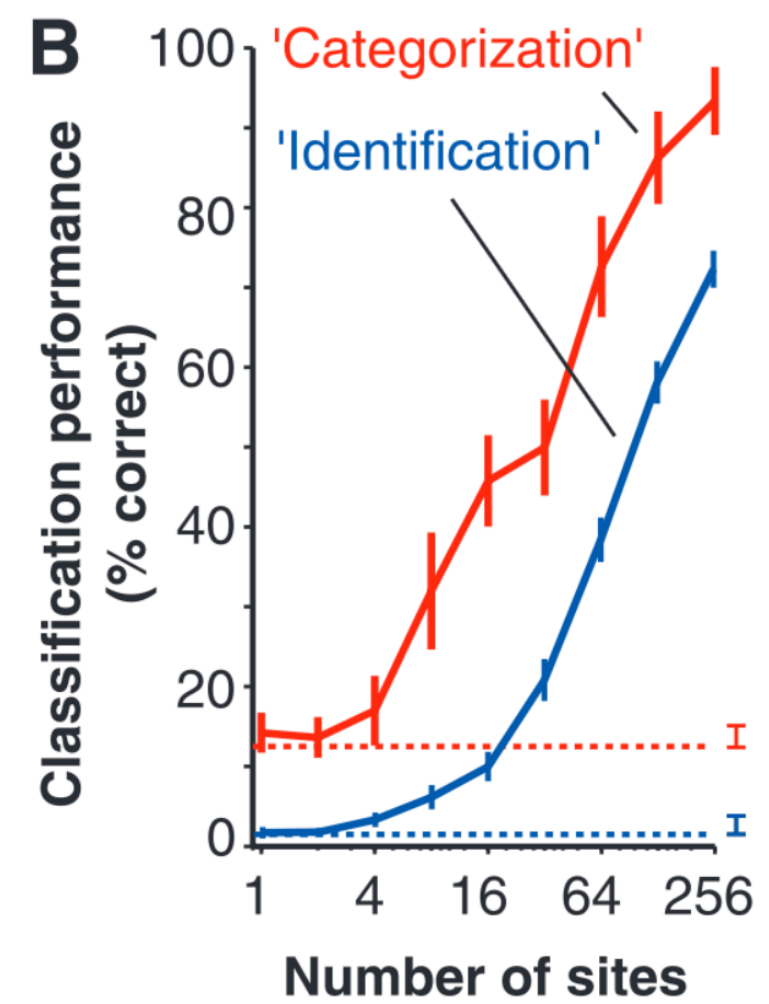
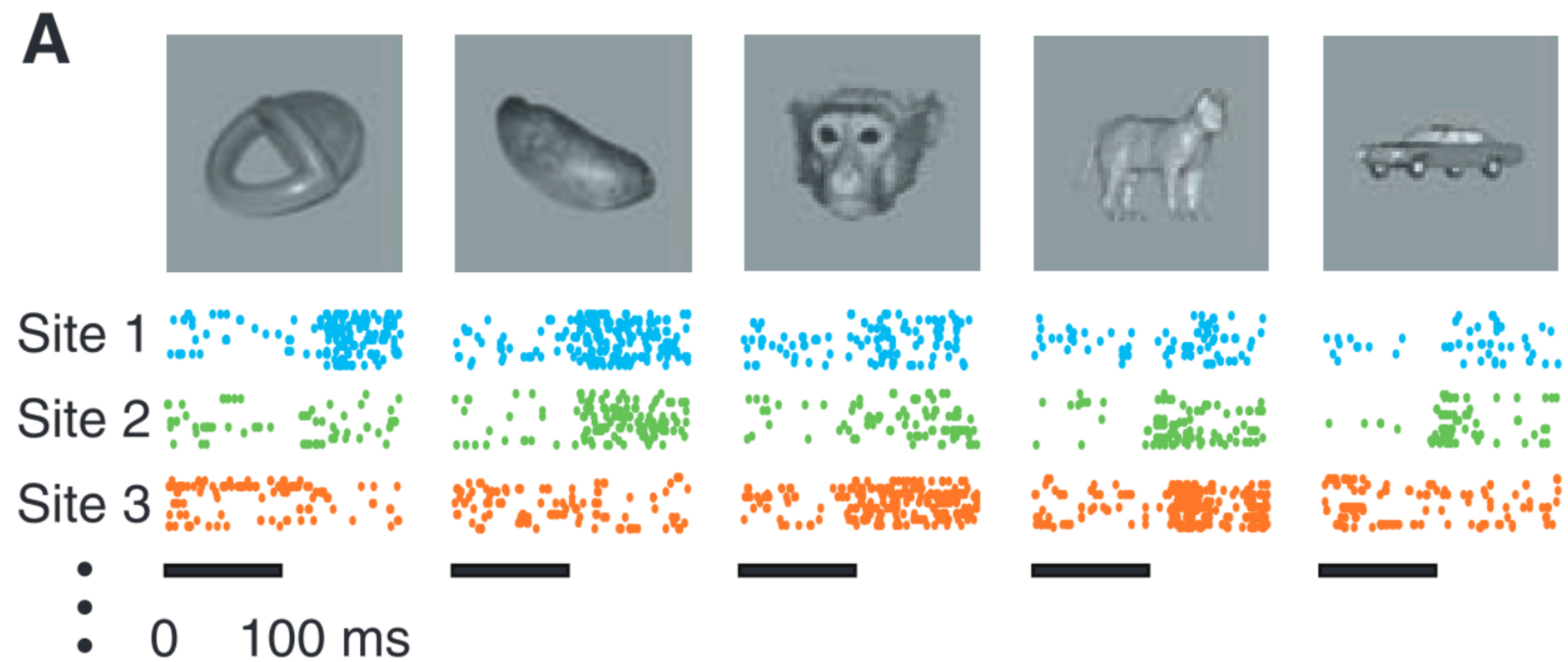
$$X \in \mathbb{R}^{n \times N_x} \text{ , } Y \in \mathbb{R}^{n \times N_y}$$

- Most methods focus on learning a linear mapping
- Let  $X$  be the model representations, and  $Y$  the neural responses to the same set of stimuli.
- **Goal:** Find the best mapping from  $X$  to  $Y$ .

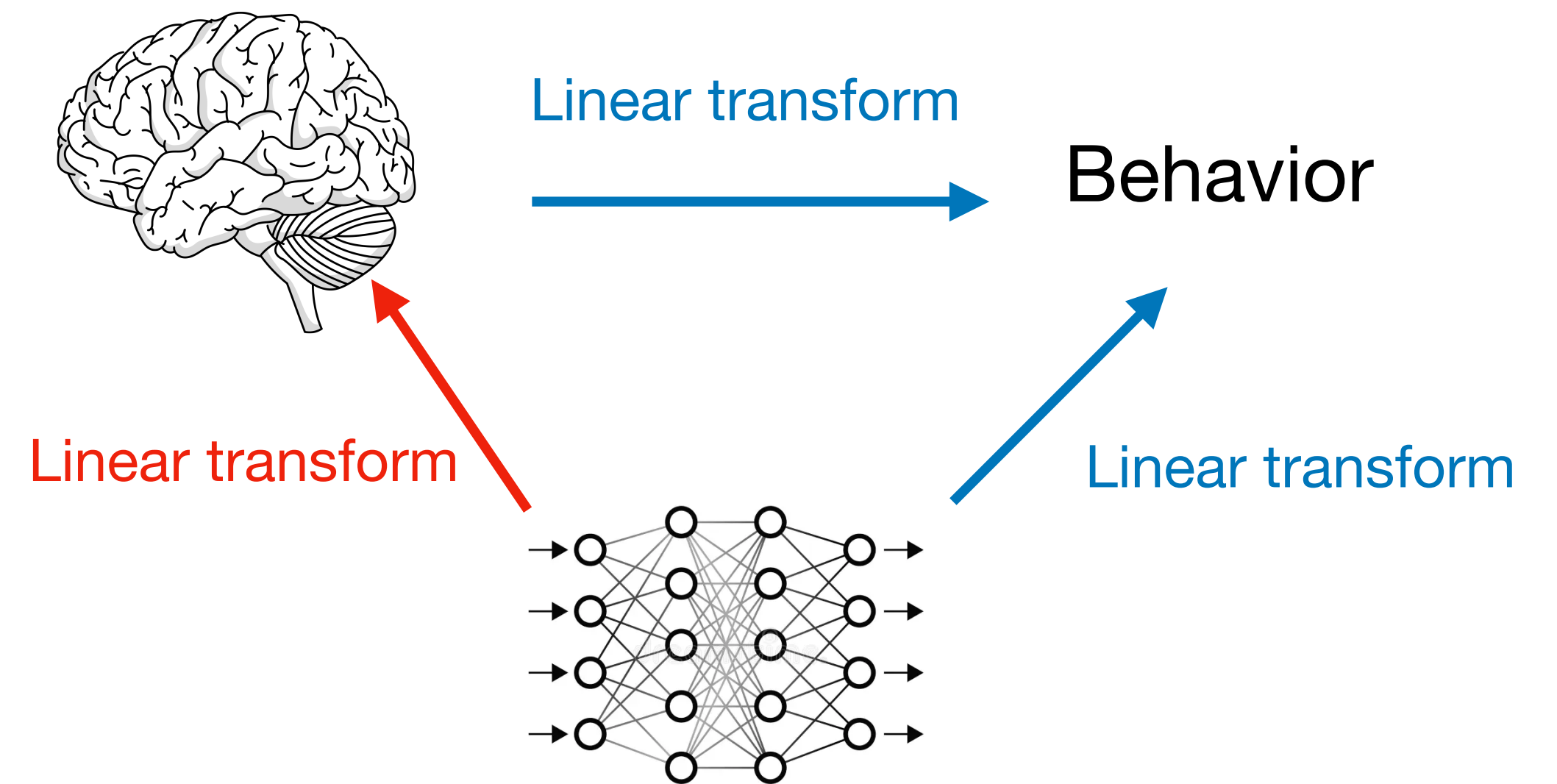
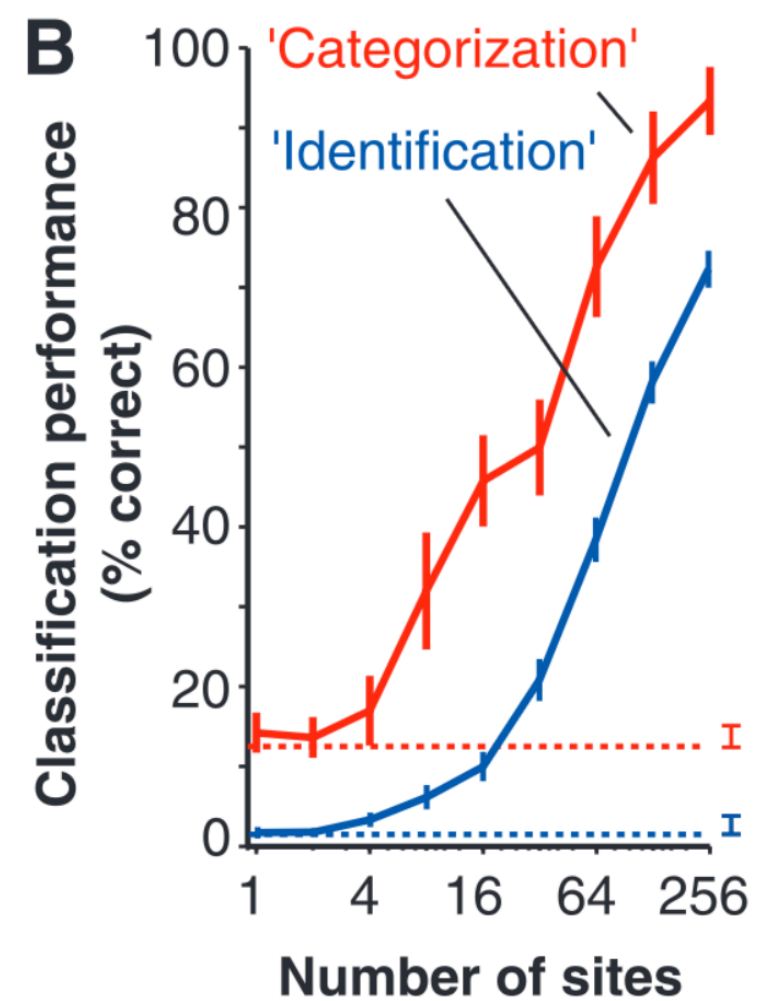
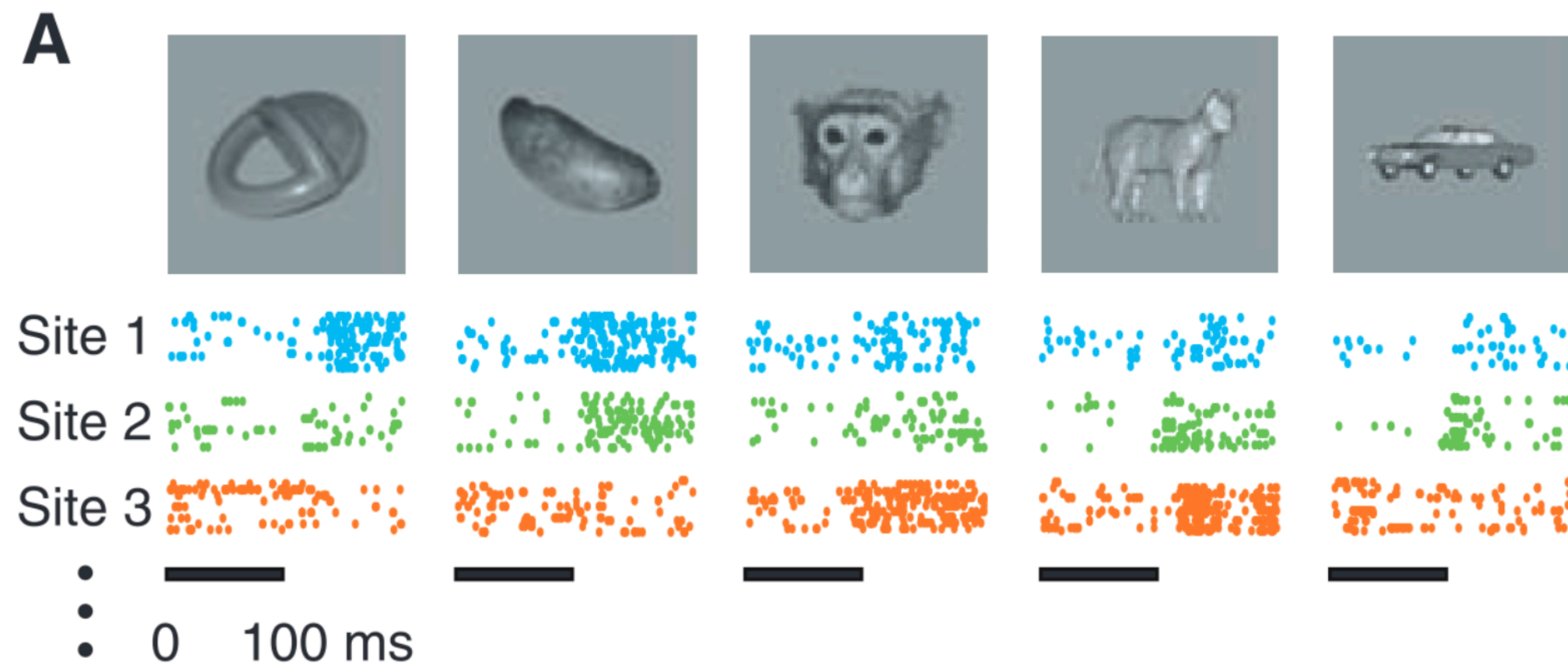
Rows = stimuli,      Columns = neurons / features



# Why Linear Mapping?

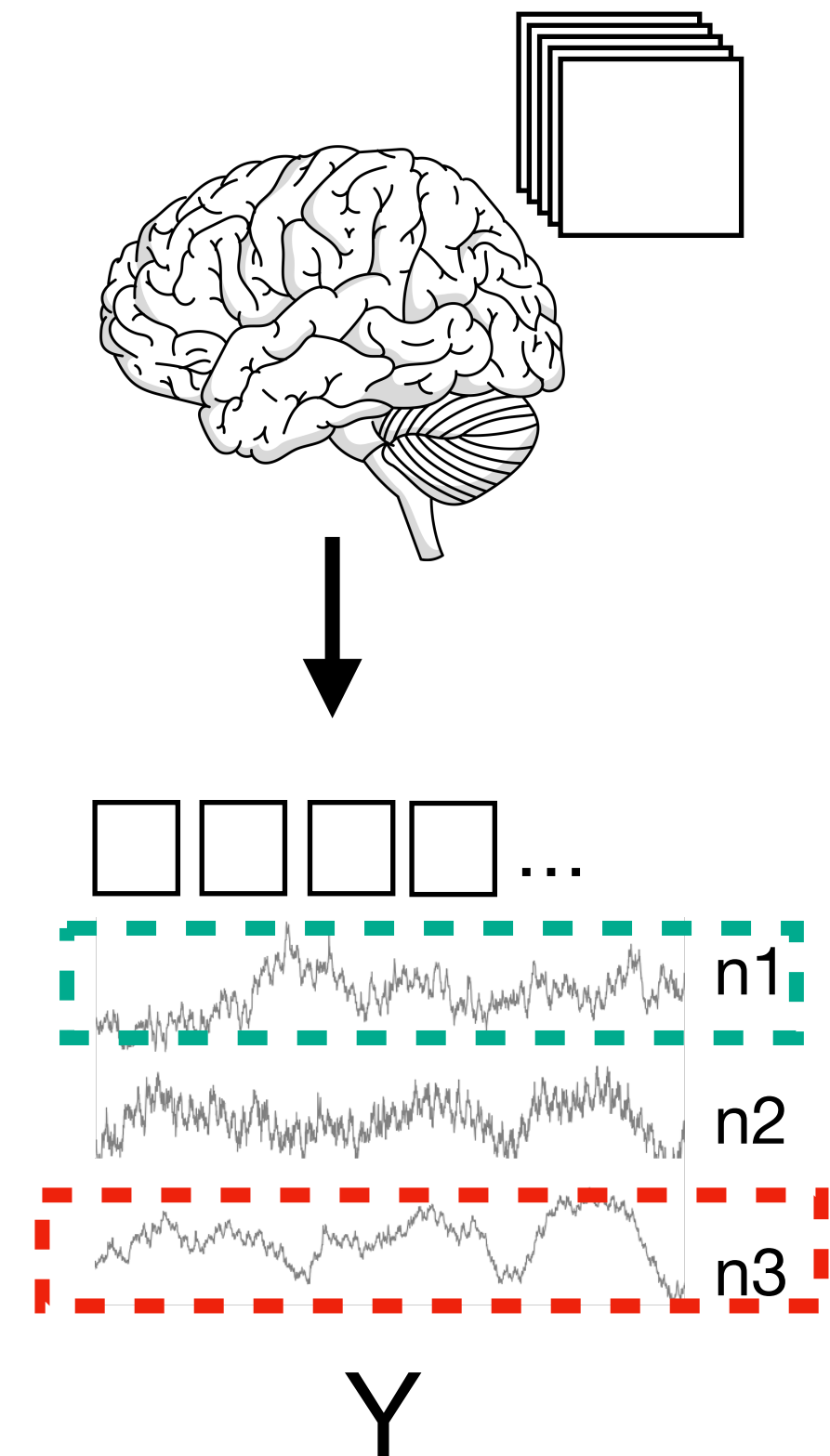
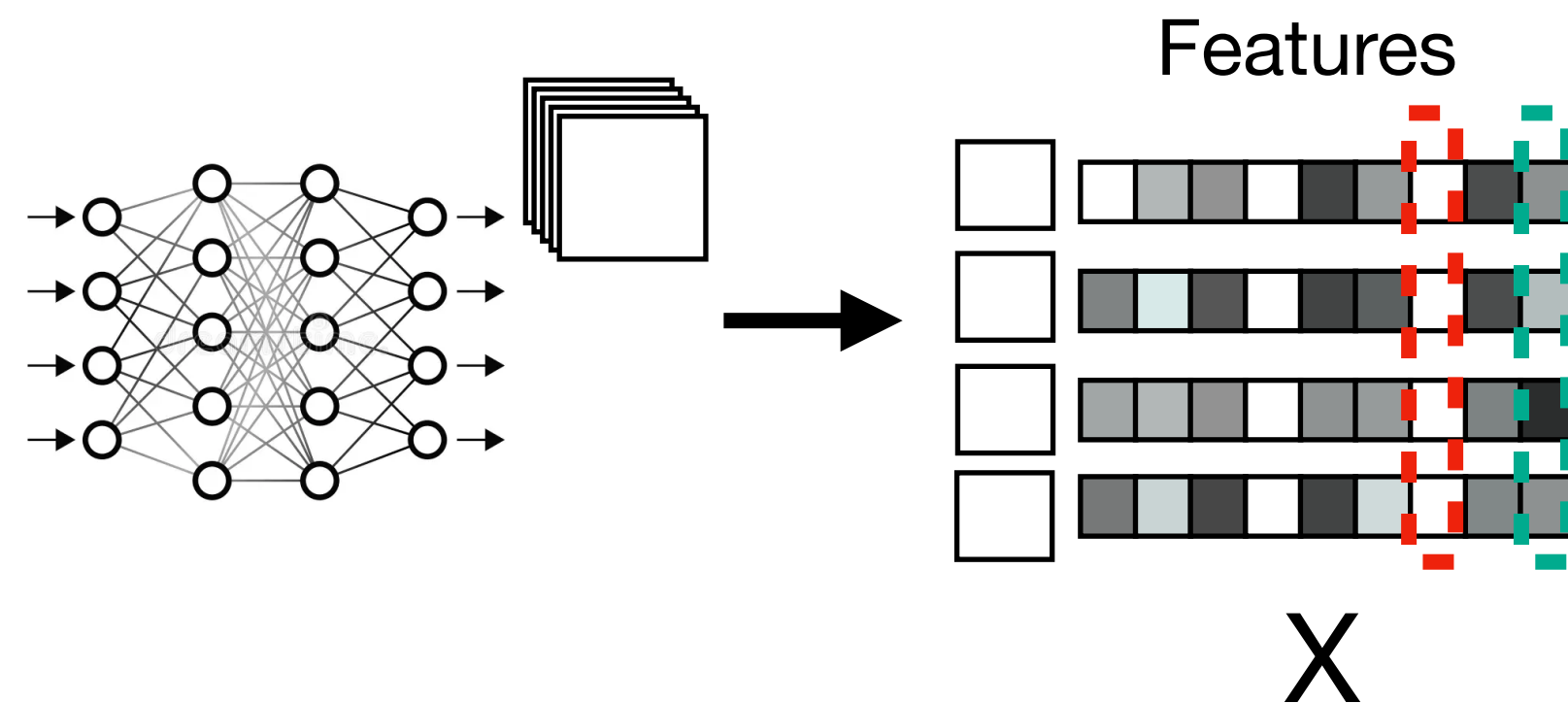


# Why Linear Mapping?



# Example: One to one mapping

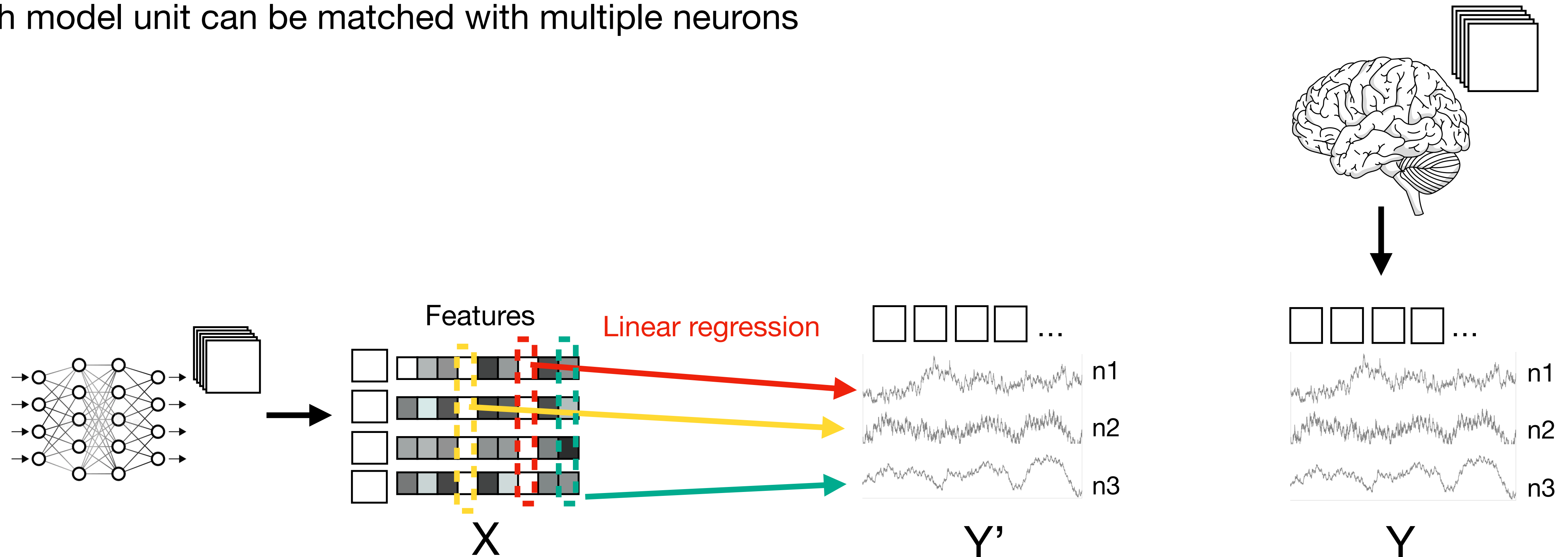
**Idea:** for 2 systems to be similar their parts should be similar



# Example: One to one mapping

**Idea:** for 2 systems to be similar their parts should be similar

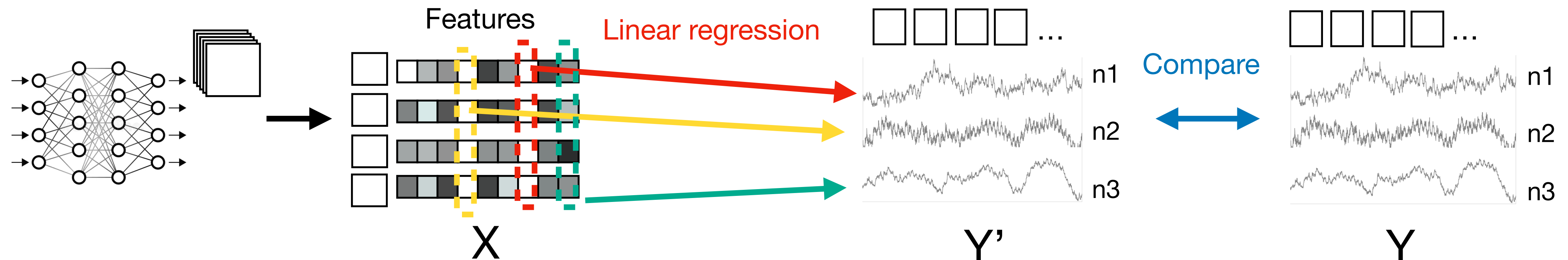
- ❖ For each neuron, find the model unit with the highest correlation, then compute the optimal linear mapping from neuron to model unit
- ❖ Each model unit can be matched with multiple neurons



# Example: One to one mapping

**Idea:** for 2 systems to be similar their parts should be similar

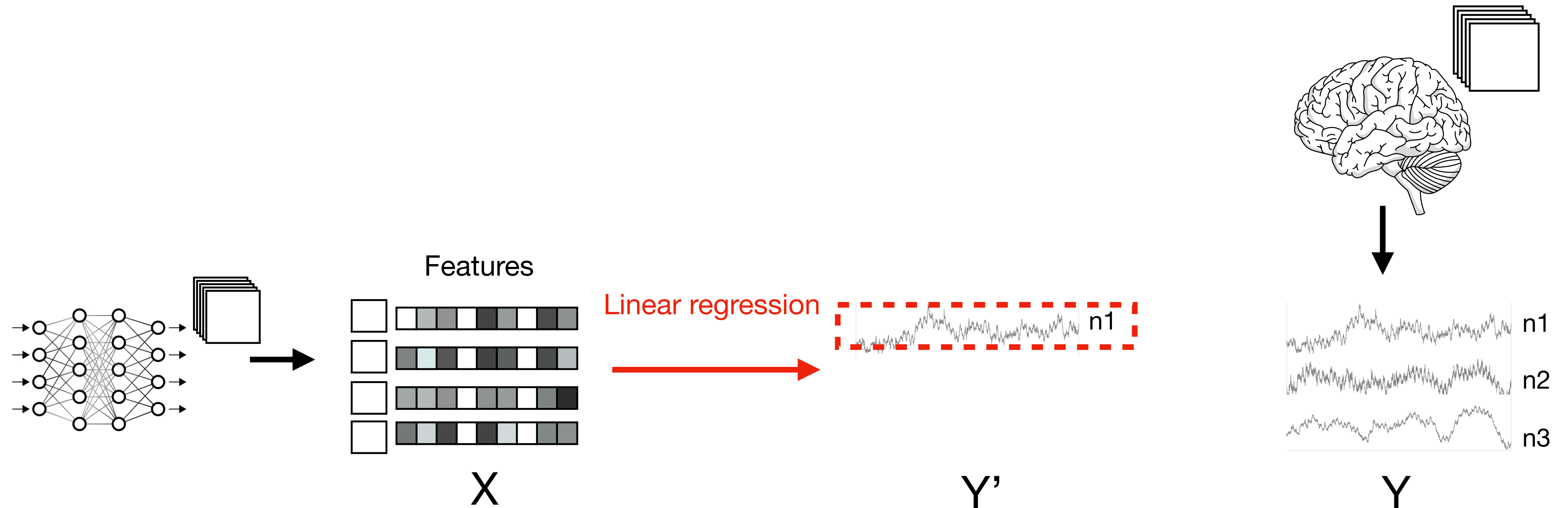
- ❖ For each neuron, find the model unit with the highest correlation, then compute the optimal linear mapping from neuron to model unit
- ❖ Each model unit can be matched with multiple neurons
- ❖ **Pros:** Simple and strict, effective for comparing very similar regions where parts of the system are consistent across individuals (ex: retina)
- ❖ **Cons:** Most brain areas don't have the exact same units in different subjects (ex: IT)



# Example: Linear Regression

**Idea:** Find linear combinations of model units that together produce a 'synthetic neuron'

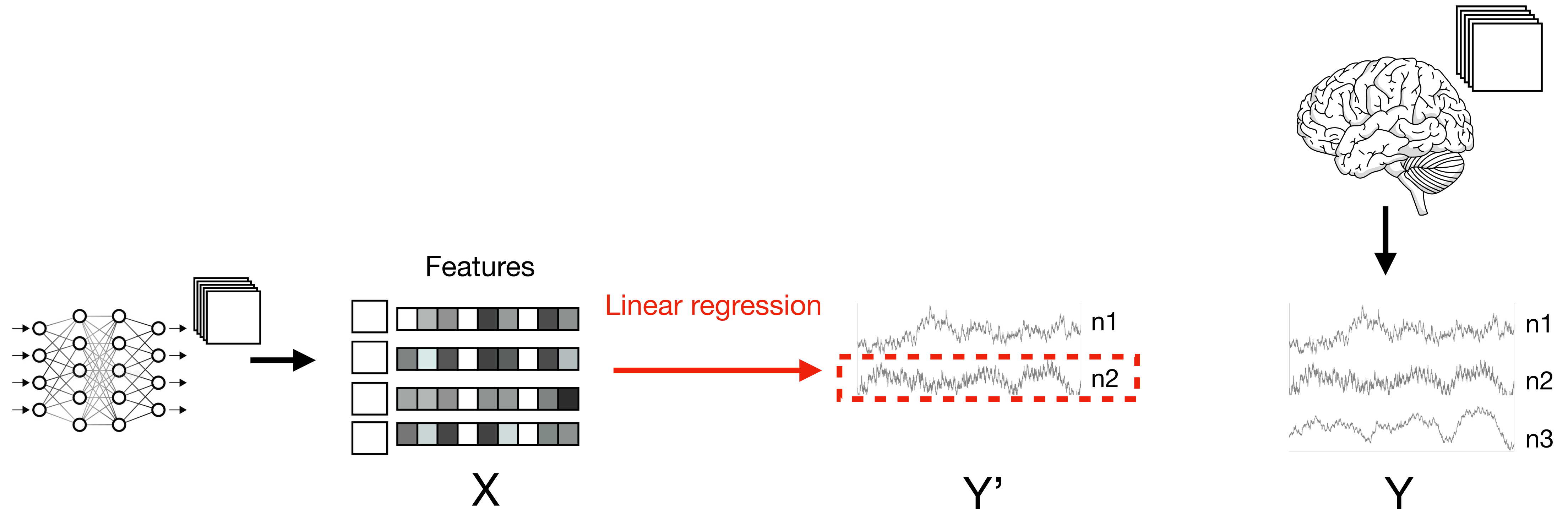
- ❖ Learn a mapping from all model units to each target neuron.



# Example: Linear Regression

**Idea:** Find linear combinations of model units that together produce a 'synthetic neuron'

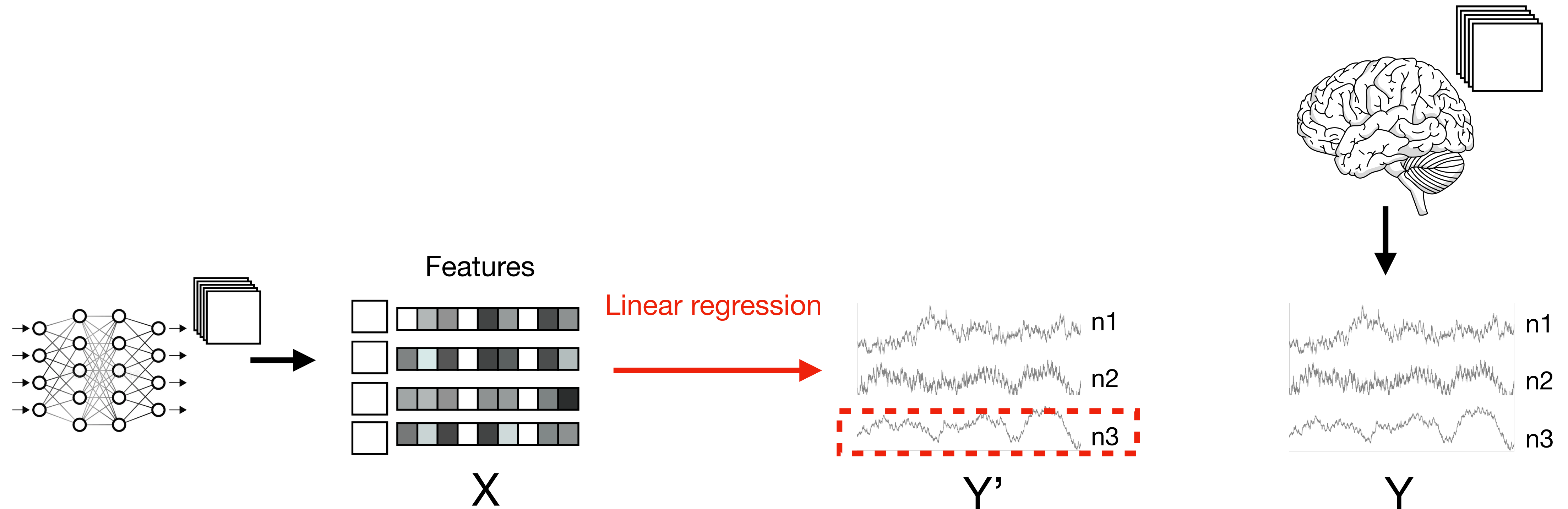
- ❖ Learn a mapping from all model units to each target neuron.



# Example: Linear Regression

**Idea:** Find linear combinations of model units that together produce a 'synthetic neuron'

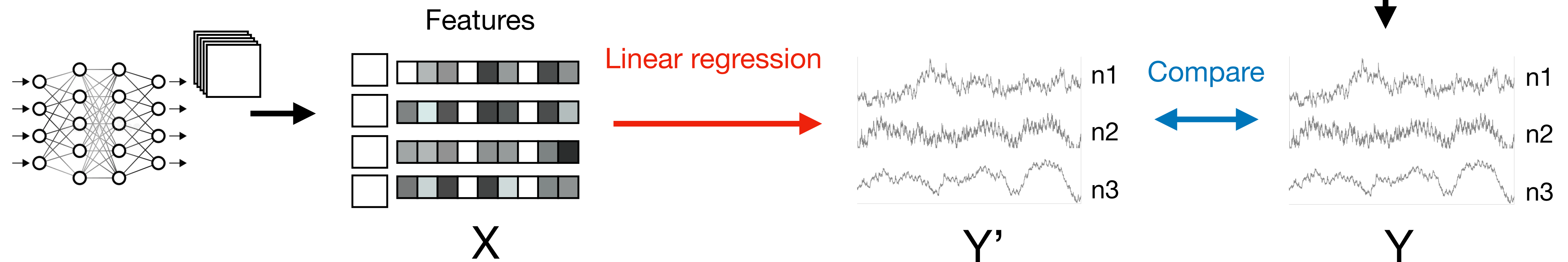
- ❖ Learn a mapping from all model units to each target neuron.



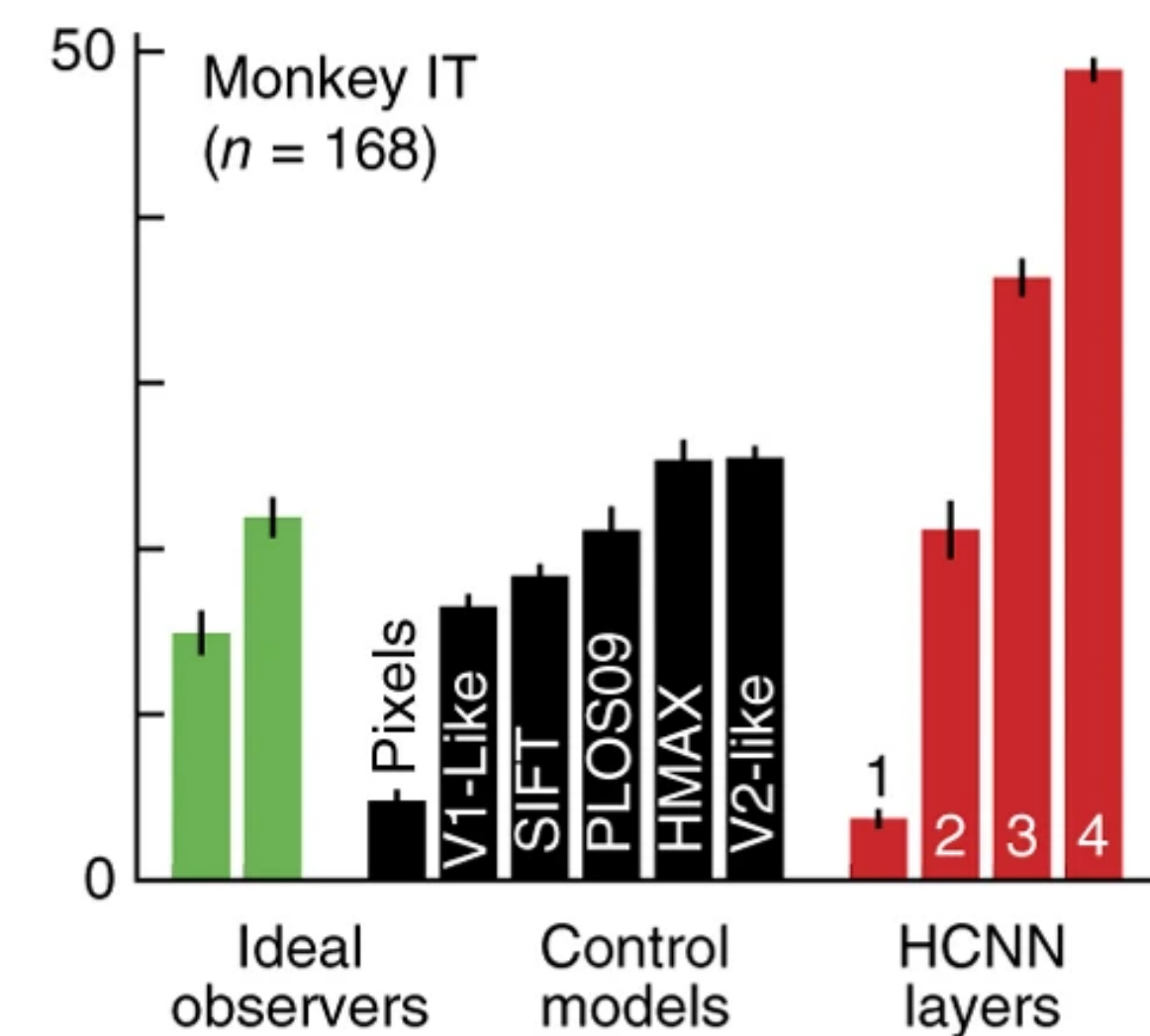
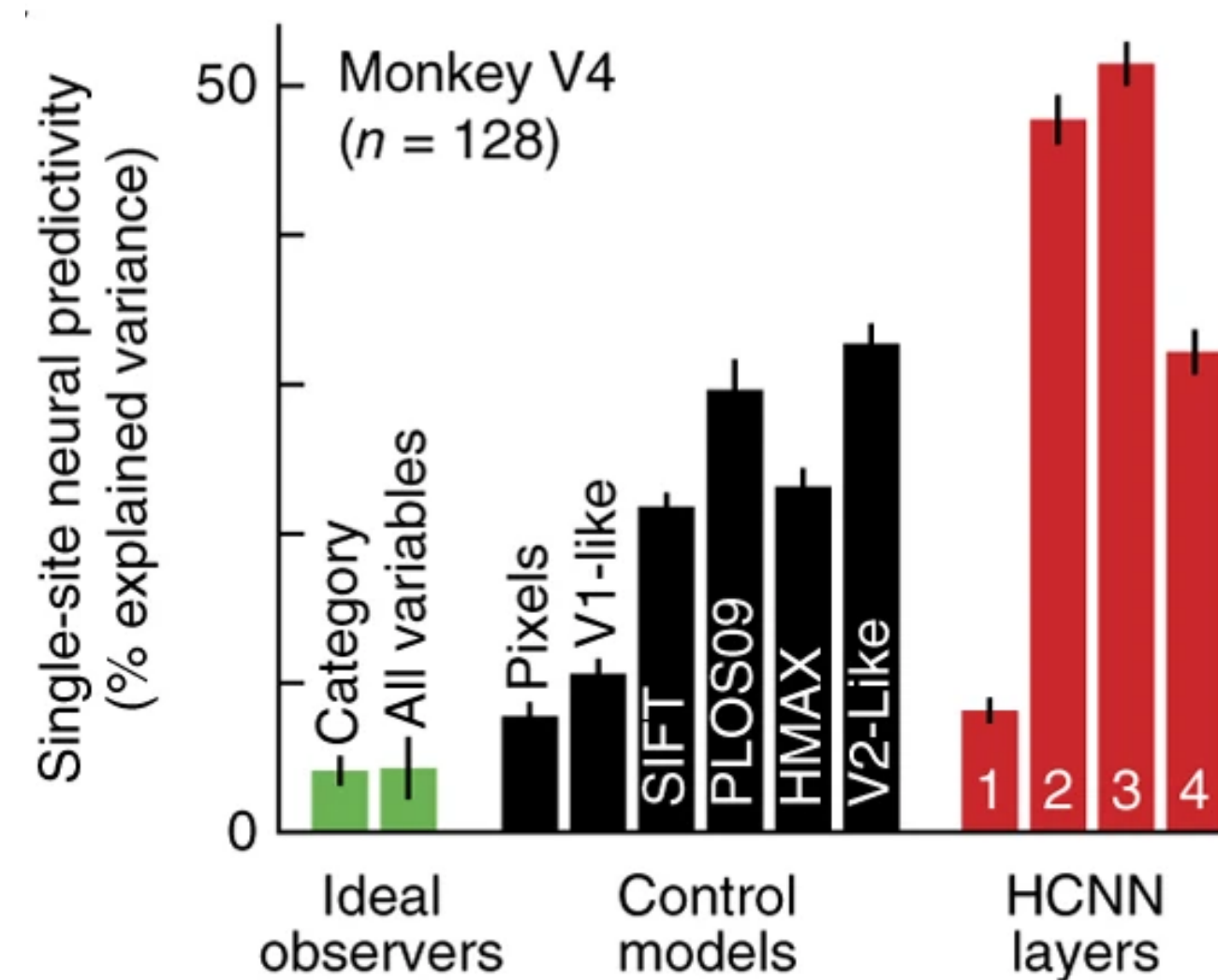
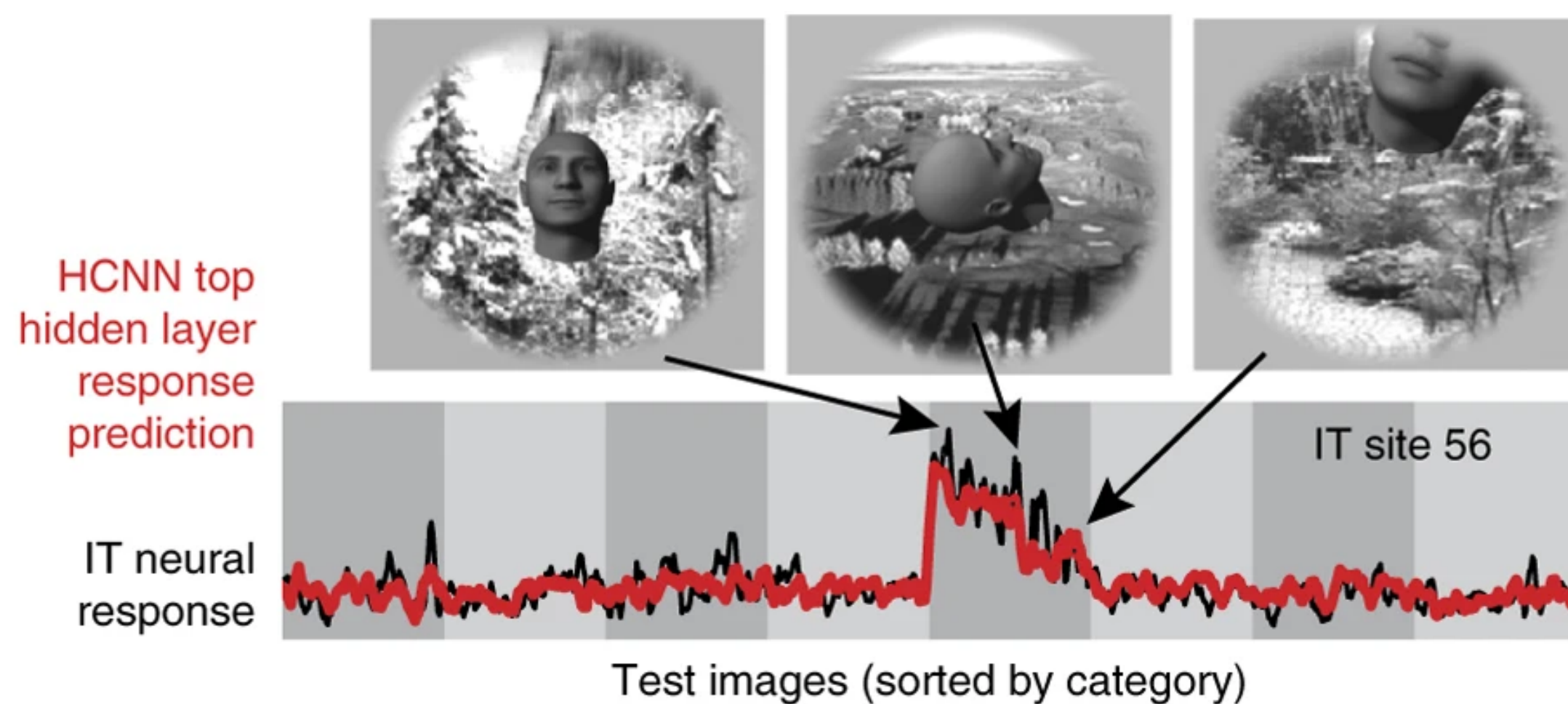
# Example: Linear Regression

**Idea:** Find linear combinations of model units that together produce a 'synthetic neuron'

- ❖ Learn a mapping from all model units to each target neuron.
- ❖ **Pros:** More flexible than RSA & one to one matching, not prone to errors when systems are similar
- ❖ **Cons:** Need to train parameters



# Example: Linear Regression

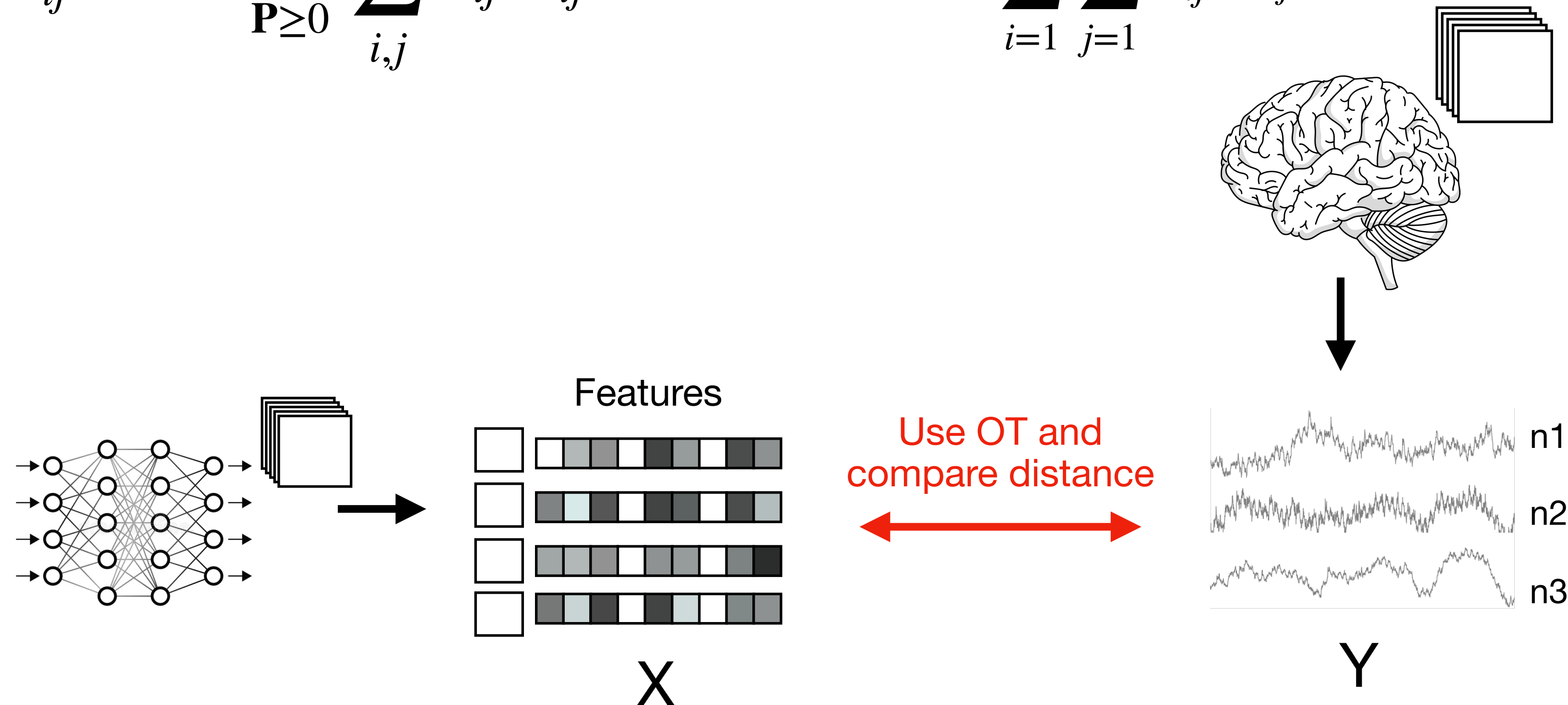


# Example: Soft Matching

**Idea:** Match individual model units to individual neurons without requiring an exact one to one match

❖ Solve an Optimal Transport (OT) problem: “how much is a source unit matched to a target unit subject to mass conservation constraints”

$$M_{ij} = 1 - \text{corr}(x_i, y_j), \quad P_{ij}^{\star} = \arg \min_{P \geq 0} \sum_{i,j} P_{ij} M_{ij} \quad \text{SMD}(X, Y) = \sum_{i=1}^{N_x} \sum_{j=1}^{N_y} P_{ij}^{\star} M_{ij}$$



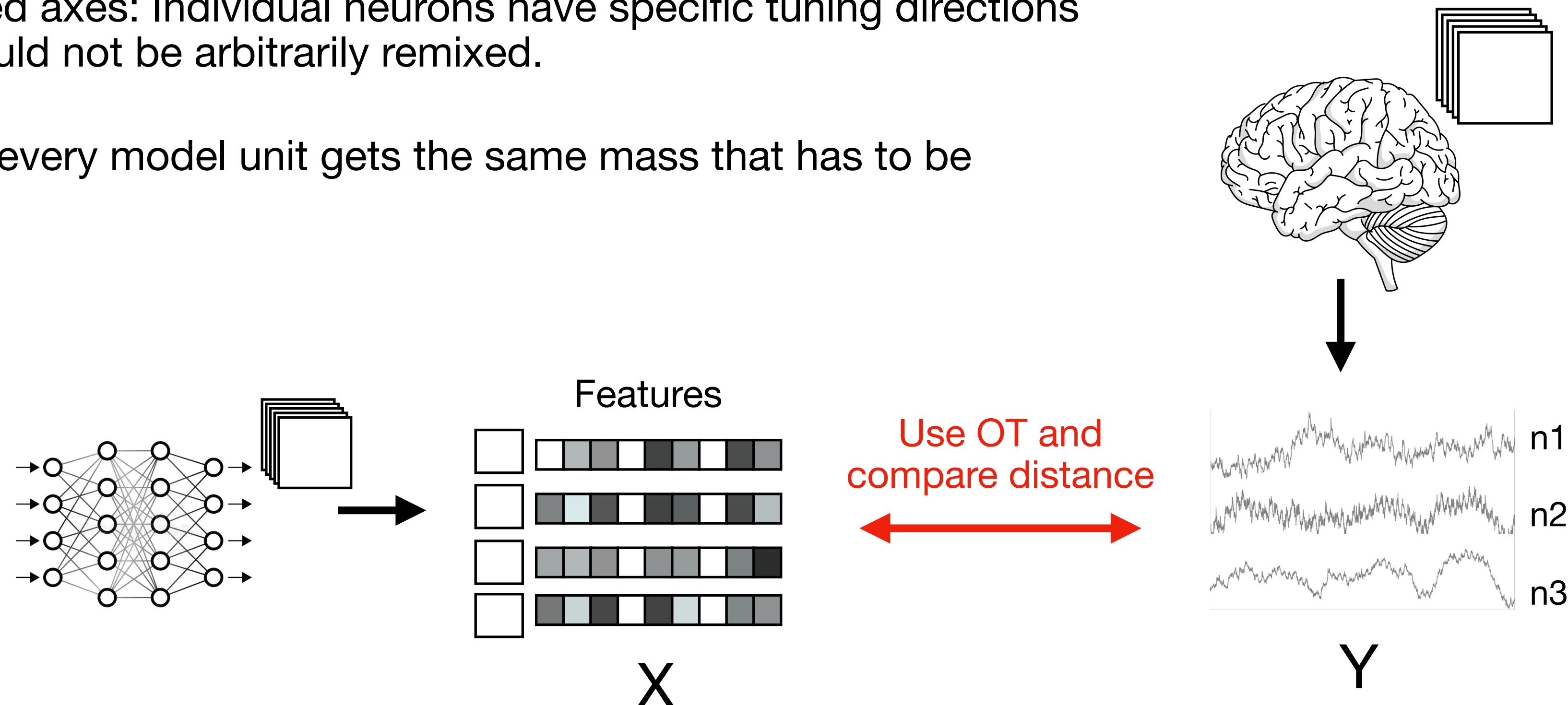
# Example: Soft Matching

**Idea:** Match individual model units to individual neurons without requiring an exact one to one match

❖ Solve an Optimal Transport (OT) problem: “how much is a source unit matched to a target unit subject to mass conservation constraints”

❖ **Pros:** Supports the idea of privileged axes: Individual neurons have specific tuning directions that matter mechanistically and should not be arbitrarily remixed.

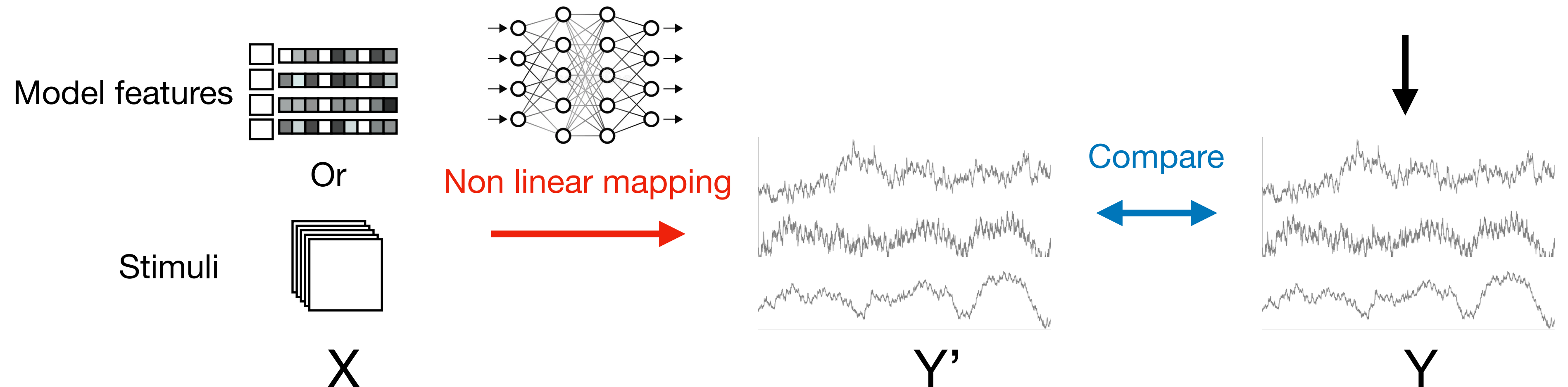
❖ **Cons:** Mass constraint means that every model unit gets the same mass that has to be distributed somewhere



# Nonlinear mapping

**Idea:** Use a neural network (transformer, convnet, etc) to learn the brain data from the stimuli or model features

- ❖ Predict neural data using back propagation
- ❖ **Pros:** Very useful for engineering purposes where explanation does not matter as much as prediction accuracy
- ❖ **Cons:** not great for forming theories and answering questions about the brain



# Selecting the right method

**What is your goal?**

**Studying the brain -> linear mapping methods, RSA, CKA, etc**

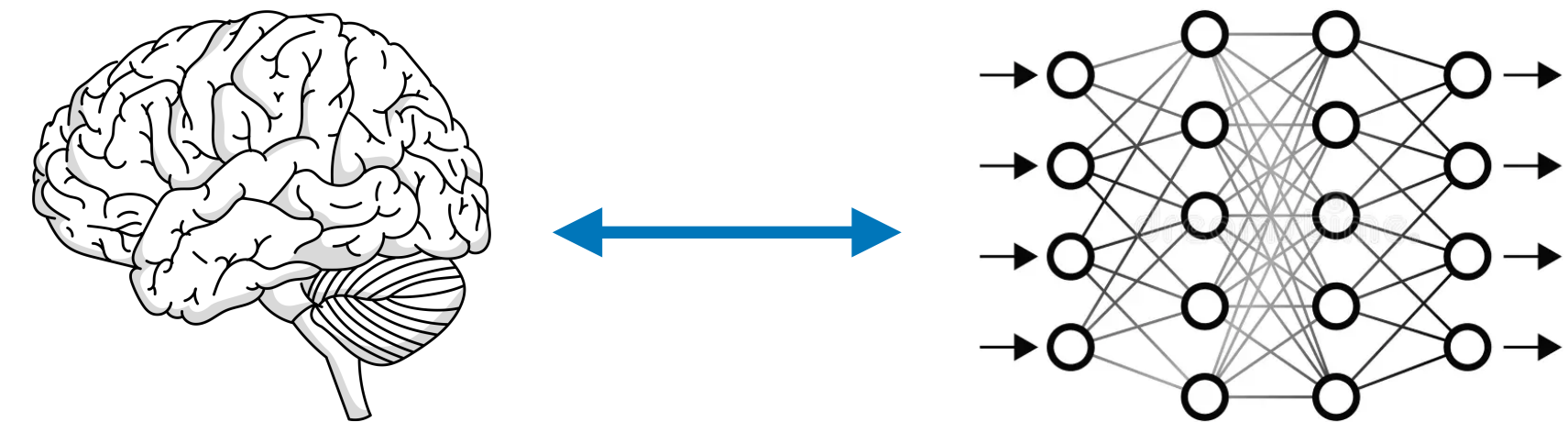
Building a model of the brain -> nonlinear mapping (brain foundation models)

# Selecting the right method (for studying the brain)

## Symmetry vs bidirectionally

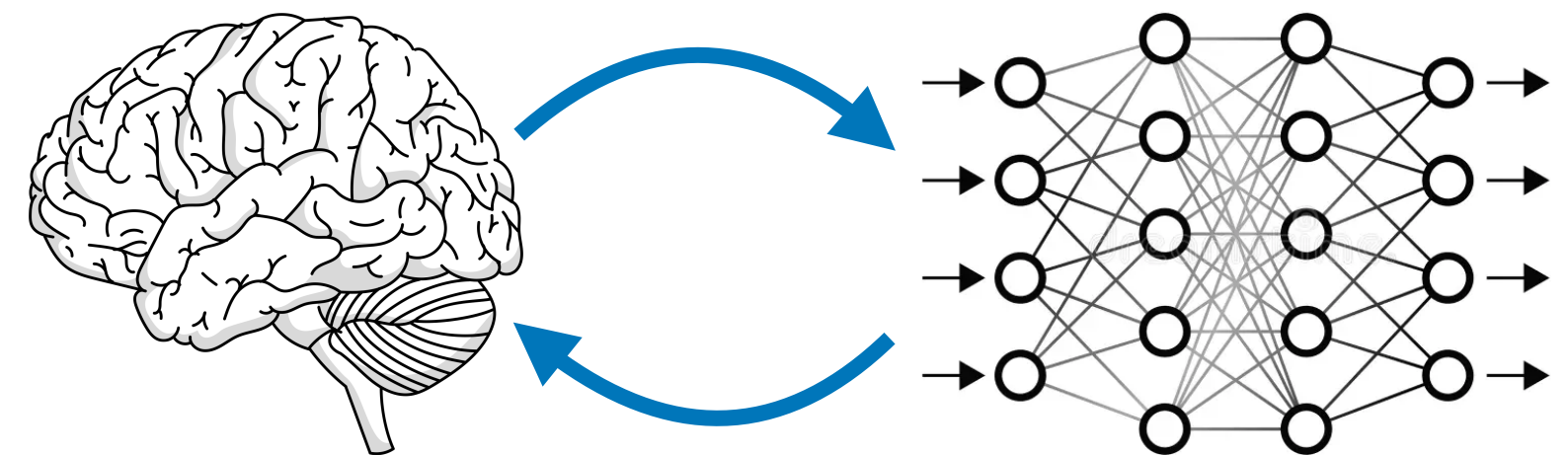
### Symmetry

- Many metrics are symmetric by definition (RSA, CKA, soft matching)
- **Problem:** we have access to all model units but often only a small amount of brain units



### Bidirectionally

- Brain-brain transform is not symmetric, why should model-brain be?



# The inter animal transform class (IATC) framework

- Identify the narrowest class of transforms that maps responses between subjects for a given brain area and species.
- The right class of transform should be:

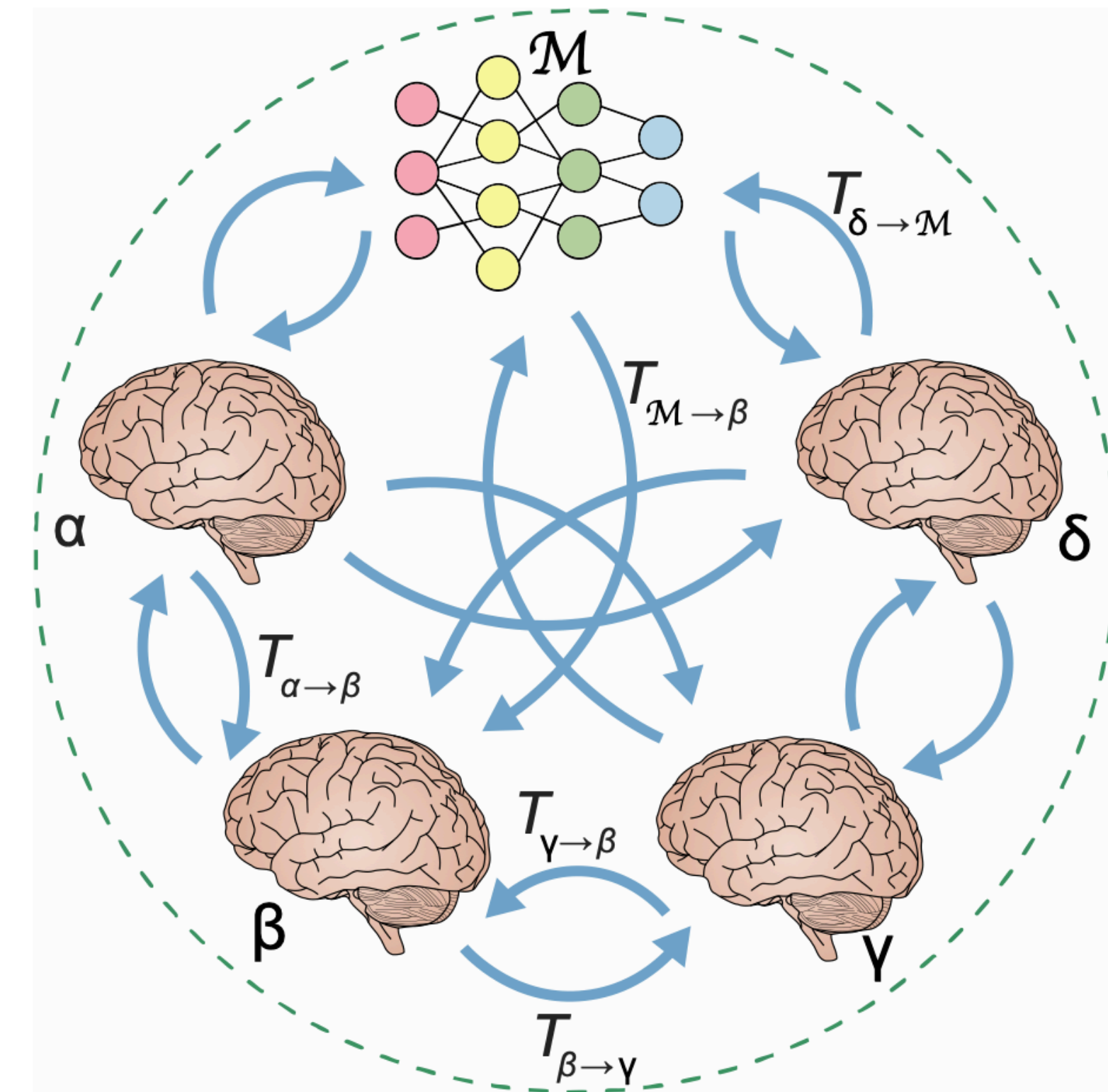
## Predictive

- Maximally predict neural responses

## Strict

- Distinguish brain areas while recognizing the same areas across subjects

*“how well can the model masquerade as a member of the population?”*



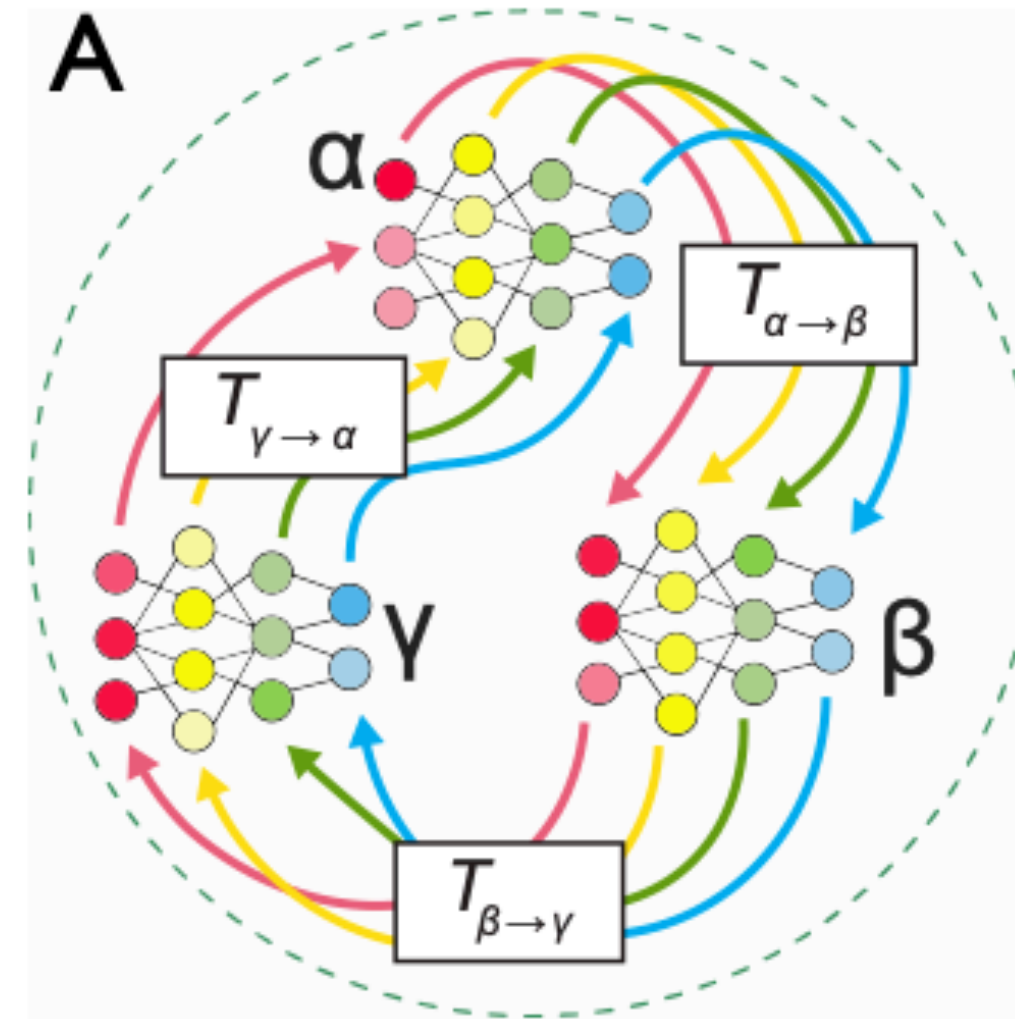
# Assessing same-area similarity in a model population

## Model: Modified AlexNet

- Trained with contrastive learning
- Softplus activation function + Poisson-like noise

## Population simulation:

vary the random seed  
controlling initialization  
and training data order.



Zippering effect

# Assessing same-area similarity in a model population

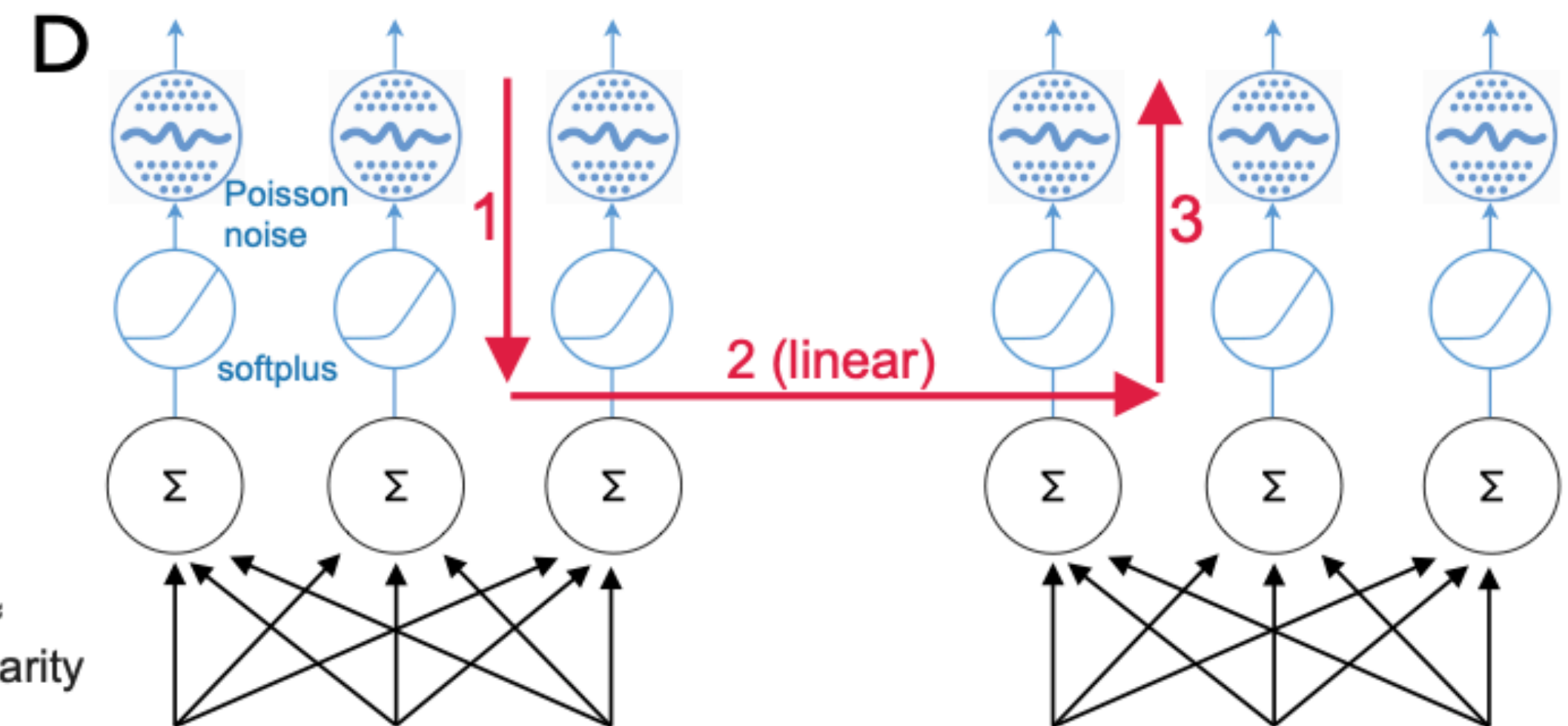
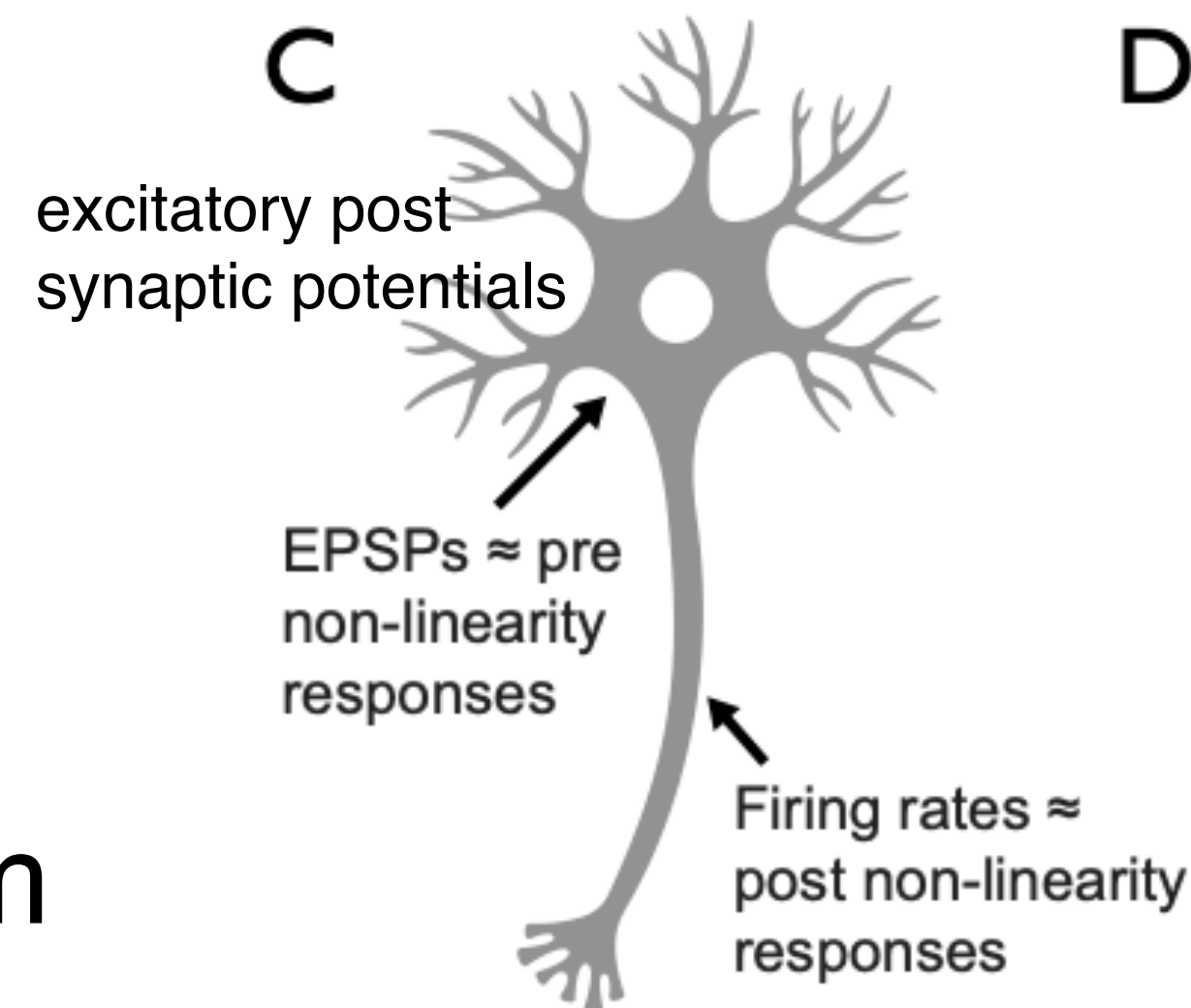
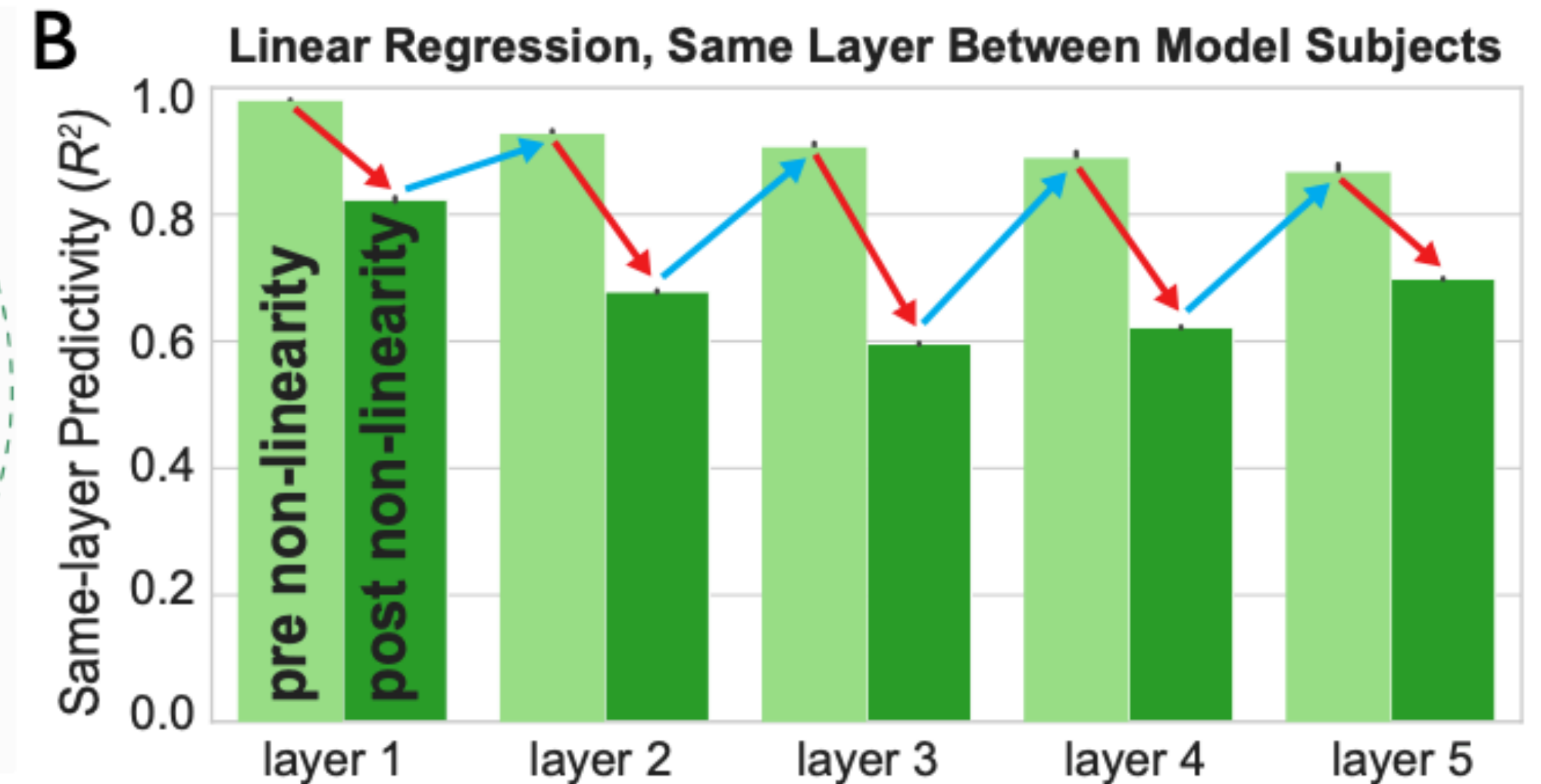
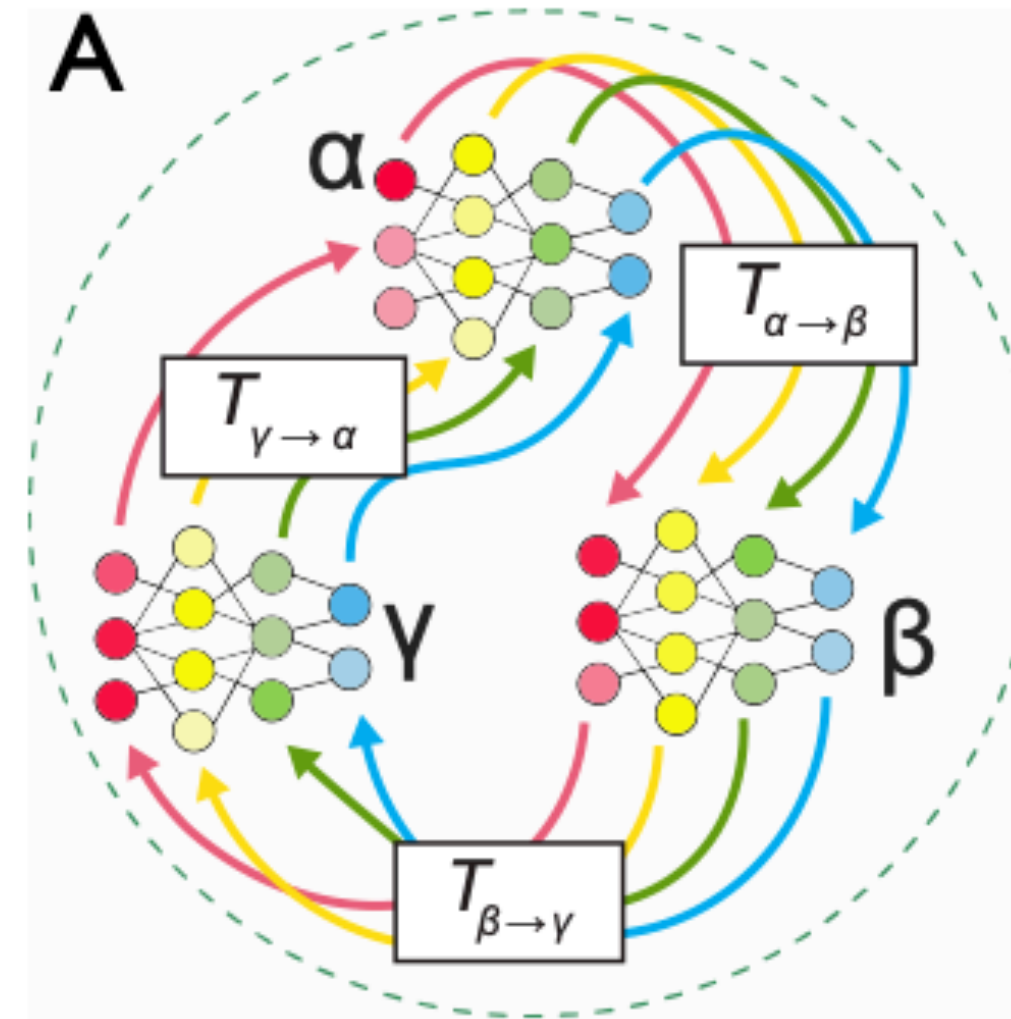
## Model: Modified AlexNet

- Trained with contrastive learning
- Softplus activation function + Poisson-like noise

## Population simulation:

vary the random seed  
controlling initialization  
and training data order.

## New transform class: The zippering transform

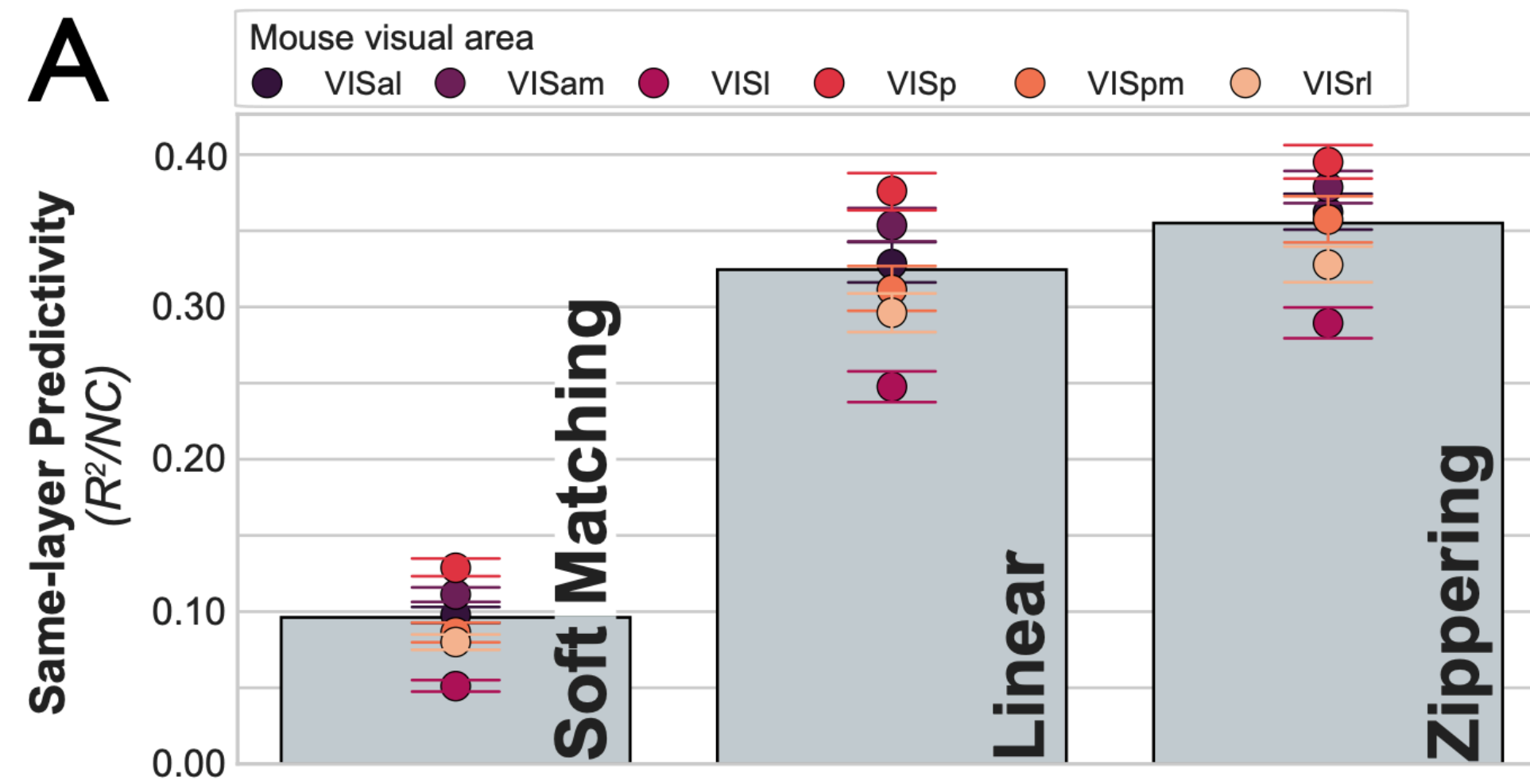


# Applying IATC to the mouse neural data

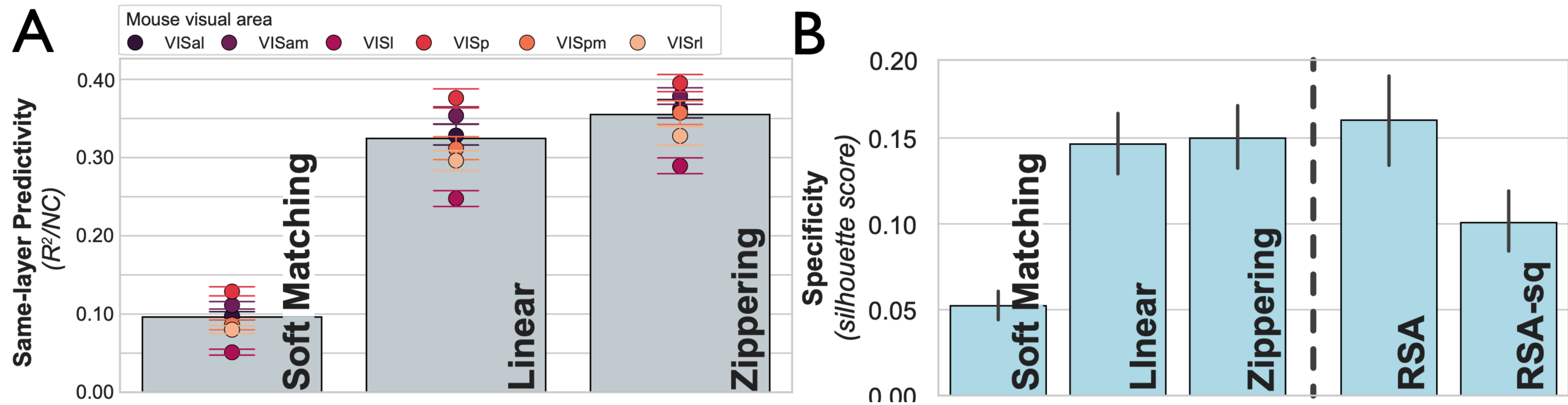
## Dataset

- Neuropixel recordings for 31 subjects
- 6 brain areas
- The mice passively viewed 118 different visual stimuli.

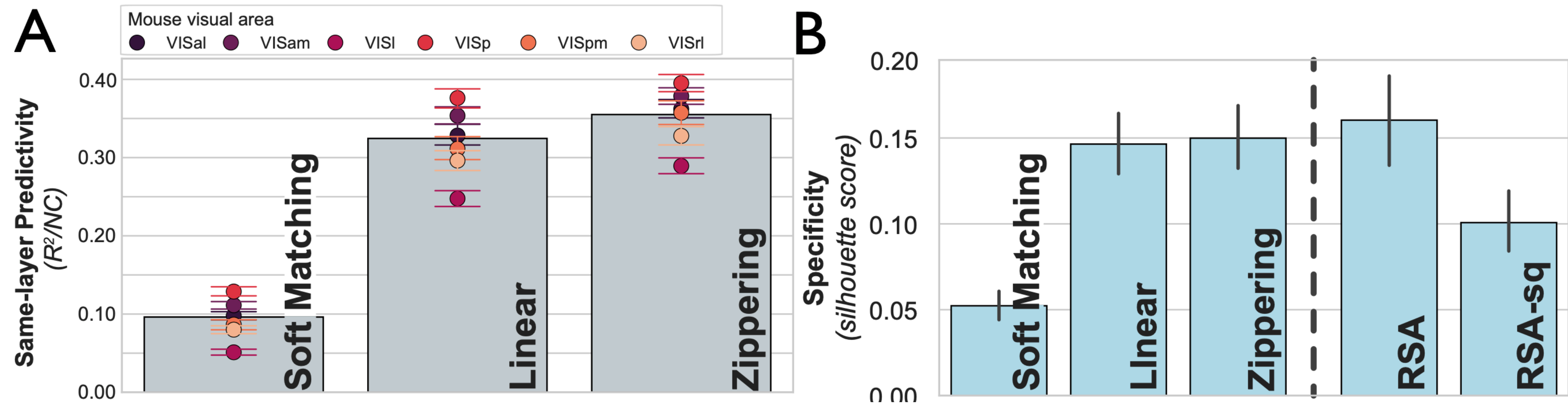
# Applying IATC to the mouse neural data



# Applying IATC to the mouse neural data



# Applying IATC to the mouse neural data



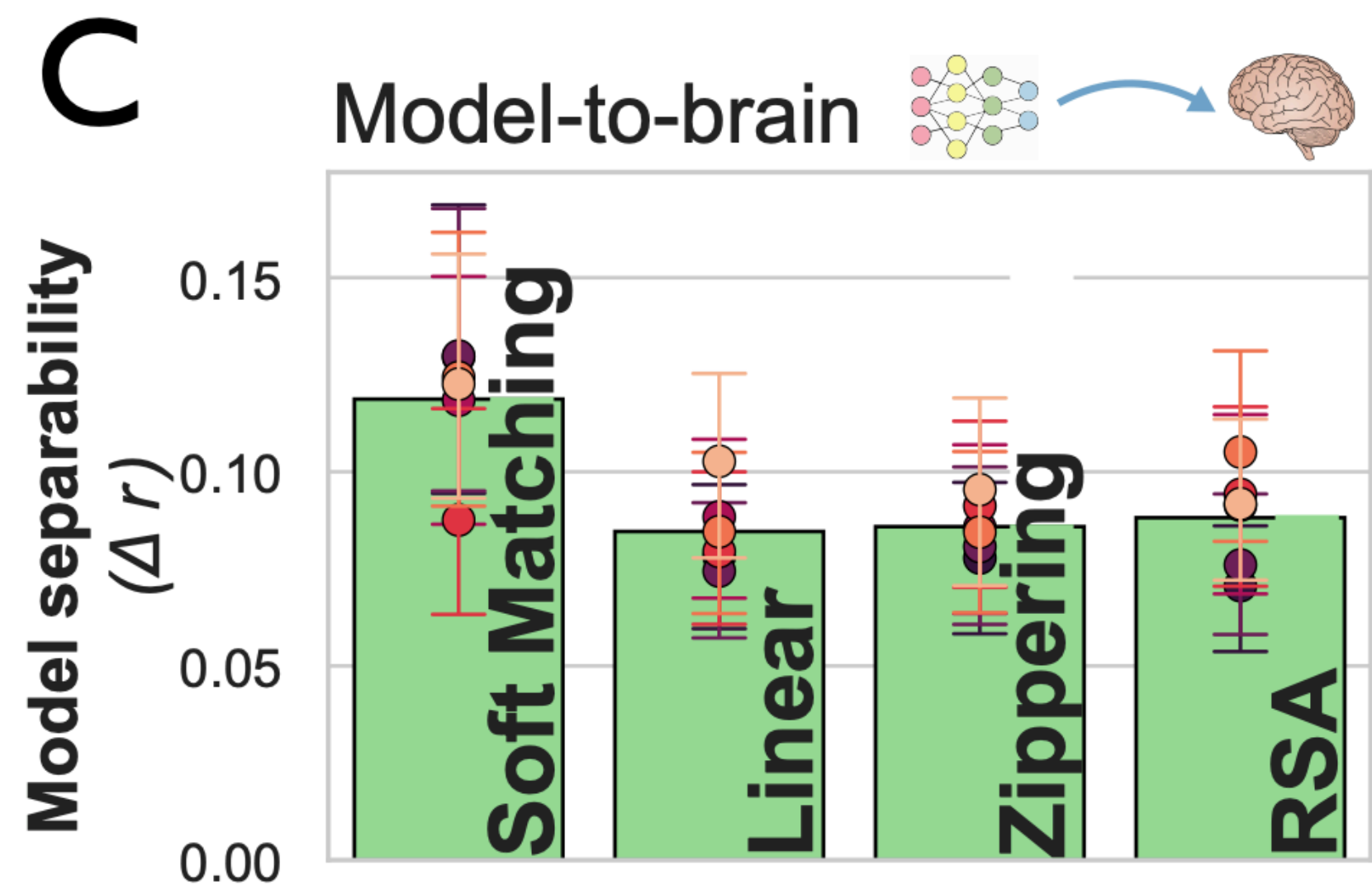
$$s(i) = \frac{b(i) - a(i)}{\max(b(i), a(i))}$$

**a(i):** within-area dissimilarity

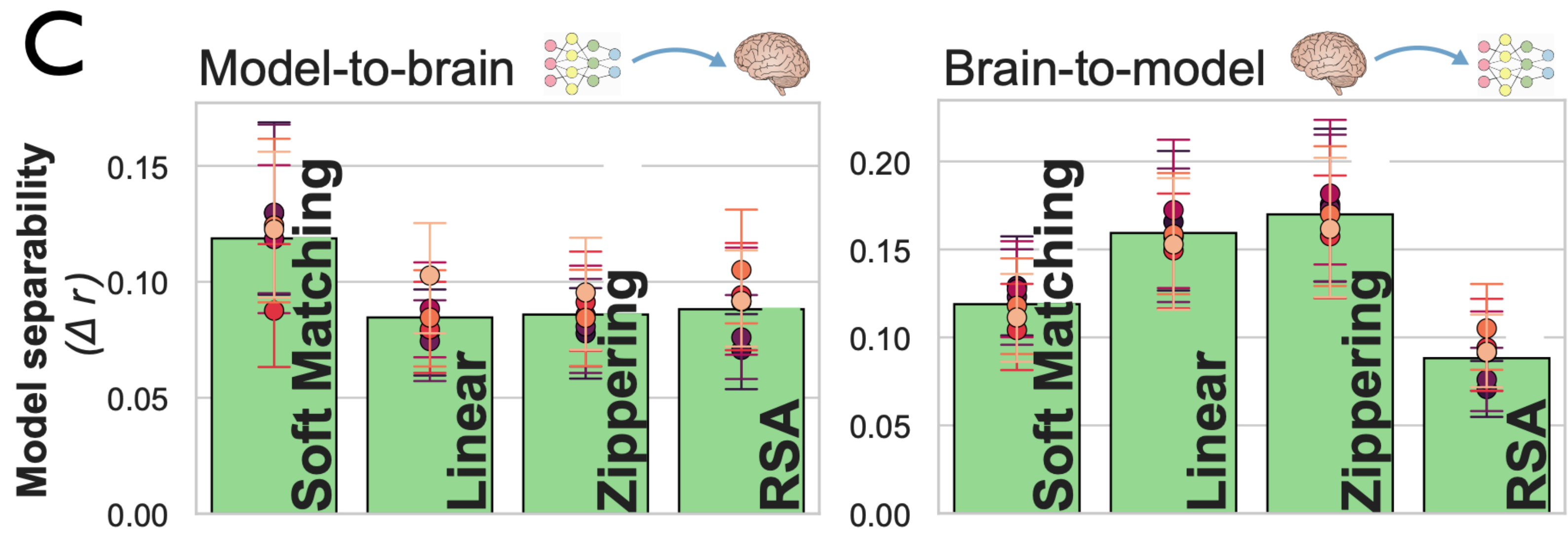
**b(i):** between-area dissimilarity

Good specificity: **low a(i)** and **high b(i)**

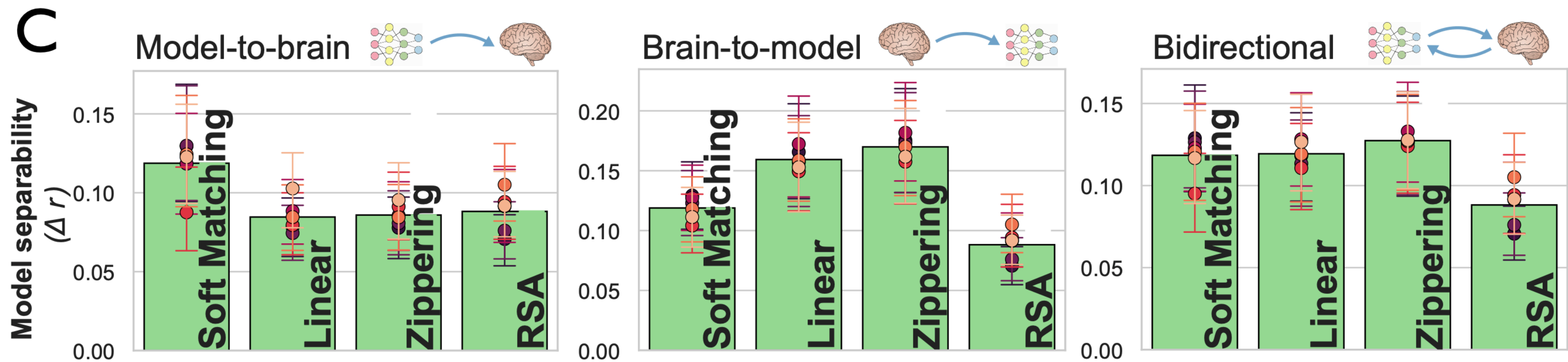
# IATC Guided Model Separability



# IATC Guided Model Separability



# IATC Guided Model Separability



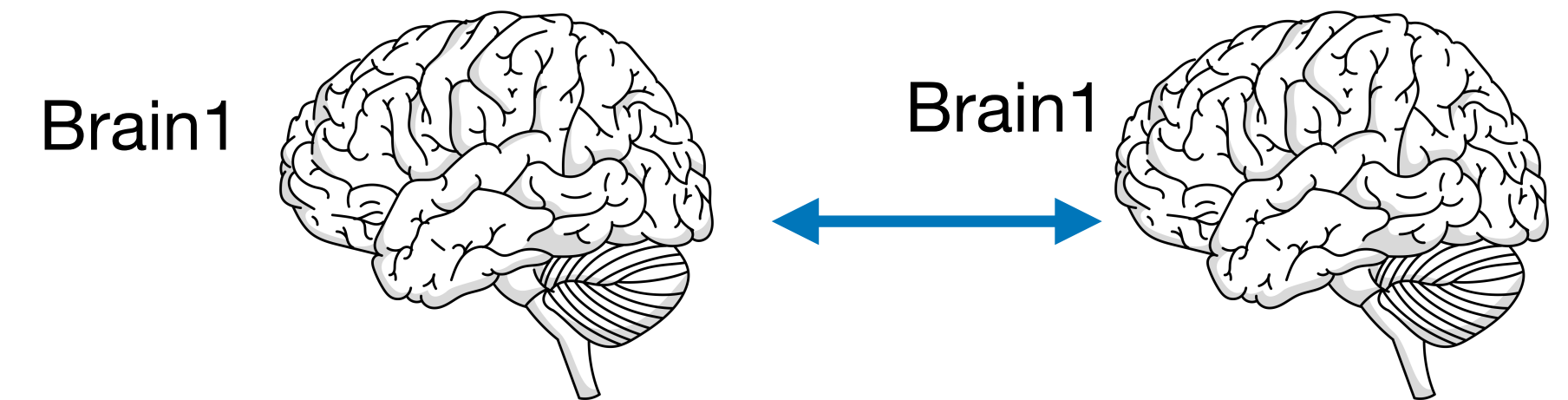
# Noise in neural data

- A fundamental challenge in evaluating the performance of NN models lies in the noise inherent in empirical data
- Examples of noise:
  - Motion artifacts (head motion)
  - Attention fluctuations
  - Arousal
  - Eye movements
- Why does this matter?
  - If a model only captures 20% variance in the data, this could indicate poor model performance.
  - If there is a high degree of noise, 20% may be as good as it gets.

# Noise ceiling estimates: Inter animal vs cross animal

## Inter animal spit-half reliability

- Common for evaluating individual participant data when you have repeats
- Repeated measurements are divided into two halves and the responses are correlated.



$$r_{\text{sh}} = \text{corr}(Y^{(1)}, Y^{(2)})$$

## Cross animal

- **Fundamental question:** How do we decide how to measure similarity across different animals?
  - Use the IATC framework to find the right class of transforms

