Object Category Structure in Response Patterns of Neuronal Population in Monkey Inferior Temporal Cortex

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1Research Group for Brain and Cognitive Sciences, School of Medicine, Shaheed Beheshti University, Tehran, Iran; 2School of Cognitive Sciences, Institute for Studies in Theoretical Physics and Mathematics, Niavaran, Tehran, Iran; 3Department of Neurobiology and Behavior, University of Washington, Seattle, Washington; 4Cognitive Brain Mapping Laboratory, RIKEN Brain Science Institute, Wako, Saitama, Japan; and 5Graduate School of Science and Engineering, Saitama University, Saitama, Saitama, Japan

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Kiani R, Esteky H, Mirpour K, Tanaka K. Object category structure in response patterns of neuronal population in monkey inferior temporal cortex. J Neurophysiol 97: 4296–4309, 2007. First published April 11, 2007; doi:10.1152/jn.00024.2007. Our mental representation of object categories is hierarchically organized, and our rapid and seemingly effortless categorization ability is crucial for our daily behavior. Here, we examine responses of a large number (>600) of neurons in monkey inferior temporal (IT) cortex with a large number (>1,000) of natural and artificial object images. During the recordings, the monkeys performed a passive fixation task. We found that the categorical structure of objects is represented by the pattern of activity distributed over the cell population. Animates and inanimates objects created distinguishable clusters in the population code. The global category of animate objects was divided into bodies, hands, and faces. Faces were divided into primate and nonprimate faces, and the primate-face group was divided into human and monkey faces. Bodies of human, birds, and four-limb animals clustered together, whereas lower animals such as fish, reptile, and insects made another cluster. Thus the cluster analysis showed that IT population responses reconstruct a large part of our intuitive category structure, including the global division into animate and inanimate objects, and further hierarchical subdivisions of animate objects. The representation of categories was distributed in several respects, e.g., the similarity of response patterns to stimuli within a category was maintained by both the cells that maximally responded to the category and the cells that responded weakly to the category. These results advance our understanding of the nature of the IT neural code, suggesting an inherently categorical representation that comprises a range of categories including the amply investigated face category.

INTRODUCTION

The mental representation of object categories has been a source of general and perpetual interest in cognitive neuroscience. Several imaging studies have investigated this representation in humans (e.g., Aguirre et al. 1998; Allison et al. 1994; Chao et al. 1999; Epstein and Kanwisher 1998; Gauthier et al. 1999, 2000; Haxby et al. 2001; Kanwisher et al. 1997; Martin et al. 1996; McCarthy et al. 1997) and suggest that several object classes such as faces, houses, animals, and tools are represented in human temporal cortex. However, the use of a limited (and potentially biased) stimulus set and posing a presumed category structure limits the scope of many of these studies. Studies at the level of single cells in non-human primates share these shortcomings. Although the existence of face-selective cells in the inferior temporal (IT) cortex of naive monkeys is well established (Bruce et al. 1981; Desimone et al. 1984; Kiani et al. 2005; Perrett et al. 1982; Perrett et al. 1982; Rolls and Tovee 1995; Tsao et al. 2006), the generality of the finding to the representation of other object categories is unknown.

In monkeys trained to categorize stimuli into a few arbitrary groups, some single cells in the prefrontal cortex show responses covering most stimuli in one of the learned categories (Freedman et al. 2001, 2002). In humans, some cells in medial temporal lobe structures such as the hippocampus respond categorically (Kreiman et al. 2000; Quiroga et al. 2005). Both prefrontal cortex and medial temporal lobe structures receive visual input about objects from IT cortex. Although the stimulus selectivity of IT cells is affected by training for visual categorization (Baker et al. 2002; Sigala and Logothetis 2002), responses of single IT cells appear to represent individual stimuli rather than the learned categories (Freedman et al. 2003; Vogels 1999). Taken together, these studies suggest that object categories could be represented in the polymodal association cortices downstream of the IT cortex. Although some types of categorization may rely on the prefrontal cortex and medial temporal structures, it is conceivable that certain classes of visual categories would be represented in the IT cortex. Given our rapid and seemingly effortless ability for categorization of natural objects (Li et al. 2002; Thorpe et al. 1996), natural categories provide plausible candidates.

A possibility, which has not been tested extensively, is that object categories are represented by response patterns over a large population of IT cells. According to this hypothesis, objects that belong to the same category would tend to elicit similar population response patterns. We therefore asked whether, for a large (>1,000) set of natural object images rather than arbitrary and limited sets of object images, responses of a population of IT cells (rather than single units) represent the categorical structure of objects. Responses to the stimuli were examined during a fixation task to investigate object categories independent of artificially imposed or task-dependent requirements. Furthermore, the use of a large stimulus set with many categories allowed us to examine the representation of object categories in a data-driven fashion without prior assumption of any particular category structure. We found that response patterns distributed over the IT cell
population represented many animate categories, as well as the structure among them.

METHODS

Three macaque monkeys (M. mulatta) were used, two for single-cell recordings and one for behavioral experiments. All experimental procedures complied with the guidelines of the National Institutes of Health and the Iranian Society for Physiology. The monkeys were raised in human houses and then in zoos before they were brought to the laboratory. Therefore it is likely that they had encountered many animate and inanimate objects in the course of their life.

Recordings and stimuli

In a preparatory aseptic surgery, a block for head fixation and recording chamber were anchored to the dorsal surface of the skull. The position of the recording chamber was determined stereotaxically referring to the magnetic resonance images (MRIs) acquired before the surgery. Action potentials of single cells were recorded extracellularly with tungsten electrodes (FHC, ME) from the IT cortex, on the right side for one monkey (monkey 1) and on the left side for the other monkey (monkey 2) while the monkeys were performing a fixation task. The electrode was advanced with an oil-driven manipulator (Narishige) from the dorsal surface of the brain through a stainless steel guide tube inserted into the brain down to 10–15 mm above the recording sites. Recording positions were evenly distributed at anterior 15–20 mm (monkey 1) or 13–20 mm (monkey 2) over the ventral bank of the superior temporal sulcus and the ventral convexity up to the medial bank of the anterior middle temporal sulcus with 1-mm track intervals (Fig. 1). The recording was not biased by response properties. The action potentials from a single neuron were isolated in real time by a template matching algorithm (Worgotter et al. 1986).

Responses of each cell were tested with 1,124 ± 71 (mean ± SD; median, 1,084) stimuli presented in a pseudorandom order. The stimulus set was repeated 9 ± 2 (median, 10) times for each recording site. The sequence of stimuli changed randomly in each repetition, and also for different recording sites, to avoid any consistent interaction between successively presented stimuli. The stimuli were colorful photographs of natural and artificial objects isolated on a gray background. The size of the larger dimension (vertical or horizontal) of each stimulus was ~7° of visual angle.

The monkey had to maintain fixation within ±2° of a 0.5° fixation spot presented at the center of the display. The eye position was measured by an infra-red eye-tracking system (http://staff.aist.go.jp/k.matsuda/eye/), which allowed a precision of 1 deg or less for the measurement of eye position. The presentation of stimulus sequence started after the monkey maintained fixation for 300 ms. Each stimulus lasted for 105 ms and was followed by another stimulus without intervening gap. The sequence stopped when 60 stimuli were presented or when the monkey broke the gaze fixation. The monkey was rewarded with a drop of juice every 1.5–2 s during the fixation. It has been shown that cells in the monkey IT cortex preserve their stimulus selectivity in rapid serial presentations as fast as 14–28 ms/stimulus (Edwards et al. 2003; Foldiak et al. 2004; Keysers et al. 2001). Also backward masking has a minimal effect on the initial part of neuronal responses when the stimulus onset asynchrony is >80 ms (Kovacs et al. 1995; Rolls and Tovee 1994).

Data analyses

The data set consisted of all the cells with reliable unit isolation throughout the stimulus presentation, regardless of the cell’s stimulus selectivity (n = 674). The spontaneous activity was measured in a 200-ms window immediately before the sequence of stimulus presentation initiated, and its SD was calculated across different sequences. We measured the neural activity for each stimulus presentation in a 140-ms window starting 71 ms and ending 210 ms after the onset of the stimulus. Responses to the last two stimuli in each sequence did not enter the analysis. To minimize the contamination of neural activity measurement by responses to the previous stimuli, we excluded presentations with large activity (exceeding the spontaneous activity by 2 × SD) in the 50-ms period immediately after the stimulus onset. This resulted in exclusion of 15% of the presentations. However, neither the contamination correction nor the exact size of the window was crucial to the results. Due to our rapid stimulus presentation paradigm we could not assess how our results might have changed by taking into account very late responses of the cells.

Similarity of response patterns measured with degree of correlation

Responses elicited in the population of cells were used to calculate a measure of similarity between stimuli (Fig. 2). First, the mean responses of a cell to the set of stimuli were arranged in a vector and were normalized by subtracting the mean response of the cell from the vector and then dividing it by its Euclidean length. The response normalization for single cells canceled the bias due to different baseline activity and different ranges of firing rates in different cells. Changes in the method of normalization did not change the basic results. Second, for any pair of stimuli, we calculated Pearson’s correlation coefficient (r) between patterns of responses evoked by the stimuli in the cell population. The distance (or dissimilarity) between two stimuli was quantified by 1 − r (neural distance). Two stimuli with similar response patterns in the cell population therefore have a small distance. The use of correlation coefficient as a measure of distance has the advantage of focusing on the population response pattern and discounting effects that nonspecifically change the firing rate of the IT population (e.g., contrast or luminance of the stimulus). Nevertheless, similar results were obtained with other distance metrics, such as Euclidean distance, which do not specifically measure the pattern similarity.

Multidimensional scaling and cluster analyses

Multidimensional scaling (MDS) (Young and Hammer 1987) was used to visualize the distribution of stimuli based on the neural
3 correlations are significant. The neural distances (Figs. 5 and S2, Tables 1 and 2, and supplemental information, with 9, and 12
A–C, and 6 of 9, and 12
A
B
C
D
E
F
G
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I
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K
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O
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X
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Z

Cluster analysis on stimulus similarity in low-level features

We also applied cluster analysis to the physical similarity between stimulus images (Fig. 6A). Physical similarity was measured by sum of absolute differences in red, green, and blue values over the pixels of two images, sum of absolute difference in intensity over all pixels, and sum of absolute differences in coefficients of Wavelet transformation of stimulus images. We used a biorthogonal wavelet (Daubechies 1992) from the Wavelet Toolbox of Matlab (bior 5.5 with 7 levels of decomposition), but similar results were obtained with other wavelets as well.

Similarity of the stimuli was also measured by outputs of a population of modeled V1 simple cells, and modeled V1 complex cells (Fig. 6A). The receptive fields of simple cells were simulated by Gabor filters of different orientations (0, 90, −45 and 45°), sizes (0.3–1.2° in steps of 0.1°), and contrast selectivity (preferring lightness or darkness at the center of the receptive field) for cells without color selectivity. To introduce color information, we replaced the intensity contrast with red-green or blue-yellow. Cells were distributed over the stimulus image with 0.04° intervals between the receptive field centers of adjacent cells. Negative values in outputs were rectified to zero. Each of the images was presented to the model separately. The receptive fields of complex cells were modeled by MAX operation (Lampl et al. 2004; Riesenhuber and Poggio 1999) on outputs of neighboring simple cells with similar orientation selectivity. Simple cells were divided into four groups based on their receptive field size (0.3–0.4, 0.4–0.6, 0.6–0.9, and 0.9–1.2°), and each complex cell pooled responses of neighboring simple cells in one of these groups. The spatial range of pooling varied in the four groups (4 × 4, 6 × 6, 9 × 9, and 12 × 12 for the four groups, respectively) (Riesenhuber and Poggio 1999). Similarity of responses evoked by two stimuli in the population of modeled cells was calculated either by the same method used for the responses in the IT cell population or by calculating the absolute value of difference in outputs of individual cells and summing over all cells. The results obtained by the latter method are shown in Fig. 6A, but similar results were obtained by the former method as well.

Cluster analysis on responses of model units tuned to randomly selected complex features

To examine whether the representation of object categories was a feature that could emerge without category knowledge in a population of units selective for complex features, we created model units with complex feature selectivity based on the hierarchical object recognition model of Riesenhuber and Poggio (1999) (Fig. 6B and the rightmost set of bars in Fig. 6A). This model effectively combines several experimental findings and consists of a hierarchy of units with increasingly complex stimulus selectivity and invariance. Units either perform template matching on their input to develop more complex pattern-specificity from simpler features (S units), or they perform a nonlinear operation (MAX) to develop invariance by pooling over units tuned to the same feature but at different positions or scales (C units). A hierarchy of units with these operations leads to C2 units, which are tuned to partially complex features and are invariant to changes in position and scale (roughly similar to V4 neurons). The model was implemented with 256 C2 units as described in Riesenhuber and Poggio (1999), except that due to the difference of image sizes in the two studies, S1 and C1 units’ receptive field sizes were similar to the simple and complex V1 cells described in the preceding

Fig. 2. Examples of correlation between response patterns evoked in the 674 cells by 3 pairs of stimuli. ● 1 of the cells, and the x and y values of each ● represent the normalized responses of the cell to the stimulus pair. The 3 pairs share a common stimulus, which is shown at the left. The Pearson’s correlation coefficient (r) was 0.35, 0.20, and −0.20 for A–C, respectively. All 3 correlations are significant.

We listed 23 intuitive object categories, based on human convention, with ≥12 category members in the stimulus set (see Table 1). For each of them and also for higher categories made of them, we examined whether there was a corresponding node in the tree. Two indices were defined for this purpose: ratio 1 = (number of category members under the node)/(total members of the category), ratio 2 = (number of category members under the node)/(total stimuli under the node).

The average of the two ratios was used as a score for the match between the category and the node. We searched for the node with the maximum score for each category. To determine the mean value and variation of the score expected by chance clustering of the stimuli we repeated the same procedure for a group of randomly selected stimuli of the same size as the category (Monte Carlo method). The match of the higher categories (see Table 2) with the nodes was examined by the same method.
text. The final stage of the hierarchy consisted of shape-tuned units (STUs), which were selective to the images in our stimulus set. Each STU received inputs from 32 C2 units that were most strongly activated by its preferred stimulus. We randomly selected 674 images from our stimulus set, and tuned the STUs to these images. The tuning width of the STUs was adjusted so that their response sparseness and response distribution matched the average of the recorded IT cells. The exact tuning width of STUs or the number of C2 units connected to each STU was not crucial for our basic results.

Selectivity of single cells for object categories

The selectivity of single cells for object categories was examined based on the 13 categories located at the lowest level in the tree of Fig. 5. We will refer to these 13 categories as “the lowest-level categories.” Responses to individual presentations of all the stimulus members within each category were pooled for the analysis. A cell was regarded category-selective if responses to the best category were significantly larger than responses to any of the other categories (Newman-Keuls post hoc, \( P < 0.05 \)) (similar to the 2 cells in Fig. 7, A and B). A cell was also regarded selective to a combination of categories if responses to any category within the combination were significantly \(( P < 0.05 )\) larger than responses to any of the categories outside of the combination (similar to the cell in Fig. 7C).

To visualize the overlap of response magnitude distributions between categories, the mean responses to individual stimuli were plotted against the normalized stimulus rank for each category (Fig. 7, right). The stimuli of each lowest-level category were ranked according to the magnitudes of mean responses, and the rank was normalized by the number of stimuli within the category. A normalized rank of one represents the stimulus that evoked the largest response within the category. To average these magnitude-rank profiles across the cells (Fig. 8, left), the mean responses to individual stimuli were first normalized by the maximum response in each cell, and then averaged across cells. In Fig. 8, the stimuli were divided into two groups, those in the preferred category (or preferred category combination) and those in the remaining categories.

In Fig. 8A, right, the normalized mean responses to the individual stimuli were averaged over 10–20 cells preferring the same category in each monkey. The magnitude-rank profiles were then averaged across categories and monkeys. In Fig. 8B, right, the normalized mean responses to individual stimuli were averaged among 11 or 20 cells selective to human faces in each monkey. The magnitude-rank profiles were then averaged for the two monkeys. Monkey and nonprimate faces were excluded in Fig. 8B to allow comparison with previous studies.

We measured the significance of differences between responses to suboptimal categories (Fig. 9) by comparing responses of a cell to different categories. For each cell, the lowest-level categories were ranked based on their average response magnitude, responses to individual presentations were pooled across stimuli belonging to each category, and the significance of difference in response magnitude was calculated for each pair of category rank (Wilcoxon test, significance defined as \( P < 0.05 \)). The proportion of cells showing a significant difference for each category-rank pair is presented for the 255 category-selective cells (Fig. 9A) and for the other 419 cells (Fig. 9B).

Correlation of mean response patterns between categories

For further evaluation of the contribution of cells without maximal responses to a category to the discrimination of the category (Fig. 10), we used a correlation analysis similar to the one employed by Haxby et al. (2001). The analysis was done in two stages. First, the members in each of the lowest-level categories were randomly divided into two equally sized groups, and the mean responses of each cell to each half-category group was calculated by averaging the normalized mean responses to individual stimuli. Mean responses of 674 cells to each half-category formed a response pattern. Second, the pairwise correlation of these response patterns was calculated for all possible pairs of categories \(( n = 91 \) ). The procedure was repeated 1,000 times with different random divisions of the categories, and the mean value of the correlation coefficient was obtained. Note that the correlation was calculated for responses to categories in this analysis, while the neural distances that were used for the previous analyses (e.g., the clustering analysis) were based on the correlations of response patterns to individual stimuli.

We performed three versions of the described correlation analysis. 1) The correlation was calculated for the responses of all 674 cells (Fig. 10, A). 2) The correlation was calculated after removing the cells that maximally responded to either of the two categories involved in each correlation calculation (Fig. 10, B). 3) Finally, the responses of the cells that did not respond maximally to either of the paired categories were decorrelated by shuffling the mean response values across cells. The cells maximally responding to either of the paired categories were not shuffled. The correlation was calculated for the combination of shuffled and nonshuffled data points (Fig. 10, C).

Spatial distribution of cells with similar categorical selectivity

To examine whether there was clustering of cells with similar category selectivity, we compared responses of cell pairs with different spatial distances in the recording region. The position of the recording sites were estimated based on the location of the guide tube, the reading of the manipulator, the estimated distribution and location of gray matter, and the depth of ventral brain surface (detected by a characteristic noise on arrival of the electrode tip to the ventral cortical surface). Cell pairs were grouped according to the distance of recording sites in the three-dimensional space: same recording sites \(( <0.5 \text{ mm}, 0.5 \sim 1 \text{ mm}, 1 \sim 0.5 \text{ mm}, \text{ and so on} \). Cell pairs were excluded from the analysis when one of the cells was located in the lower bank of STS and the other one in the convexity of IT.

The similarity of the cells’ responses was quantified by the coefficient of correlation between mean responses to individual stimuli (stimulus correlation, Fig. 11, left) or between mean responses to the lowest-level categories (category correlation, Fig. 11, right). The mean response to a category was obtained by averaging mean responses to individual stimuli in the category.

Behavioral experiment

A third monkey was used for a preliminary behavioral examination of the monkey's perceptual categories (Figs. 12 and S4). Prior to the experiment, the monkey was trained extensively for more than a year on a serial delayed matching-to-sample task with stimuli that were not included in our stimulus set. For the behavioral test, we selected 44 stimuli from the 1,084 in our set. The monkey, sitting comfortably in a monkey chair without head restraint, started each trial by pressing a lever. A fixation point appeared for 400 ms at the center of the display, followed by a sample stimulus that was shown for 300–500 ms. After a 1,000- to 1,300-ms delay, a test stimulus appeared for 300–500 ms. If the test stimulus was identical to the sample, the monkey was given a 1,000- to 1,300-ms delay, a test stimulus appeared for 300–500 ms. If the test stimulus was identical to the sample, the monkey had to release the lever, within 700 ms of the test stimulus onset, to obtain reward. If the test stimulus did not match the sample, the monkey had to keep pressing the lever until the appearance of a third stimulus, which was always identical to the sample. The monkey was rewarded only based on the identity of the stimuli not their membership in a particular category. We gradually introduced the 44 stimuli to the monkey and started the data collection when the mean performance reached 80%. During the data collection, we adjusted the length of the stimulus presentation and delay periods, within the specified ranges, to keep the monkey’s performance at 80–85%. Because the focus of...
our analysis was on nonmatch trials (see following text), such trials were presented slightly more frequently (50–65%, average: 55%). We calculated the probability of correct discrimination for each nonmatch stimulus pair, as an estimate of their perceptual distance for the monkey (Sands et al. 1982). There were 13–48 repetitions (mean, 29.7) for each stimulus pair in the data set. Kruskal’s nonmetric multidimensional scaling was performed on the resulting discrimination matrix (Fig. S4); it is equivalent to running MDS on the confusion matrix (Sands et al. 1982). Note that, our choice of a small subset (n = 44) of the stimuli is dictated by practical limitations. Testing all possible nonmatch pairs of the 1,084 stimuli would require several years of data collection.

RESULTS

We recorded activity from 674 neurons, in multiple data-collection sessions, in the anterior IT cortex of two macaque monkeys (Fig. 1) while the monkeys performed a fixation task. Responses of each neuron were examined with >1,000 colorful photographs and paintings of natural and artificial objects. The monkeys had not been trained for any categorization task previously.

We will first explain that response patterns distributed over the IT cell population represented our intuitive category structure. Then the distributed nature of the category representation will be shown.

IT response patterns form category clusters

Different cells in IT cortex increased or decreased their firing rate to different stimuli so that each stimulus elicited a particular pattern of response over the population of recorded cells. The response pattern is defined by the set of response magnitudes in the 674 cells. Stimuli that are closer to each other in our hierarchical category structure elicited more similar response patterns. This tendency is illustrated by the scatter plots in Fig. 2 for three exemplar stimulus pairs. For each pair of stimuli, the similarity of population response patterns was measured by Pearson’s correlation coefficient (r) of normalized response patterns evoked by the two stimuli (see METHODS). The coefficient was 0.35, 0.20, and −0.20 for the pairs shown in Fig. 2, A–C, respectively. The correlation coefficients varied from −0.31 to 0.54 across the stimulus set (Fig. 3). Generally, animate and inanimate objects evoked negatively correlated responses while animate objects evoked positively correlated responses. Within the group of animate objects, the highest correlations belonged to stimuli in the same intuitive category.

To visualize the relationship between the similarity of stimuli in our intuitive category structure and the similarity of the neural response patterns, we first used a MDS analysis. MDS has been used to infer the internal representation of stimuli based on neuronal responses or behavioral data (Cutzu and Edelman 1998; Hasselmo et al. 1989; Op de Beeck et al. 2001; Sugihara et al. 1998). MDS allows us to generate a low-dimensional layout of the stimuli based on the similarity of response patterns. We used 1 − r as a measure of distance between two stimuli (neural distance). Using a correlation coefficient has the advantage of focusing on the population patterns of responses rather than nonspecific response changes. The 1,084 stimuli were plotted in a low-dimensional space with interstimulus distances approximating the original neural distances.

The stimuli were roughly divided into four category clusters—faces, bodies, hands, and inanimate objects—that can be appreciated even in a two-dimensional (2D) projection of the space (Fig. 4A). Each of these clusters was further divided into smaller groups in other projections: faces were divided into human, monkey, and nonprimate animal faces (Fig. 4B), and bodies were also divided into several subgroups (Fig. 4C). The Scree plot (the percentage of unexplained variance plotted against the number of dimensions) indicates that 2D projections of the space can explain only ≲35% of the variance in the data (Fig. S1). Therefore it is important to note that each 2D map captured only a small part of the structure that appeared in a higher dimensional space.

IT categories resemble human-intuitive categories

To better understand the organization of the objects in the high dimensional space and to further examine the category structure reconstructed from the neural distances, we conducted an agglomerative cluster analysis of the neural distances. The analysis started with 1,084 nodes corresponding to 1,084 stimuli that were consistently used in the experiments. The nodes were connected to each other step by step to make larger nodes. In each step, the two nodes with the smallest distance were connected to make a new node, and all the stimuli were connected to a single node after 1,083 steps. The whole reconstructed tree is shown as a supplementary program. A one-dimensional alignment of the stimuli based on the tree is also shown in Fig. S2. The tree shows several levels of organization. In the first branching, the stimulus set was divided into animate and inanimate object groups. The animate object group was further divided into several meaningful categories, as expected from the MDS analysis.

To objectively determine which categories appeared in the tree, we listed 23 intuitive categories, based on human convention (see Table 1), that had ≥12 category members in the stimulus set and examined whether they and their combinations had significantly corresponding nodes in the tree. For each category, we calculated two ratios for each node in the tree: the fraction of category members that were located under the node (Ratio 1) and the fraction of stimuli under the node that were members of the category (Ratio 2). We then selected the node that gave the maximum averaged value of the two ratios. The
match was regarded as significant if the value exceeded by 4 SD the chance value calculated by a Monte Carlo method for randomly selected stimulus groups of the same size as the category and more than half of the category members were under the node (Ratio 1 > 0.5).

Many animate object categories at several hierarchical levels had significantly matching nodes (Fig. 5 and Tables 1 and 2). The tree also reconstructed positional relations among the animate object categories in our intuitive category structure: the global category of animate objects was divided into bodies, hands, and faces, and the categories of bodies and faces were divided into meaningful subcategories. Faces were divided into primate and nonprimate faces, and the primate face group was divided into human and monkey faces. For bodies, human, birds, and four-limb animals clustered together, whereas lower animals such as fish, reptile, and insects made their own cluster. Thus the cluster analysis formally showed that the similarity of population response patterns reconstructed a large cluster. Thus the cluster analysis formally showed that the similarity of population response patterns reconstructed a large cluster.

### Table 1.

<table>
<thead>
<tr>
<th>Category</th>
<th>Ratio 1</th>
<th>Ratio 2</th>
<th>Score</th>
<th>Chance</th>
<th>No. of Stimuli</th>
</tr>
</thead>
<tbody>
<tr>
<td>Animal face</td>
<td>0.38</td>
<td>0.94</td>
<td>0.66*</td>
<td>0.52</td>
<td>42</td>
</tr>
<tr>
<td>Monkey face</td>
<td>0.97</td>
<td>0.81</td>
<td>0.89*</td>
<td>0.52</td>
<td>39</td>
</tr>
<tr>
<td>Human face</td>
<td>0.97</td>
<td>0.98</td>
<td>0.84*</td>
<td>0.53</td>
<td>64</td>
</tr>
<tr>
<td>Hand*</td>
<td>0.93</td>
<td>1.00</td>
<td>0.96*</td>
<td>0.52</td>
<td>27</td>
</tr>
<tr>
<td>Bird body</td>
<td>0.16</td>
<td>1.00</td>
<td>0.58*</td>
<td>0.53</td>
<td>56</td>
</tr>
<tr>
<td>4-limb animal body*</td>
<td>0.71</td>
<td>0.88</td>
<td>0.73*</td>
<td>0.55</td>
<td>103</td>
</tr>
<tr>
<td>Human body*</td>
<td>0.95</td>
<td>0.93</td>
<td>0.94*</td>
<td>0.52</td>
<td>40</td>
</tr>
<tr>
<td>Butterfly*</td>
<td>0.53</td>
<td>1.00</td>
<td>0.76*</td>
<td>0.51</td>
<td>17</td>
</tr>
<tr>
<td>Other insects</td>
<td>0.19</td>
<td>1.00</td>
<td>0.59*</td>
<td>0.52</td>
<td>27</td>
</tr>
<tr>
<td>Reptile*</td>
<td>0.84</td>
<td>0.41</td>
<td>0.63*</td>
<td>0.52</td>
<td>19</td>
</tr>
<tr>
<td>Fish*</td>
<td>0.87</td>
<td>1.00</td>
<td>0.93*</td>
<td>0.51</td>
<td>15</td>
</tr>
<tr>
<td>Car*</td>
<td>0.87</td>
<td>0.83</td>
<td>0.85*</td>
<td>0.52</td>
<td>23</td>
</tr>
<tr>
<td>Tree</td>
<td>0.15</td>
<td>1.00</td>
<td>0.58*</td>
<td>0.51</td>
<td>13</td>
</tr>
<tr>
<td>Leaf</td>
<td>1.00</td>
<td>0.02</td>
<td>0.51</td>
<td>0.51</td>
<td>12</td>
</tr>
<tr>
<td>Flower</td>
<td>0.14</td>
<td>1.00</td>
<td>0.57</td>
<td>0.52</td>
<td>22</td>
</tr>
<tr>
<td>Fruit</td>
<td>0.12</td>
<td>1.00</td>
<td>0.56</td>
<td>0.51</td>
<td>17</td>
</tr>
<tr>
<td>Vegetable</td>
<td>0.11</td>
<td>1.00</td>
<td>0.56</td>
<td>0.51</td>
<td>18</td>
</tr>
<tr>
<td>Food</td>
<td>0.07</td>
<td>1.00</td>
<td>0.53*</td>
<td>0.52</td>
<td>44</td>
</tr>
<tr>
<td>Furniture</td>
<td>0.16</td>
<td>1.00</td>
<td>0.58*</td>
<td>0.52</td>
<td>25</td>
</tr>
<tr>
<td>Lamp</td>
<td>0.19</td>
<td>1.00</td>
<td>0.60*</td>
<td>0.51</td>
<td>21</td>
</tr>
<tr>
<td>Common tool</td>
<td>0.11</td>
<td>1.00</td>
<td>0.56</td>
<td>0.52</td>
<td>27</td>
</tr>
<tr>
<td>Kitchen utensil</td>
<td>1.00</td>
<td>0.03</td>
<td>0.52</td>
<td>0.51</td>
<td>19</td>
</tr>
<tr>
<td>Home appliance</td>
<td>0.15</td>
<td>1.00</td>
<td>0.58*</td>
<td>0.51</td>
<td>13</td>
</tr>
</tbody>
</table>

Ratio 1 = (number of category members under the node)/(number of all members in the category). Ratio 2 = (number of category members under the node)/(number of all stimuli under the node). Score = (Ratio 1 + Ratio 2)/2, with the asterisk indicating that the value is significantly larger than the chance value. Chance = chance score estimated by random clustering of stimuli, estimated by Monte-Carlo simulation. No. of stimuli = number of stimuli belonging to the category. The asterisk after category name indicates that the match is satisfying both of the following criteria: the score is significantly larger than the chance value, and Ratio 1 > 0.5.

categories for the monkey and magnifies the significance of the represented animate categories.

The tree in Fig. 5 was reconstructed based on responses of all 674 cells recorded from the two monkeys. The trees that were constructed for individual monkeys showed all the basic properties seen here: the first division into animate and inanimate objects, subdivision of animate objects into faces and bodies and further subdivision of faces and bodies into subcategories. The scores for categories, listed in Table 1 for the combined data, were well correlated between the two monkeys ($r = 0.8, P < 10^{-6}$). Hereafter, the 13 categories located at the

### Table 2.

<table>
<thead>
<tr>
<th>Category</th>
<th>Ratio 1</th>
<th>Ratio 2</th>
<th>Score</th>
<th>Chance</th>
</tr>
</thead>
<tbody>
<tr>
<td>Primate face*</td>
<td>0.97</td>
<td>0.91</td>
<td>0.94*</td>
<td>0.55</td>
</tr>
<tr>
<td>Face*</td>
<td>0.86</td>
<td>0.89</td>
<td>0.87*</td>
<td>0.57</td>
</tr>
<tr>
<td>Face + hand*</td>
<td>0.91</td>
<td>0.75</td>
<td>0.83*</td>
<td>0.58</td>
</tr>
<tr>
<td>4-limb animal + bird body*</td>
<td>0.83</td>
<td>0.83</td>
<td>0.83*</td>
<td>0.57</td>
</tr>
<tr>
<td>4-limb animal + bird + human body*</td>
<td>0.87</td>
<td>0.85</td>
<td>0.86*</td>
<td>0.59</td>
</tr>
<tr>
<td>Insect</td>
<td>0.41</td>
<td>0.90</td>
<td>0.65*</td>
<td>0.52</td>
</tr>
<tr>
<td>Fish + reptile*</td>
<td>0.85</td>
<td>0.74</td>
<td>0.80*</td>
<td>0.52</td>
</tr>
<tr>
<td>Insect + fish + reptile*</td>
<td>0.71</td>
<td>0.85</td>
<td>0.78*</td>
<td>0.54</td>
</tr>
<tr>
<td>Body*</td>
<td>0.86</td>
<td>0.90</td>
<td>0.88*</td>
<td>0.64</td>
</tr>
<tr>
<td>Animal*</td>
<td>0.94</td>
<td>0.90</td>
<td>0.92*</td>
<td>0.72</td>
</tr>
<tr>
<td>Inanimate*</td>
<td>0.94</td>
<td>0.91</td>
<td>0.92*</td>
<td>0.76</td>
</tr>
</tbody>
</table>

Legends are similar to Table 1.
Low-level features cannot account for the categorical structure

To test whether the reconstruction of category structure was due to similarity of stimuli in low-level features, we applied the cluster analysis to low-level physical similarity (e.g., color) of the stimuli or similarity in responses of modeled V1 cell population. The modeled V1 cells had size and orientation selectivity, and some of them were color selective.

The trees that were built based on these similarity measures failed to show the categories (Fig. 6A). Therefore the representation of category structure in the population responses of IT cells appears to be a result of visual information processing after V1.

Randomly selected complex features cannot account for the categorical structure

We then examined whether the representation of category structure emerges trivially in a population of units tuned to various, but randomly selected, complex features. We tested this possibility by creating 674 STUs based on the hierarchical object recognition model of Riesenhuber and Poggio (1999). Each STU was tuned to a randomly selected stimulus, from our stimulus set, through adjusting its input connections from V4-like units (C2 units) in the model. The broadness of tuning of STUs was adjusted to match that of actual IT cells. Hence for individual STUs, the sparseness of responses and the information about the identity of stimuli were comparable with those of actual IT cells. We did not necessarily regard this model as the best model to simulate the real monkey IT; we used it only to create 674 model units tuned to complex images.

The population response patterns of STUs did not show any meaningful grouping of stimuli as demonstrated by the mixed distribution of categories in a 2D stimulus projection based on the MDS analysis (Fig. 6B). The tree reconstructed from the responses of the STUs also failed to represent the categories (Fig. 6A, rightmost set of bars). Similar results were obtained for the C2 population in the model. Because of the position and size invariance of C2 units and STUs, the failure cannot be attributed to the lack of invariance. These results suggest that the reconstructed object category structure based on the responses of actual IT cells reflects something about the monkey IT cortex beyond the representation of a randomly selected set of complex features.

Category membership can be read out by means of a linear classifier

Different categories elicited easily separable population response patterns in IT cortex. Full classification of the stimuli into different categories in the tree can be performed reliably even by a linear classifier. We used a simple two-layer perceptron network (Duda et al. 2001) with 674 cells in the input layer and the significant lowest-level categories in the output layer. The magnitudes of mean responses to each stimulus were introduced to the input units, and the output unit yielding the largest value was taken as the classification result. The network was trained by adjusting its connection weights until it achieved a perfect classification for a training data set.

After training with a random selection of 50% of the stimuli, the network correctly classified 86% of the remaining stimuli. This performance is not simply the result of a large degree of freedom in the parameters of linear classifier or a result of the high dimensionality of the response space. When stimuli were randomly assigned to 10 groups of the same sizes as the 10 categories, the performance of the linear classifier decreased to 50% [the 50% chance performance is because one group corresponding to the “other inanimate objects” included about a half of the stimuli (Table 1)].

Single-cell responses are less clearly categorical than population responses

To examine properties of the category representation in IT cell population, we determined the selectivity of individual
cells to the lowest-level categories or their combinations. Ten of the lowest-level categories were significant in the tree of Fig. 5, and the rest were added to indicate the category combinations significantly matching the higher nodes.

Some cells (184/674) discriminated stimuli in one of the categories from those in any of the other categories: responses to the category were significantly larger than responses to any other category (Newman-Keuls post hoc, \( P < 0.05 \); Table 3). Figure 7 illustrates responses of a cell that responded selectively to human bodies (A) and another cell that responded selectively to bodies of four-limb animals (B).

In addition to selectivity for the lowest-level categories, many cells discriminated a combination of categories from others (145 cells, 186 combinations): responses to any category within the combination were significantly larger than responses to any of the categories outside the combination (\( P < 0.05 \)). Many of these combinations (67/186) matched significant higher-level nodes in the tree shown in Fig. 5 (Table 3), although these nodes were only a very small fraction (1.3%) of all possible combinations of the lowest-level nodes. Figure 7C shows responses of a cell that selectively responded to bodies of humans, birds, and four-limb animals.

In summary, a total of 255 cells (38% of 674 cells) were selective to a category or a combination of categories. The number is smaller than the sum of the two abovementioned category-selective groups (184 + 145) because 74 cells showed selectivity to both a single category and a combination of categories (as did the cell in Fig. 7B).

The categorical selectivity of single cells was imperfect compared with that of the cell population as a whole. For single cells, the distribution of response magnitudes for the preferred category (or category combination) largely overlapped with the distribution of responses to other categories (Figs. 7, rightmost column for the example cells, and 8A, left). This was true even for the cells selective to the face categories (Fig. 8B, left), although a previous study (Tsao et al. 2006) has shown that there may be a cluster of cells in more posterior parts of monkey IT cortex with near perfect face-selectivity. Unlike this overlap in single-cell responses, when responses to individual stimuli were averaged over 10–20 categorical cells preferring the same category (tested for monkey faces, human faces, human bodies, or hands), the overlap largely disappeared (Fig. 8, right). These results suggest that partial deficits in categorical selectivity of individual cells can be compensated by averaging responses over a small population of cells preferring the same category.

<table>
<thead>
<tr>
<th>Category</th>
<th>Number of Cells</th>
<th>Averaged Maximum Response, Spikes/s*</th>
<th>Average Spontaneous Response, Spikes/s</th>
</tr>
</thead>
<tbody>
<tr>
<td>Animal face</td>
<td>13</td>
<td>37.3</td>
<td>6.3</td>
</tr>
<tr>
<td>Monkey face</td>
<td>19</td>
<td>32.4</td>
<td>6.3</td>
</tr>
<tr>
<td>Human face</td>
<td>54</td>
<td>39.5</td>
<td>7.6</td>
</tr>
<tr>
<td>Hand</td>
<td>26</td>
<td>38.8</td>
<td>5.5</td>
</tr>
<tr>
<td>Bird</td>
<td>2</td>
<td>51.0</td>
<td>7.8</td>
</tr>
<tr>
<td>Four-limb animal</td>
<td>5</td>
<td>32.8</td>
<td>4.7</td>
</tr>
<tr>
<td>Human body</td>
<td>36</td>
<td>28.3</td>
<td>4.6</td>
</tr>
<tr>
<td>Butterfly</td>
<td>10</td>
<td>53.7</td>
<td>11.5</td>
</tr>
<tr>
<td>Other insects</td>
<td>3</td>
<td>49.1</td>
<td>2.8</td>
</tr>
<tr>
<td>Reptile</td>
<td>3</td>
<td>40.6</td>
<td>6.1</td>
</tr>
<tr>
<td>Fish</td>
<td>5</td>
<td>52.4</td>
<td>6.4</td>
</tr>
<tr>
<td>Car</td>
<td>8</td>
<td>28.2</td>
<td>5.6</td>
</tr>
<tr>
<td>Other inanimate objects</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Primate face</td>
<td>10</td>
<td>40.8</td>
<td>6.1</td>
</tr>
<tr>
<td>Face</td>
<td>37</td>
<td>39.3</td>
<td>6.7</td>
</tr>
<tr>
<td>Face + hand</td>
<td>4</td>
<td>62.5</td>
<td>12.1</td>
</tr>
<tr>
<td>Four-limb + bird body</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Four-limb + bird + human body</td>
<td>5</td>
<td>35.8</td>
<td>8.9</td>
</tr>
<tr>
<td>Insect</td>
<td>2</td>
<td>22.9</td>
<td>1.2</td>
</tr>
<tr>
<td>Fish + reptile body</td>
<td>3</td>
<td>41.3</td>
<td>5.0</td>
</tr>
<tr>
<td>Insect + fish + reptile body</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Body</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Animale (face + body + hand)</td>
<td>6</td>
<td>49.9</td>
<td>10.3</td>
</tr>
<tr>
<td>Inanimate</td>
<td>0</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*The mean firing rate in the 140-ms response window to the best stimulus for individual cells was averaged over the cells that responded selectively to the category.
Responses to suboptimal categories contribute to the category representation

Another aspect of the distributed nature of category representation in IT cortex is significant difference of neural responses even for categories that a neuron is not maximally responsive to. For each cell, we compared the magnitude of responses (Wilcoxon test) for all pairs of the lowest-level categories after ranking the categories according to the cell’s mean category responses. The proportion of cells with significant difference ($P < 0.05$) for each rank pair is shown for the 255 category-selective cells and for the other cells in Fig. 9. Significant differences were widely distributed over different rank combinations. For example, >50% of cells showed significantly different responses for a rank difference of five categories after ranking the categories according to the cell’s mean category responses. The proportion of cells with significant difference ($P < 0.05$) for each rank pair is shown for the 255 category-selective cells and for the other cells in Fig. 9. Significant differences were widely distributed over different rank combinations. For example, >50% of cells showed significantly different responses for a rank difference of five.

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**Fig. 7.** Three examples of category-selective cells. Example responses to individual members of the preferred category (left), the averaged responses to the lowest-level categories (middle), and the magnitude of responses to individual stimuli of the lowest-level categories plotted against the normalized stimulus rank within each category (right) for a cell preferring human bodies (A), 4-limb animal bodies (B), and the combination of human bodies, 4-limb animal body, and bird bodies (C). Left and middle: horizontal bars indicate the stimulus presentation period. Middle and right: the best categories are shown in red. The categories that evoked responses significantly smaller than the best category but significantly larger than other categories (bird and reptile in B; reptile, fish, and other insects in C) are shown in blue. Gray indicates other categories. Normalized rank of 1 indicates the stimulus that evoked the largest response within the category.

**Fig. 8.** The overlap of average response magnitudes of stimuli in the preferred category (black lines) with responses to other stimuli (gray lines) for all the categorical cells (A) and for cells selective to human faces (B). A and B, left: mean responses to individual stimuli were normalized by the maximum mean response in each cell, and then responses to stimuli of the same normalized rank were averaged across cells. Normalized rank of 1 indicates the largest response in the stimulus group. A, right: normalized mean responses to the same stimulus were averaged over 10–20 cells preferring the same category in each monkey (performed for monkey faces, human faces, human bodies, or hands). The resulting magnitude-rank curves were then averaged across categories and monkeys for the figure. B, right: similar to A but the normalized mean responses to individual stimuli were averaged among 11 or 20 cells selective to human faces in each monkey. Monkey faces and nonprimate faces were excluded in B to allow comparison with previous studies.

**Fig. 9.** Many IT cells showed significant differences in their responses to suboptimal categories. For each cell, the lowest-level categories were ranked based on the average response magnitude, and the significance of difference in response magnitudes was calculated for each pair of category ranks (Wilcoxon test, significance defined as $P < 0.05$). Individual trial responses pooled for all the stimuli belonging to each category were used for the comparison. The proportion of cells that showed a significant difference for each category-rank pair is color-coded for the 255 category-selective cells (A) and the remaining 419 cells (B).
between categories among the category-selective cells (Fig. 9A). This high probability of significant difference implies that not only responses to the best categories but also those to other categories, ranging from suboptimal to the worst, carry information. For the cells without sharp category selectivity, a rank difference of eight was enough to achieve significant difference in 50% of cells (Fig. 9B).

The presence of categorical information in responses to suboptimal categories was also demonstrated by examining the correlations in the population response patterns to each category. In this analysis, each lowest-level category was randomly divided into two groups of equal size, responses of each cell to individual stimuli were averaged across stimuli in each half-category, and the correlation of averaged responses were calculated for all possible pairs of categories across the cell population. The response to a half-category was more similar to, or more correlated with, responses to the other half of the same category than another category (Fig. 10, □). This was true even after removing the cells that maximally responded to either of the categories in the pair (Fig. 10□), meaning that a category can be potentially discriminated from another one even in the absence of maximally responsive cells. For example, images of four-limb animal bodies were discriminated from images of fish even without the cells selective to either category. As a complementary test, when we shuffled the mean responses of cells with suboptimal responses to the two categories, ranging from suboptimal to the worst, carry information. For the cells without sharp category selectivity, a rank correlation values (Fig. 11, left) compared with the previous finding of local clustering of cells with similar stimulus selectivity (Fujita et al. 1992; Tamura et al. 2005; Tsunoda et al. 2001; Wang et al. 1996, 1998; Yamane et al. 2006). The category correlation had a similar tendency, but showed stronger correlation values (Fig. 11, right) compared with stimulus correlations.

Cells with similar category selectivity were usually found in the same penetration, and clusters of cells in neighboring penetration sites (1-mm interval on our grid system) were rarely selective to the same category. This is reflected in the small change in the category correlations beyond 1 mm in Fig. 11. Instead of creating a big spatial cluster, cells with similar category selectivity appeared in multiple small clusters distributed over the recorded region, as shown for the global face category and global body category in Fig. S3. This may be another aspect of the distributed representation of object categories in the IT cortex. However, the number of cells recorded in each hemisphere (322 cells in the 1st monkey and 352 cells in the 2nd one) was not large enough to let us draw strong conclusions about the topography of categorical representation in IT. It is also important to note that our analysis in Fig. 11 measured the spatial extent of clusters by making the assumption that such clusters were spherical. It is possible that the topography of IT consists of nonspherical neural clusters with similar category selectivity that extends in one spatial dimension for distances >1 mm. A finer and more extensive sampling of IT cortex is required to test this possibility.
The monkey’s perceptual categories were examined by analyzing the probability of confusion between stimuli for a third monkey in a delayed matching-to-sample task. The task required discrimination of stimulus pairs based on their identity. We chose 44 stimuli from the stimulus set that was used in the recording experiments. The frequency of the erroneous responses (confusion) in this task can be influenced by the categorical similarity between stimuli (Sands et al. 1982): stimuli belonging to the same category or to closely related categories can be more often confused with each other.

The pattern of confusions between the stimuli corroborates the results based on IT responses. The 44 stimuli included 11 categories at the lowest level of the tree in Fig. 5: human faces, monkey faces, nonprimate animal faces, four-limb-animal bodies, bird bodies, fish, insects, reptiles, and other inanimate objects. Each category had four stimuli except the category of other inanimate objects, which had eight stimuli. The small number of stimuli reflects practical limitations in the length of experiments (see METHODS). We collected 51,806 trials in 27 sessions of data collection. Figure 12 shows the neural distances between stimulus pairs plotted against the probability of correct discrimination in nonmatch trials. Note that the probability of correct discrimination equals 1 − (probability of confusion). The probability of a correct discrimination was larger for stimulus pairs that elicited more distinct response patterns in IT ($r = 0.44, P < 10^{-6}$).

Using the MDS analysis, we projected the 44 stimuli on a 2D space according to the correct discrimination rates (Fig. S4). The stimuli in each of the lowest-level categories occupied a smaller region compared with the region occupied by the higher-level face or body categories.

**DISCUSSION**

**Representation of categories and category structure**

We found that the similarity of population response patterns evoked by object images in the monkey IT cortex was correlated with the distance between their categories in the intuitive category structure. The images of objects selected from the same category tended to evoke similar response patterns, whereas those of objects belonging to more distant categories evoked disparate patterns. The response pattern here is defined by the distribution of magnitudes of responses over the cell population. The correlation between the category distance and the response pattern similarity was strong enough for us to find the category structures in the distribution of stimuli plotted in a low-dimensional space according to the degree of similarity of the response patterns (Fig. 4). The reconstruction of category structure was verified by an agglomerative clustering analysis (Figs. 5 and S2 and supplementary program). The match between the category and stimuli under the node was as large as 0.86 on average for the significant lowest-level categories in Fig. 5 (Table 1). The information about categories can be easily read out from the activity of IT cells. For example, when a linear classifier, in which the 674 IT cells were connected to 10 output units with different connection weights, was trained with a half of the stimuli, it classified the remaining object images with nearly 90% accuracy.

Hung et al. (2005) have recently shown that membership of stimuli in a set of predefined categories can be extracted from responses of a population of IT cells. However, the small size and stimulus homogeneity of the categories and, more importantly, the pose of a predefined set of categories leave it unclear whether an inherently categorical representation of objects exists in IT. We provide the evidence for such a categorical representation for animate objects and show that responses of a population of IT cells represent both the individual categories and the intuitive relationship of the categories.
It is important to note that the MDS and cluster analyses revealing the grouping and cluster tree were data-driven in that they did not require prior specification of any category structure. The intuitive category structure was only used post hoc for quantitative assessment of the similarity between the IT stimulus clusters and our intuitive categories. The processes to make the MDS maps and the tree were completely data-driven. When we view the arrangements, we immediately notice that the clustering of objects appear to match with our intuitive categories. To quantify this, we listed 23 intuitive categories existing in the stimulus set with ≥12 category members and examined the match of the categories and their higher categories with the nodes in the tree. Categories at several hierarchical levels, especially those of animate objects, had significantly matching nodes. We conclude that response patterns over a large population of the monkey IT cortex reconstruct intuitive object categories and their structure. We do not intend to assume that the monkeys had all of the categories identified in the tree. It might be unlikely that the monkeys had the category of cars, for example. However, Sands et al. (1982) showed evidence suggesting that monkeys have the categories of human faces and monkey faces, discriminating them from each other and also combining them as a higher category and discriminating human and monkey faces from other objects. Our own preliminary behavioral test performed on a monkey with a subset of the stimuli \((n = 44)\) demonstrates that the distance of the stimulus pairs in the response space of IT is correlated significantly with the probability of confusing the stimuli in a delayed match-to-sample task (Fig. 12). The behavioral test also suggests that monkeys have a category structure similar to that of humans for animate objects (Fig. S4). These results indicate that the category structure of monkeys correlates, at least partly, with that of humans.

Distributed nature of the representation

Although ~40% of IT cells showed significantly larger responses to stimuli in one category (or a combination of categories) than to stimuli in any of the remaining categories, the distribution of responses of each cell to the preferred category and other categories overlapped substantially (Fig. 8, left). The information that each cell carried about object categories was limited. The overlap in the magnitude of responses largely disappeared when responses to individual stimuli were averaged over 10–20 cells preferring the same category (Fig. 8, right). The averaging was effective because the mismatch between the cells’ stimulus selectivity profile and the preferred category was different in different cells. In other words, cells that were selective to a category complemented each other. Pooling of the responses to individual stimuli among cells with similar category selectivity, therefore, increased the information about the category membership of the stimuli (Vogels 1999). Because cells with similar category selectivity also clustered locally in the cortex (Fig. 11), the increase of information could be achieved by pooling the neural responses based on cortical position of the cells.

The results in the present study also indicate that the information about categories is largely distributed over the cell population. Single cells showed significantly different magnitudes of responses between many pairs of nonpreferred categories (Fig. 9). For example, some cells that maximally responded to human faces significantly discriminated bird bodies from inanimate objects. Correspondingly, the similarity of response patterns to stimuli in a category was maintained not only by the cells that maximally responded to the category but also by other cells that responded to the category with medium and weak responses (Fig. 10). Such suboptimal category selectivity helped the tree in Fig. 5 capture the appropriate relative positions of categories, e.g., fish were farther from monkey faces than four-limb animal bodies. The suboptimal category selectivity may also help the simultaneous classification of an individual stimulus at multiple category levels. For instance, although a human face is classified into the human-face category by the strongest responses in human-face cells, it can be classified into the global-face category based on submaximal responses in cells tuned to other face categories. Moreover, responses distributed over cells that are tuned to various animate categories can be used to classify the stimulus into the animate-object category.

How does the category structure emerge from feature selectivity?

Previous studies have shown that individual IT cells respond to moderately complex features of object images (Brincat and Connor 2004; Desimone et al. 1984; Fujita et al. 1992; Ito et al. 1994, 1995; Kobatake and Tanaka 1994; Tanaka et al. 1991). An important question that arises is whether responding to a set of moderately complex features by IT cells would automatically result in the representation of the category structure. To examine this possibility, we tuned a population of shape-tuned model units (STUs), which simulate monkey IT cells (Riesenhuber and Poggio 1999), to a set of randomly selected images from the stimulus set used in the present study. Both the MDS and clustering analysis failed to reconstruct the category structure from outputs of the STUs (the rightmost set of bars in Fig. 6A and the distribution in Fig. 6B). This result suggests that the monkey’s IT cortex does something more than just respond to a random selection of moderately complex features. There are a huge number of such features that IT cells could potentially be tuned for. However, IT cells do not randomly select their favorite features. Instead, they may select the features that are useful for the purposes of the monkey’s behavior. Categorical discrimination of object images may be one of the factors that dictate what features the cells should be tuned for.

The images of objects belonging to the same category or close categories in Fig. 5 may appear more similar to each other than those of objects belonging to distant categories. This intuitive impression, however, has to be more formally defined. The failure of reconstructing the category structure from the similarity of images in low-level features indicates that the similarity of stimuli in low-level features did not underlie the success of reconstruction based on IT response patterns. The failure with the outputs of STUs tuned to randomly selected object images also suggests that the similarity of stimulus images in terms of randomly selected complex features did not underlie the categorical clustering of IT response patterns. Our results suggest that the monkey IT specifically adapts complex features for the purpose of object categorization. The visual system may have found these features, through postnatal ex-
perience and possibly through evolutionary processes, and have implemented them in the selectivity of neurons in the IT cortex and its afferent stages (Baker et al. 2002; Kobatake et al. 1998; Logothetis et al. 1995; Miyashita 1988; Sakai and Miyashita 1994; Sigala and Logothetis 2002).

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GRANTS

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REFERENCES


Figure S1. Scree plot of the MDS analysis. The percentage of unexplained variance of the neural distances between stimulus pairs is plotted against the number of dimensions used in the MDS analysis. The arrows indicate values for one to ten dimensions.
**Fig. S2.** The arrangement of 1084 stimuli, which were consistently used in our experiments, according to the agglomerative clustering analysis. The analysis produced a 2D tree structure with stimuli arranged along one dimension and the branching of the tree represented in the orthogonal dimension. The stimuli are ordered from left to right in each row and then from top to bottom. The complete tree is available as a supplementary program.
Figure S3. Distribution of category selectivity in the 3D cortical space. The $d'$ index was calculated for each cell for the discrimination of the global face category from other stimuli, and also for the discrimination of the global body category from other stimuli. The $d'$ index was calculated based on the following formula

$$d' = \frac{[\mu(g) - \mu(ng)]}{\sqrt{\frac{\sigma^2(g) + \sigma^2(ng)}{2}}}$$

where $\mu(g)$ and $\sigma^2(g)$ are the mean and variance of responses to stimuli in the category and $\mu(ng)$ and $\sigma^2(ng)$ are those to other stimuli. The mean and variance for each stimulus group were calculated from mean responses to individual stimuli. A larger $d'$ implies better discrimination. AP, the anterior-posterior position relative to the ear bar; ML, the medial-
lateral position (right hemisphere in monkey K and left hemisphere in monkey S); Depth, the distance from the horizontal plane traversing the dorsal apex of the brain. We used a grid system with 1mm spacing between the holes to control the AP and ML position of the guide tube and electrode. The circles in the figure mark some of the category-selective neural clusters. Cells with similar category selectivity were usually clustered tightly in space. The selectivity to the same category sometimes spread across penetrations to form larger clustering, but, as a whole, the selectivity to the categories appeared repeatedly at several segregated positions distributed over a large part of the recorded region.
Fig. S4. Two-dimensional arrangement of the 44 stimuli used in the behavioral experiment. The MDS was performed on the probability of correct discrimination for the non-match stimulus pairs.
Supplementary Program. The supplementary program shows the detail structure of the tree created by agglomerative clustering analysis on the responses of IT cells (see Methods). It can be downloaded from

http://www.shadlen.org/roozbeh/Research/Categorization/tree.zip

The program runs on Windows operating system. Expand the zip file “tree.zip” in a folder. Make sure that the executable file “clusterviewer.exe” and the data file “HCA.dat” both show in the folder, and the images are in the subfolder “ImageSet”. Run the executable file. You will see the structure of the tree on the left pane of the program. The highest node of the tree is shown on the top. Below the highest node are the tree branches which lead to individual images. Individual image nodes are indicated by 8 digit numbers. Other nodes are assigned 4-digit numbers. The numbers are used just to facilitate differentiation of the nodes from each other, and to allow easy navigation through the tree. They don’t carry any special meaning. You can expand or collapse each branch of the tree to see its full or simplified structure. When you click on each of the nodes on the tree the list of images under the node will be shown on the right pane. It may take a few seconds to a couple of minutes to load the images, depending on the speed of your computer. The images are arranged from left to right in each row and the rows are ordered from top to bottom. Clicking on the highest node produces a list similar to Fig. S2.