

1 Evidence that the ventral stream codes the errors used in hierarchical 2 inference and learning

3 Running title: Error coding in the ventral stream

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12 AUTHOR CONTRIBUTIONS

13 E.B.I. and J.J.D. designed the experiments. E.B.I. carried out the experiments and performed the data
14 analysis. E.B.I. and C.F.C. designed the models. E.B.I., C.F.C., and J.J.D. wrote the manuscript.

15 ABSTRACT

16 Hierarchical feedforward processing makes object identity explicit at the highest stages of the
17 ventral visual stream. We leveraged this computational goal to study the fine-scale temporal
18 dynamics of neural populations in posterior and anterior inferior temporal cortex (pIT, aIT)
19 during face detection. As expected, we found that a neural spiking preference for natural over
20 distorted face images was rapidly produced, first in pIT and then in aIT. Strikingly, in the next 30
21 milliseconds of processing, this pattern of selectivity in pIT completely reversed, while selectivity
22 in aIT remained unchanged. Although these dynamics were difficult to explain from a pure
23 feedforward perspective, a model class computing errors through feedback closely matched the
24 observed neural dynamics and parsimoniously explained a range of seemingly disparate IT
25 neural response phenomena. This new perspective augments the standard model of online
26 vision by suggesting that neural signals of states (e.g. likelihood of a face being present) are
27 intermixed with the error signals found in deep hierarchical networks.
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32 INTRODUCTION

33 The primate ventral visual stream is a hierarchically organized set of cortical areas
34 beginning with the primary visual cortex (V1) and culminating with explicit (i.e. linearly
35 decodable) representations of objects in inferior temporal cortex (IT) (1) that quantitatively
36 account for invariant object discrimination behavior (2). Consistent with a feedforward flow of
37 processing from V1 to V2 to V4 to IT, neurons at higher cortical stages are more selective for
38 object shape and identity while being more tolerant to changes in object size and position (3)(4).
39 Formalizing object recognition as the result of a series of feedforward computations yields
40 models that achieve impressive performance on basic object categorization tasks (5)(6) similar
41 to the level of performance achieved by IT neural populations (7)(8). Importantly, these models
42 are only optimized to solve invariant object recognition tasks and are not trained on neural data,
43 yet their intermediate layers are highly predictive of time-averaged V4 and IT neural responses
44 supporting the longstanding belief that the computational purpose of the ventral stream is to
45 solve such tasks. Thus, the feedforward inference perspective provides a simple but powerful,
46 first-order framework for studying core invariant object recognition (9).
47

48 However, visual object recognition behavior may not simply be the result of a single
49 feedforward neural processing pass (a.k.a. feedforward inference), as evidenced by the

50 observed dynamics of neural responses in IT (10)(11)(12)(13). From a theoretical perspective,
51 feedback in a recurrent network might substantially improve online inference or support learning
52 (14)(15). For example, inferring that a particular object is present in an image in the presence of
53 local ambiguities (e.g. ambiguity due to noise, missing local information, or local information that
54 is incongruent with the whole) can be achieved by integrating top-down with bottom-up
55 information through feedback (16)(17)(18). Another possibility is that feedback from high level
56 representations to lower level representations is used to drive changes of the synaptic weights
57 of neural networks (i.e. learning) through unsupervised or supervised algorithms (e.g. error
58 backpropagation) so as to improve future feedforward inference in performing tasks such as
59 object recognition (19). These two possibilities are not mutually exclusive, and several
60 computational models aim to integrate both in a natural way by building feedback pathways onto
61 a feedforward architecture (see **Fig. 7** and **Discussion**).

62
63 These architectures constitute a broad class of recurrent networks that all have an error
64 computation which compares top-down predictions or targeted outputs against bottom-up inputs
65 to compute gradients for driving sensory inference and learning (19)(20). Though the need for
66 errors and subsequent gradient signals is theoretically well motivated, we do not know if such
67 computations (e.g. error backpropagation and hierarchical Bayesian inference) are implemented
68 in the brain. Currently, little conclusive neurophysiological evidence exists in support of these
69 models. For example, while it has been suggested that end-stopped tuning of V1 cells is the
70 result of feedback loops that “explain away” extended edges (21), this feedback-based
71 interpretation cannot be disambiguated from standard alternative interpretations utilizing local
72 mechanisms like lateral inhibition (which builds selectivity for shorter edges) (22) or
73 normalization (which reduces responses to large stimuli such as extended edges) (23). Thus,
74 these neural measurements taken from a single cortical area in which the mapping to a visual
75 behavior is unclear, only weakly constrain computational models, and an open question is
76 whether the dynamics of propagation of neural signals across stages of the visual hierarchy
77 performs anything beyond feedforward inference.

78
79 Here, to disambiguate between the different types of inference that can be implemented in
80 a hierarchy through feedforward, lateral, or feedback connections, we measured the temporal
81 dynamics of neural signals across three stages of face processing in macaque IT. We found
82 that many neurons in the intermediate processing stages reversed their initial preference – they
83 rapidly switched from face preferring to anti-face preferring. Standard feedforward models
84 including those employing local recurrences such as adaptation, lateral inhibition, and
85 normalization could not fully capture the dynamics of face selectivity in our data. Instead, our
86 modeling revealed that the reversals of face selectivity in intermediate processing stages are a
87 natural dynamical signature of a family of hierarchical models that use feedback connections to
88 implement “error coding.” We interpret this very good fit to our data as evidence that the ventral
89 stream is implementing a model in this family. If correct, this model family informs us that we
90 should not interpret neural spiking activity at each level of the ventral stream as only an explicit
91 representation of the variables of interest (e.g. is a face present?) but should interpret much of
92 spiking activity as representing the necessary layer-wise errors propagated across the
93 hierarchy. In addition, we find that, without additional parameter modifications, the same error
94 coding hierarchical model family explains seemingly disparate IT neural response phenomena,
95 hence unifying our results with previous findings under a single computational framework.

96
97
98 **RESULTS**

99 Neural recordings were made across the posterior to anterior extent of IT in the left
100 hemispheres of two monkeys (**Fig. 1**; example neurophysiological map in one monkey).
101 Recording locations were accurately localized *in vivo* and co-registered across penetrations
102 using a stereo microfocal x-ray system (~400 micron *in vivo* resolution)(24). The face versus
103 non-face object selectivity of each site was measured using a standard screen image set, and
104 neural maps of face versus non-face object selectivity were used to physiologically define
105 subregions of IT containing an enrichment of face preferring sites (a.k.a. face patches) (25).
106 Sites were assigned as belonging to a face-preferring cluster by their distance to the center, a
107 purely spatial rather than functional criterion. The identified subregions were present in both
108 monkeys and encompassed at least three hierarchical stages of face processing in posterior
109 (pIT), central (cIT), and anterior (aIT) IT (**Fig. 1**). We asked how neural signals are transformed
110 from the earliest (pIT) to the latest stage (aIT) of IT face processing.

111

112 **Images with typical versus atypical arrangements of face parts**

113 Converging lines of evidence from correlative and causal studies implicate the face patch
114 subnetwork in face detection (25)(26). However, these studies relied on images testing general
115 face versus non-face object discrimination (a.k.a. face detection) that can be solved using
116 differences across multiple feature dimensions (local contrast, spatial frequency, and number of
117 features). We sought images that would provide a more stringent test of the face subnetwork
118 near the limits of its detection abilities. Specifically, if these regions are performing accurate face
119 detection, they should specifically respond to the configuration of a face even when all other
120 features are matched. Thus, we probed the face selectivity of IT with face-like images that only
121 differed in the configuration of the face parts such that there could be conflicting local (parts)
122 and global (configuration) evidence as to whether a face was truly present (**Fig. 2a**). For
123 example, images with an eye presented in an unnatural, centered position have a very different
124 global interpretation (“cyclops”) than when the eye is presented in its natural lateralized position
125 in the outline even though these images have very similar local information (eye presented in an
126 outline in the upper visual field) (see examples in **Fig. 2a** and **Supplementary Fig. 1b**). These
127 images create an “aperture problem” because they are difficult to distinguish based on local
128 information alone and must be disambiguated based on the surrounding context (27). This
129 image set thus poses a more stringent challenge of face detection ability than standard screen
130 sets which vary along many stimulus dimensions (i.e. faces vs bodies and non-face objects; see
131 **Fig. 1**). Consistent with previous work showing that neurons in pIT can be driven by images like
132 the ‘cyclops’ which contain information that is globally inconsistent with a face (28), we identified
133 13 atypical face part configurations that drove neurons to produce an early response that was
134 $\geq 90\%$ of their response to a correctly configured whole face (**Fig. 2a**). Because these images
135 drove a high feedforward response, we view them as being relatively well matched in their low-
136 level image properties and capable of activating additional processing in face responsive
137 regions of IT. These images were comparable to whole faces from our screen set in driving
138 responses with short latency (median latency: whole face= 70.0 ± 0.4 ms, typical arrangement of
139 parts= 69.5 ± 0.5 ms, atypical arrangements of parts= 70.0 ± 0.5 ms, $n_{\text{images}}=10, 8,$ and 13,
140 respectively; $p < 0.01$ for atypical versus typical and atypical versus whole face, $n=115$ sites),
141 and firing rates were comparable in strength to whole faces even over a broad analysis window
142 (median normalized response over 50-200 ms: whole face exemplars= 1.22 ± 0.23 , typical
143 arrangement= 1.36 ± 0.04 , atypical arrangements= 1.37 ± 0.03 ; $p > 0.05$ for all two-way
144 comparisons, $n=115$ sites). Thus, building on the insight from previous work that pIT does not
145 perform full face detection and is susceptible to the “aperture problem” even though it passes
146 the basic face versus non-face object test, we constructed images that allowed us to ask how a
147 difficult face detection task might be solved over time in pIT or if a solution is produced in

148 downstream areas.

149

150 **Time course of responses in posterior, central, and anterior IT for images with typical** 151 **versus atypical arrangements of face parts**

152 When we examined the dynamics of neural responses in pIT, we found that an unexpected
153 preference for atypical arrangements of face parts emerged over time (see example sites in **Fig.**
154 **2b**). Of the sites showing a significant change in their preference over time, a majority showed a
155 decreasing preference over time for typical arrangements of face parts (43 of 51 sites or 84%; p
156 < 0.01 criterion for significant change at the site level, $n = 115$ sites) (**Fig. 2c**). This surprising
157 trend -- decreasing responses for images with typical as compared to atypical arrangements of
158 face parts -- was strong enough that it reversed the small, initial preference for images of
159 normally arranged face parts over the population (median d' : 60-90 ms = 0.11 ± 0.02 vs. 100-
160 130 ms = -0.12 ± 0.03 , $p < 0.01$, $n=115$ sites), and this trend was observed in both monkeys
161 when analyzed separately ($p < 0.01$, one-tailed test for decreased d' between 60-90 ms and
162 100-130 ms in both monkeys, $n_{M1} = 43$, $n_{M2} = 72$ sites; **Fig. 3a**). Even at the individual site level,
163 a complete and rapid reversal of the expected selectivity profile of face neurons could be
164 observed (33 of 51 sites changed from face preferring to non-face preferring and only 3 of 51
165 sites changed to face preferring after being non-face preferring) (**Fig. 4a**, left). As a result, the
166 majority of pIT sites responded more strongly to atypical images over typical images in the late
167 response phase (prefer typical arrangement: 60-90 ms = 66% vs. 100-130 ms = 34%; $p < 0.01$,
168 $n = 115$) (**Fig. 4b**, light green bars).

169

170 In the anterior face-selective regions of IT which are furthest downstream of pIT and
171 reflect additional stages of feedforward processing (see block diagram in **Fig. 1**), we did not
172 observe these strong reversals in selectivity profile -- 98% of sites (39 of 40) did not change
173 their relative preference for typical vs. atypical face features ($p < 0.01$ criterion for significant
174 change at the site level). Rather, we observed a stable selectivity profile over time in aIT
175 (median d' : 60-90 ms = 0.13 ± 0.03 vs. 100-130 ms = 0.17 ± 0.03 , $p > 0.05$, $n=40$ sites) with a
176 slight but gradual accumulation (increasing preference for normal arrangements of the face
177 parts) rather than a reversal of preference (**Fig. 4a**, right). As a result, the majority of anterior
178 sites preferred images with typical arrangement of the face parts in the late phase of the
179 response (prefer typical: 60-90 ms = 78% of sites vs. 100-130 ms = 78% of sites; $p > 0.05$, $n =$
180 40 sites) despite only a minority of upstream sites in pIT preferring these images in their late
181 response (**Fig. 4b**, black bars). This suggests that spiking responses of individual aIT sites
182 resolve images as expected from a computational system whose purpose is to detect faces, as
183 previously suggested (29). Finally, in cIT whose anatomical location is intermediate to pIT and
184 aIT, we observed a selectivity profile over time that was intermediate to that of pIT and aIT
185 consistent with its position in the ventral visual hierarchy (**Fig. 4a,b**, dark green). The very
186 different patterns of dynamics in pIT, cIT, and aIT neurons are surprising given the intuition that
187 in a feedforward network selectivity for the preferred stimulus class is maintained or built
188 contrary to the complete reversal of rank-order selectivity observed in the lower stages but not
189 the highest stage of IT.

190

191 **Controls for low-level image variation and for overall activity**

192 To validate our findings against potential confounding factors, we checked the robustness of our
193 surprising observation that preference for typical over atypical arrangements of face parts
194 reverses in pIT. Since the number of parts varied between the two image classes tested, we
195 recomputed selectivity using only images with the same number of parts thus limiting
196 differences in the contrast, spatial frequency and retinal position of energy in the image (see

197 examples in **Fig. 3b**). We found that pIT face selectivity was still consistently reversed across
198 subsets of images containing a matched number of one, two, or four parts ($n=5, 30,$ and 3
199 images, respectively). Here, we quantify reversals as a decrease in d' from the early (60-90 ms
200 post-image onset) to the late (100-130 ms post-image onset) response phase where a negative
201 d' indicates a preference for atypical face part arrangements (see **SI Methods**). For all three
202 image subsets controlling the number of face parts to be one, two, or four, d' began positive on
203 average (i.e. preferring typical face part arrangements) (median d' for 60-90 ms = $0.13 \pm 0.05,$
204 $0.05 \pm 0.02,$ 0.33 ± 0.09 for one, two, and four parts) and significantly decreased in the next
205 phase of the response becoming negative on average (median d' for 100-130 ms: $-0.27 \pm 0.06,$
206 $-0.14 \pm 0.02,$ -0.04 ± 0.12 ; $p < 0.01$ for d' comparisons between 60-90 ms and 100-130 ms, $n =$
207 $115, 115, 76$ sites) (**Fig. 3b**). A similar reversal in face selectivity was observed when we re-
208 tested single part images at smaller (3°) and larger (12°) image sizes suggesting a dependence
209 on the relative configuration of the parts and not on their absolute retinal location or absolute
210 retinal size (median d' for 60-90 ms vs. 100-130 ms: three degrees = 0.51 ± 0.09 vs. $-0.29 \pm$
211 $0.14,$ twelve degrees = 0.07 ± 0.14 vs. -0.11 ± 0.14 ; $n = 15$; $p < 0.01$ for three degree condition
212 only) (**Fig. 3c**).

213
214 We also tested the hypothesis that absolute initial firing rates, which were not perfectly
215 matched, were somehow responsible for producing the pIT image preference reversal. We
216 found no support for this hypothesis -- the observed change in firing rate over time ($\Delta r_{\text{pIT}} = r_{\text{pIT late}} -$
217 $r_{\text{pIT early}}$) was weakly correlated with the strength of the initial response ($\rho_{\text{pIT early}, \Delta \text{pIT}} = -0.24 \pm$
218 $0.15,$ $p = 0.044,$ $n = 20$ images; for these firing rate controls, the original whole face image drove
219 a much higher response than the synthetic images we created, and being a firing rate outlier,
220 we excluded this image). Instead, firing rate changes over time were strongly correlated with the
221 class (typical versus atypical) of the image ($\rho_{\text{class}, \Delta \text{pIT}} = -0.77 \pm 0.04,$ $p < 0.01,$ $n = 20$ images). In
222 other words, responses to images with normally arranged face parts were specifically weaker by
223 18% on average in the next phase of the response (Δrate (60-90 vs 100-130 ms) = $-18\% \pm 4\%,$
224 $p < 0.01$; $n = 7$ images), but responses to images with unnatural arrangements of face parts,
225 which also drove high initial responses, did not experience any firing rate reduction in the next
226 phase of the response (Δrate (60-90 vs 100-130 ms) = $2 \pm 1\%,$ $p > 0.05$; $n = 13$ images). This
227 dependence on the image class and not on initial response strength argues against
228 explanations such as rate-driven adaptation that solely depend on a unit's activity to explain
229 decreasing neural responses over time. Indeed, we found that the late phase firing rates in PIT
230 could not be predicted from early phase pIT firing rates ($\rho_{\text{pIT early}, \text{pIT late}} = 0.07 \pm 0.17,$ $p > 0.05$; n
231 $= 20$ images). In contrast, we found that pIT late phase firing rates were better predicted by
232 early phase firing rates in downstream regions cIT and aIT ($\rho_{\text{cIT early}, \text{pIT late}} = -0.52 \pm 0.11,$ $p <$
233 0.01 ; $\rho_{\text{aIT early}, \text{pIT late}} = -0.36 \pm 0.14,$ $p = 0.012$; $n_{\text{pIT}}=115,$ $n_{\text{cIT}}=70,$ $n_{\text{aIT}}=40$ sites). That is, for images
234 that produced high early phase responses in cIT and aIT, the following later phase responses of
235 units in the lower level area (pIT) tended to be low, consistent with the hypothesis that feedback
236 from those areas is producing the pIT selectivity reversals. Finally, the relative speed of
237 selectivity reversals in pIT (~ 30 ms peak-to-peak) makes explanations based on fixational eye
238 movements or shifts in attention (e.g. from behavioral surprise to unnatural arrangements of
239 face parts) unlikely as saccades and attention shifts occur on slower timescales (hundreds of
240 milliseconds) (30).

241

242 **Computational models of neural dynamics in IT**

243 Given the above observations of a non-trivial, dynamic selectivity reversal during face detection,
244 we next proceeded to build formal models of gradually increasing complexity to determine the

245 minimal set of assumptions that could capture our empirical findings. We used a linear
246 dynamical systems modeling framework to evaluate dynamics in different hierarchical
247 architectures (**Supplementary Fig. 1 a,b** and see **SI Methods**). A core principle of feedforward
248 ventral stream models is that object selectivity is built by feature integration from one cortical
249 area to the next cortical area in the hierarchy leading to low dimensional representations at the
250 top of the hierarchy. Here, we take the simplest feature integration architecture where a unit in a
251 downstream area linearly sums the input from units in an upstream area to produce greater
252 downstream selectivity than any upstream input alone. This generic encoding model
253 conceptualizes the idea that different types of evidence, local (i.e. parts) and global (i.e.
254 arrangement of parts), have to converge and be integrated to separate face from non-face
255 images in our image set. Dimensionality reduction as performed in this network is a key
256 computation specified by most network architectures whether unsupervised (i.e. autoencoder)
257 or supervised (i.e. backprop) allowing these networks to learn an abstracted, low dimensional
258 representation from the high-dimensional input layer. Here, we performed dimensionality
259 reduction in linear networks as monotonic nonlinearities can be readily accommodated in our
260 framework (14)(21). First, we focused on two stage models to use the simplest configuration
261 possible and gain intuition since stacks of two processing stages can be used to generate a
262 hierarchical system of any depth. In this framing, the activity of the unit in the output stage
263 corresponds to aIT which integrates activity from units in the deepest hidden stage measured
264 corresponding to pIT (**Figure 4**, top row, first five models, and **Supplementary Fig. 1 a,b**)
265 bearing in mind that aIT is actually a hidden stage of processing with respect to the next
266 processing stage in the larger cortical stack. When an external step input is applied to such a
267 system, it will of course produce a (lagging) step response in each of the two stages. We here
268 sought to determine how adding recurrent connectivity to this basic feedforward architecture
269 could generate internal dynamics beyond those simple dynamics, and to compare those
270 dynamics with the observed IT neural dynamics.

271
272 Based on previous ideas in the literature, we considered lateral inhibition within a stage,
273 normalization within a stage (23), and cortico-cortical feedback (14). Adding recurrent lateral
274 inhibitory connections leads to competition within a stage which can limit responses over time to
275 a strong driving stimulus but not to weaker stimuli. Similarly, normalization scales down
276 responses over time to strong driving stimuli and can be implemented by a leak term that scales
277 adaptation (the degree of decay in the response) and is controlled recursively by the summed
278 activity of the network (23). Besides lateral connections and normalization within a processing
279 stage, feedback connections between stages are a remaining available mechanism for driving
280 dynamics. To constrain our choice of a feedback-based model, we took a normative approach
281 minimizing a quadratic reconstruction cost between stages. While there are many possible
282 layer-wise cost functions to consider (i.e. input to output mappings to optimize), we minimized a
283 classical reconstruction cost as this term is at the core of an array of hierarchical generative
284 models including hierarchical Bayesian inference (31), Boltzmann machines (32), analysis-by-
285 synthesis networks (14), sparse coding (20), predictive coding (21), and autoencoders in
286 general (33). Optimizing a quadratic loss results in feedforward and feedback connections that
287 are symmetric -- reducing the number of free parameters -- such that inference on the
288 represented variables at any intermediate stage is influenced by both bottom-up sensory
289 evidence and current top-down interpretations. Critically, a common feature of this large model
290 family is the computation of between-stage error signals via feedback which is distinct from
291 state-estimating model classes (i.e. feedforward models) that do not compute or propagate
292 errors (see **Figure 7** for a comparison of state and error estimating model classes). A dynamical
293 implementation of such a network uses leaky integration of error signals which, as shared

294 computational intermediates, guide gradient descent of the values of the represented variables
295 (Δ activity of each neuron \Rightarrow inference) to a previously learned target value or descend the
296 connection weights (Δ synaptic strengths \Rightarrow learning) to values that give the best behavior, here
297 defined as a reconstruction goal (similar results were found using other goals and networks;
298 **Supplementary Fig. 2**).

299
300 When we fit each of the models to our neural data, they were all able to produce an
301 increase in selectivity from the first stage of the network to the second stage of the network.
302 This increase is not surprising because all models had converging feedforward connections
303 from the first to second stages (**Figure 5a**, first five columns, compare green and black curves).
304 However, we found that neither the lateral inhibition model, nor the normalization model, could
305 capture the observed selectivity reversal phenomenon in pIT. Instead, the selectivity of these
306 models simply increased to a saturation level set by the leak term (shunting inhibition) in the
307 system (**Figure 5a**, first five columns). Similar behavior was present when we tried a nonlinear
308 implementation of the normalization model that more powerfully modulated shunting inhibition
309 (23). That the normalization models performed poorly can be explained by the fact that
310 responses to a strong stimulus even when normalized can meet but not fall below those to a
311 stimulus that was initially weak. Thus, a complete reversal in stimulus preference on average
312 across a population of cells (i.e. Figures 2-4, PIT data) is not possible when only using
313 normalization mediated by surround suppression.

314
315 In contrast to the above models, we found that the feedback model capable of computing
316 hierarchical error signals naturally displayed a strong reversal of selectivity in a sub-component
317 of its first processing stage -- qualitatively similar behavior to the selectivity reversal that we
318 observed in many pIT neural sites. Specifically, this model showed reversal dynamics in the
319 magnitude of its reconstruction error signals but not in its state signals (the states of the world
320 inferred by the model) (**Figure 5a**, compare fifth and sixth columns). These error signals
321 integrate converging state signals from two stages -- one above and one below (see **SI**
322 **Methods**). The term "error" is thus meaningful in the hidden processing stages where state
323 signals from two stages can converge. The top nodes of a hierarchy receive little descending
324 input and hence do not carry additional errors with respect to the desired computation; rather,
325 top node errors closely approximate the feedforward-driven state estimates (put another way,
326 top stages drive the formation of errors below). This behavior in the higher processing stages is
327 consistent with our observation of explicit representation of faces in aIT in all phases of the
328 response (**Fig. 4**) and with similar observations of decodable identity signals by others in all
329 phases of aIT responses for faces (34) and objects (1)(2).

330
331 Finally, we asked whether our results generalized to larger networks of increasing depth.
332 We found similar results for three layer versions of the models described above. Specifically, the
333 dynamics of error signals in a three-layer model produced a good match to our data collected
334 from three successive cortical areas (**Fig. 5a**, seventh column), while state signals in three layer
335 model networks did not produce the observed IT face selectivity dynamics (**Fig. 5b**, right set of
336 bars).

337 338 **Predictions of an error coding hierarchical model**

339 While our neural observations at multiple stages of the IT hierarchy led us to the error coding
340 hierarchical model above, a stronger test of the idea of error coding is whether it predicts other
341 IT neural phenomena. To identify stimulus regimes that would lead to insightful predictions, we
342 asked in what way would the behavior of error-estimating hierarchical models differ most from

343 the behavior of generic feedforward state-estimating models. Because our feedback-based
344 model uses feedforward inference at its core, it behaves similarly to a state-estimating
345 hierarchical feedforward model when the statistics of inputs match the learned feedforward
346 weight pattern of the network (i.e. 'natural' images drawn from everyday objects and scenes)
347 since in these settings feedforward inferences derived from the sensory data are aligned with
348 top-down expectations. Thus, predictions of feedback-based models that could distinguish them
349 from feedforward-only models are produced when the natural statistics of images are altered so
350 that they differ from the feedforward patterns previously learned by the network. We have
351 (above) considered one such type of alteration: images where local face features are present
352 but altered from their naturally occurring (i.e. statistically most likely) arrangement. Next, we
353 tested two other image manipulations from recent physiology studies which yielded novel neural
354 phenomena that lacked a principled, model-based explanation (35)(13). To test whether the
355 error coding hierarchical model family displays these behaviors, we fixed the architectural
356 parameters derived from our fitting procedure in **Figure 5** and simply varied the input to this
357 network, specifically the correlation between inputs and the network's weight pattern, in order to
358 match the nature of the image manipulations performed in prior experiments.

359
360 *Sublinear integration of the face features.* In face-selective areas in pIT and cIT, the sum of the
361 responses to the face parts exceeds the response to the whole face (35)(28), and this behavior
362 increases from linear to more sublinear over the timecourse of the response (ratio of sum of
363 responses to parts vs. response to whole: 60-90 ms = 1.5 ± 0.1 , 100-130 ms = 4.6 ± 0.3 ; $p <$
364 0.01 , $n = 33$ sites) (**Fig. 6a**, left panel). This result runs counter to what would be expected in a
365 model where selectivity for the whole face is built from the conjunction of the parts. In such a
366 model, the response to the whole face would be at least as large if not greater (superlinear) than
367 the summed responses to the individual features. To test whether an error coding model
368 exhibited the phenomenon of sublinear feature integration, we compared the response with all
369 inputs active (co-occurring features) to the sum of the responses when each input was activated
370 independently (individual features). The reconstruction errors in our feedback-based model
371 showed a strong degree of sublinear integration of the inputs such that the response to the
372 simultaneous inputs (whole) was much smaller than what would be predicted by a linear sum of
373 the responses to each input alone (parts), and the model's sublinear integration behavior
374 qualitatively replicated the time course observed in pIT without any additional fitting of
375 parameters (**Fig. 6a**, right panel).

376
377 *Evolution of neural signals across time.* Neural responses to familiar images are known to
378 rapidly attenuate in IT when compared to responses to novel images (36)(37)(38). This
379 observation seems to contradict what would be predicted by simple Hebbian potentiation for
380 more exposed stimuli. Furthermore, familiar images show much sharper temporal dynamics
381 than responses to novel images when presented repeatedly (13). These qualitatively different
382 dynamics for familiar versus novel images are surprising given that stimuli are drawn from the
383 same distribution of natural images and are thus matched in their image-level statistical
384 properties (color, spatial frequency, contrast). To test whether our network displayed these
385 different dynamical behaviors, we simulated familiar inputs as those that match the learned
386 weight pattern of a high-level detector and novel inputs as those with the same overall input
387 level but with weak correlation to the learned network weights (here, we have extended the
388 network to include two units in the output stage corresponding to storage of the two familiarized
389 input patterns to be alternated; conceptually, we consider these familiar pattern detectors as
390 existing downstream of IT in a region such as perirhinal cortex which has been shown to code
391 familiarized image statistics and memory-based object signals (39)). We repeatedly alternated

392 two familiar inputs or two novel inputs and found that model responses in the hidden processing
393 stage were temporally sharper for familiar inputs that matched the learned weight patterns
394 compared to novel, unlearned patterns of input, consistent with the previously observed
395 phenomenon (**Fig. 6b**; data reproduced with permission from Meyer et al., 2014 (13)). Model
396 responses reproduced additional details of the neural dynamics including a large initial peak
397 followed by smaller peaks for responses to novel inputs and a phase delay in the oscillations of
398 responses to novel inputs compared to familiar inputs. Intuitively, these dynamics are composed
399 of two phases. After the initial response transient, familiar patterns lead to lower errors and
400 hence lower neural responses than random patterns (see **Fig. 6b**, red curve drops below the
401 blue curve after the onset response), similar to the observed weaker response to more familiar
402 face-like images present in our data (**Fig. 2d**). When the familiar pattern *A* is switched to
403 another familiar pattern *B*, this induces a short-term error in adjusting to the new pattern (**Fig.**
404 **6b**, red curve briefly goes above the blue curve during pattern switch and then decreases).
405 Because unfamiliar patterns are closer together in the high-level encoding space than two
406 learned patterns (**Fig. 6b**, top right panel), the switch between two learned patterns introduces
407 more shift in top-down signals and hence greater change in error signals. This result
408 demonstrates that our model, derived from fitting only the first 70 ms (60-130 ms post image
409 onset) of IT responses to face images, can extend to much longer timescales and may
410 generalize to studies of images besides face images.

411

412 **Dynamical properties of neurons across cortical lamina**

413 In the large family of state-error coding hierarchical networks, a number of different cortical
414 circuits are possible (**Fig. 7**). A key distinction of two such circuit mapping hypotheses
415 (predictive coding versus error backpropagation) is the expected laminar location of state coding
416 neurons. Specifically, predictive coding asserts that superficial layers contain error units and
417 that errors are projected forward to the next cortical level (21)(40)(41) as opposed to typical
418 neural network implementations where the feedforward projecting neurons in superficial lamina
419 are presumed to encode estimates about states of the visual world (e.g. is a face present or
420 not?). State and error signals can be distinguished by their dynamical signatures in our leading
421 model, which was fit on error signals but produces predictions of the corresponding state signals
422 underlying the generation of errors. Since state units are integrators (see **SI Methods**), they
423 have slower dynamics than error units leading to longer response latencies and a milder decay
424 in responses (**Fig. 6a**, right panel). To test this prediction, we localized our recordings relative to
425 the cortical mantle by co-registering the x-ray determined locations of our electrode (~400
426 micron *in vivo* accuracy) to structural MRI data (see **SI Methods**). When we separated units into
427 those at superficial depths closer to the pial surface (1/3 of our sites; corresponds to
428 approximately 0 to 1 mm in depth) versus those in the deeper layers (remaining 2/3 of sites, ~1
429 to 2.5 mm in depth), we found a longer latency and less response decay in superficial units
430 consistent with the expected profile of state units (**Fig. 6a**, left panel). Importantly, the latency
431 difference between cortical lamina within pIT (deep vs superficial: 66.0 ± 1.7 vs 76.0 ± 1.7 ms, p
432 < 0.01) was greater than the conduction delay from pIT to cIT (i.e. from superficial layers of pIT
433 to the deeper layers of cIT) (superficial pIT vs deep cIT: 76.0 ± 1.7 vs 75.5 ± 2.1 ms, $p > 0.05$)
434 even though the distance traveled between cortical stages pIT and cIT is larger than laminar
435 distances within pIT. Thus, instead of a simple conduction delay accounting for latency
436 differences across lamina, our model suggests that temporal integration of inputs, consistent
437 with the behavior of state units implemented in standard feature coding rather than predictive
438 coding schemes, may drive the lagged dynamical properties of neurons in superficial lamina.

439

440

441

442 **DISCUSSION**

443 We have measured neural responses during a difficult face detection task across the IT
444 hierarchy and demonstrated that the initial feedforward preference for faces in the intermediate
445 (a.k.a hidden) processing stages reverses over time – that is, after the initial wave of face-
446 selective neural responses, responses at lower levels of the hierarchy (pIT and cIT) rapidly
447 evolve to not prefer typical face part arrangements. This behavior was inconsistent with a pure
448 feedforward model, even when we included strong nonlinearities in these models, such as
449 normalization. However, we showed that augmenting the feedforward model so that it
450 represents the errors generated during hierarchical processing produced the observed neural
451 dynamics (**Fig. 5**). This view argues that a fraction of cortical neurons codes error signals. Using
452 this new modeling perspective, we went on to generate a series of predictions consistent with
453 observed IT neural phenomena (**Fig. 6**). Importantly, this perspective provides an alternative
454 interpretation to prior suggestions that IT neurons are not tuned for typical faces but are instead
455 tuned for atypical faces (35). Under the present hypothesis, some IT neurons are preferentially
456 tuned to typical arrangements of face features, and many other IT neurons are involved in
457 coding errors with respect to those typical arrangements. We believe that these intermixed state
458 estimating and error coding neuron populations are both sampled in standard neural recordings
459 of IT, even though only state estimating neurons are truly reflective of the tuning preferences of
460 that IT processing stage.

461

462 The precise fractional contribution of errors to neural activity is difficult to estimate from
463 our data. Under the primary image condition tested, not all sites significantly decreased their
464 selectivity (~60%). We currently interpret these sites as coding state (feature) estimates (**Fig.**
465 **2c**, gray and black dots), and we did observe evidence of emergence of state-like signals in our
466 superficial neural recordings (**Fig. 6c**). Alternatively, at least some of the non-reversing sites
467 might be found to code errors under other image conditions than the one that we tested.
468 Furthermore, while in our primary image condition selectivity reversals only accounted for 20%
469 of the overall spiking modulation (**Fig. 2d**), we found larger modulations in late phase neural
470 firing (50-100%) under other image conditions tested (**Fig. 6 a,b**). At a computational level, the
471 absolute contribution of error signals to spiking may not be the critical factor as even a small
472 relative contribution may have important consequences in the network.

473

474 **Error signals generated across different hierarchical inference and learning models**

475 The notion of error is inherent to many existing models in the literature that go beyond the basic
476 feedforward, feature estimation class. These models use errors for guiding top-down inference
477 by computing errors implicitly (hierarchical Bayesian inference (14)(31); **Fig. 7**, second row) or
478 by representing errors explicitly (predictive coding (21); **Fig. 7**, third and last row). Alternatively,
479 errors can be used specifically for unsupervised learning (autoencoder (33); **Fig. 7**, fourth row)
480 or specifically for supervised learning (classic error backpropagation (19); **Fig. 7**, fifth row).
481 Finally, recent models incorporate aspects of both inference and learning (42)(43) (**Fig. 7**;
482 bottom two rows). A key, unifying feature across inference and learning models is the need to
483 compute an error signal between processing stages. This error signal can be in the form of a
484 generative, reconstruction cost (stage n predicting stage $n-1$) or a discriminative, construction
485 cost (stage $n-1$ predicting stage n). Regardless, this across-stage “performance” error term is
486 used in all models, is typically the only term combining signals from different model layers, and
487 is distinct from within-stage “regularization” terms (i.e. sparseness or weight decay) in driving
488 network behavior. The present study provides evidence that such errors are not only computed,
489 but that they are explicitly encoded in spiking rates. To test the robustness of this claim across

490 different model implementations, we tested models with different performance errors
491 (reconstruction, nonlinear reconstruction, and discriminative) and found similar population level
492 error signals across these networks (**Supplementary Fig. 2**). Thus, errors as broadly construed
493 in the state-error coding hierarchical model family provide a good approximation to IT population
494 neural dynamics, and future work examining the specifics of error signals and interactions
495 between error and state units at the single-neuron level may distinguish among the various
496 computational algorithms which depend on error signals for their implementation.

497

498 **Comparison to previous neurophysiology studies in the ventral stream**

499 Prior work in IT has shown that responses of face-selective cells are stronger for atypical faces
500 than for typical faces and are stronger when the parts are presented individually than when
501 presented together in a whole face (35)(28). One possible interpretation of these data is that IT
502 neurons are not tuned for faces but are instead tuned for atypical face features (i.e. extreme
503 feature tuning) (35); however, our data suggest an alternative interpretation of this finding and,
504 more deeply, an extended computational purpose of IT dynamics. In that prior work, the
505 response preference of each neuron was determined by averaging over a long time window
506 (~200 ms). By looking more closely at the fine time scale dynamics of the IT response, we
507 suggest that this same “extreme coding” phenomenon can instead be interpreted as a natural
508 consequence of networks that have an actual tuning preference for typical faces (as evidenced
509 by an initial response preference for typical faces in pIT, cIT, and aIT; **Fig. 4b**) but that also
510 compute error signals with respect to that preference. The hierarchical error coding framework
511 proposed here provides a single, unifying account of many other reliable but previously
512 unexplained phenomena in IT: sublinear integration of multiple inputs (35)(28) (**Fig. 6a**), tuning
513 for extreme features in faces (35) (**Fig. 2**; positional shifts of the eye position correspond to
514 extremes of the face space used in Freiwald et al., 2009), tuning for novel stimuli over familiar
515 stimuli (36)(38)(37) (**Fig. 6b**, response to first presentation is larger for the novel inputs), and
516 rapid response dynamics for familiar over novel images (13) (**Fig. 6b**, larger oscillation
517 amplitude to repeated presentation of familiar inputs). Thus, we have provided a parsimonious
518 framework that can account for these disparate neural phenomena by naturally extending the
519 purely feedforward model in a way suggested by prior computational work. The perspective that
520 many IT neurons code error signals reflecting deviations from the naturally learned statistics of
521 images suggests that natural joint statistics of features will lead to suppressed responses on
522 average in the hidden processing stages. This perspective may extend across the ventral visual
523 stream including V1 where there is causal evidence of a suppressive role for feedback in
524 producing end-stopping (44), and it has been suggested that end-stopping is the result of error-
525 like computations (21).

526

527 **Computational utility of coding errors in addition to states**

528 In error-computing networks, errors provide control signals for guiding learning giving these
529 networks additional adaptive power over basic feature estimation networks. This property helps
530 augment the classical, feature coding view of neurons which, with only feature activations and
531 Hebbian operations, does not lead to efficient gradient descent and may provide insight into
532 how more intelligent unsupervised and supervised learning algorithms such as backpropagation
533 could be plausibly implemented in the brain. A potentially important contribution of this work is
534 the suggestion that gradient descent algorithms are facilitated by using an error code so that
535 efficient learning is reduced to a simple Hebbian operation at synapses and efficient inference is
536 simply integration of inputs at the cell body. This representational choice, to code the
537 computational primitives of gradient descent in spiking activity, simply leverages existing neural
538 machinery for inference and learning.

539
540 While we provide evidence that IT hidden units code error signals, the precise
541 computational use of those error signals remains to be empirically determined. The most likely
542 downstream, causal impact of error signals could be in online inference (updating neural firing
543 rates), offline inference (updating synaptic weights, a.k.a. learning), or both. We expect that
544 optimizing the weights and optimizing the states (feature estimates) are both implemented in
545 cortical circuits as there is a large body of evidence on inference and learning in cortex including
546 work in IT showing unsupervised learning of novel images within a few hundred presentations
547 (37)(38)(45), work in IT showing neural changes after visual discrimination training (46), and
548 work in IT showing that neural responses evolve over short timescales to produce improved
549 estimates online (10)(11)(47). If both online and offline optimization mechanisms are used, their
550 impact on the animal's behavioral output could ground their relative importance. Conversely, the
551 animal's engagement in a task may guide the computation of error signals, through addition of a
552 top-down supervision term as in classical error backpropagation, for improved performance
553 under a new behavioral goal. As new tools become available to specifically disrupt cortico-
554 cortical feedback, they can be brought to bear on these key questions in hierarchical cortical
555 computation.

556
557

558 MATERIALS & METHODS

559
560 **Animals and surgery.** All surgery, behavioral training, imaging, and neurophysiological
561 techniques are identical to those described in detail in our previous work²⁸. Two rhesus
562 macaque monkeys (*Macaca mulatta*) weighing 6 kg (Monkey 1, female) and 7 kg (Monkey 2,
563 male) were used. A surgery using sterile technique was performed to implant a plastic fMRI
564 compatible headpost prior to behavioral training and scanning. Following scanning, a second
565 surgery was performed to implant a plastic chamber positioned to allow targeting of
566 physiological recordings to posterior, middle, and anterior face patches in both animals. All
567 procedures were performed in compliance with National Institutes of Health guidelines and the
568 standards of the MIT Committee on Animal Care and the American Physiological Society.

569
570 **Behavioral training and image presentation.** Subjects were trained to passively fixate a
571 central white fixation dot during serial visual presentation of images at a natural saccade-driven
572 rate (one image every 200 ms). Although a 4° fixation window was enforced, subjects generally
573 fixated a much smaller region of the image (<1°)²⁸. Images were presented at a size of 6° except
574 for control tests at 3° and 12° sizes (**Fig. 3c**), and all images were presented for 100 ms duration
575 with 100 ms gap (background gray screen) between each image. Up to 15 images were
576 presented during a single fixation trial, and the first image presentation in each trial was
577 discarded from later analyses. Five repetitions of each image in the general screen set were
578 presented, and ten repetitions of each image were collected for all other image sets. The screen
579 set consisted of a total of 40 images drawn from four categories (faces, bodies, objects, and
580 places; 10 exemplars each) and was used to derive a measure of face versus nonface object
581 selectivity. Following the screen set testing, some sites were tested using an image set
582 containing images of face parts presented in different combinations and positions. We first
583 segmented the face parts (eye, nose, mouth) from a monkey face image. These parts were then
584 blended using a Gaussian window, and the face outline was filled with pink noise to create a
585 continuous background texture. A face part could appear on the outline at any one of nine
586 positions on an evenly spaced 3x3 grid. Although the number of possible images is large ($4^9 =$
587 262,144 images), we chose a subset of these images for testing neural sites (n=82 images).

588 Specifically, we tested the following images: the original whole face image, the noise-filled
589 outline, the whole face reconstructed by blending the four face parts with the outline, all possible
590 single part images where the eye, nose, or mouth could be at one of nine positions on the
591 outline ($n=3 \times 9=27$ images), all two part images containing a nose, mouth, left eye, or right eye
592 at the correct outline-centered position and an eye tested at all remaining positions ($n=4 \times 8-$
593 $1=31$ images), all two part images containing a correctly positioned contralateral eye while
594 placing the nose or mouth at all other positions ($n=2 \times 8-2=14$ images), and all correctly
595 configured faces but with one or two parts missing ($n=3+4=7$ images). The particular two-part
596 combinations tested were motivated by prior work demonstrating the importance of the eye in
597 early face processing²⁸, and we sought to determine how the position of the eye relative to the
598 outline and other face parts was encoded in neural responses. The three and four part
599 combinations were designed to manipulate the presence or absence of a face part for testing
600 the integration of face parts, and in these images, we did not vary the positions of the parts from
601 those in a naturally occurring face. In a follow-up test on a subset of sites, we permuted the
602 position of the four face parts under the constraint that they still formed the configuration of a
603 naturally occurring face (i.e. preserve the 'T' configuration, $n=10$ images; **Fig. 3b**). We tested
604 single part images at 3° and 12° sizes in a subset of sites ($n=27$ images at each size; **Fig. 3c**).
605 Finally, we measured the responses to the individual face parts in the absence of the outline
606 ($n=4$ images; **Fig. 6a**).
607

608 **MR Imaging and neurophysiological recordings.** Both structural and functional MRI scans
609 were collected in each monkey. Putative face patches were identified in fMRI maps of face
610 versus object selectivity in each subject. A stereo microfocal x-ray system²⁴ was used to guide
611 electrode penetrations in and around the fMRI defined face-selective subregions of IT. X-ray
612 based electrode localization was critical for making laminar assignments since electrode
613 penetrations are often not perpendicular to the cortical lamina when taking a dorsal-ventral
614 approach to IT face patches. Laminar assignments of recordings were made by co-registering x-
615 ray determined electrode coordinates to MRI where the pial-to-gray matter border and the gray-
616 to-white matter border were defined; based on our prior work estimating sources of error (e.g.
617 error from electrode tip localization and brain movement), registration of electrode tip locations
618 to MRI brain volumes has a total of <400 micron error which is sufficient to distinguish deep
619 from superficial layers⁴⁸. Multi-unit activity (MUA) was systematically recorded at 300 micron
620 intervals starting from penetration of the superior temporal sulcus such that all sites were tested
621 with a screen set containing both faces and nonface objects, and a subset of sites that were
622 visually driven were further tested with our main image set manipulating the position of face
623 parts. Although we did not record single-unit activity, our previous work showed similar
624 responses between single-units and multi-units on images of the type presented here²⁸, and our
625 results are consistent with observations in previous single-unit work in IT³⁵. Recordings were
626 made from PL, ML, and AM in the left hemisphere of monkeys 1 and 2 and from AL in monkey
627 2. AM and AL are pooled together in our analyses forming the aIT sample.
628

629 **Neural data analysis.** The face patches were physiologically defined in the same manner as in
630 our previous study²⁸. Briefly, we fit a graded 3D sphere model (linear profile of selectivity that
631 rises from a baseline value toward the maximum at the center of the sphere) to the spatial
632 profile of face versus nonface object selectivity across our sites. We tested spherical regions
633 with radii from 1.5 to 10 mm and center positions within a 5 mm radius of the fMRI-based
634 centers of the face patches. The resulting physiologically defined regions were 1.5 to 3 mm in
635 diameter. Sites which passed a visual response screen (mean response in a 60-160 ms window
636 $>2 \times \text{SEM}$ above baseline for at least one of the four categories in the screen set) were included

637 in further analysis. All firing rates were baseline subtracted using the activity in a 25-50 ms
638 window following image onset averaged across all repetitions of an image. Finally, given that
639 the visual response latencies in monkey 2 were on average 13 ms slower than those in monkey
640 1, we applied a single latency correction (13 ms shift to align monkey 1 and monkey 2's data)
641 prior to averaging across monkeys. This was done to so as not to wash out any fine timescale
642 dynamics by averaging though similar results were obtained without using this latency
643 correction, and this single absolute adjustment was more straightforward than the site-by-site
644 adjustment used in our previous work (similar results were obtained using this alternative
645 latency correction)²⁸. The observed selectivity dynamics (**Fig. 2**) were found in each monkey
646 analyzed separately (**Fig. 3a**). Images that produced an average population response ≥ 0.9 of
647 the initial response (60-100 ms) to a face-like image were analyzed further (**Figs. 2-4**). In follow-
648 up analyses, we specifically limited comparison to images with the same number of parts (**Fig.**
649 **3b**). For example, for single part images, we used the image with the eye in the upper,
650 contralateral region of the outline as a reference and found that four other images of the 27
651 single-part images elicited a response at least as large as 90% of the response to this standard
652 image. For images containing all four face parts, we used the whole face as the standard and
653 found nonface-like arrangements of the four face parts that drove at least 90% of the early
654 response to the whole face (2 images out of 10 tested). Individual site d' measures were
655 computed using $d' = (u_1 - u_2) / ((var_1 + var_2) / 2)^{1/2}$ where variance was computed across all trials for
656 that image class (i.e. all presentations of all typical face images). A positive d' implies a stronger
657 response to more naturally occurring (typical) arrangements of face parts while a negative d'
658 indicates a preference for unnatural (atypical) arrangements of the face parts.

659

660 **Dynamical models**

661 *Modeling framework and equations.* To model the dynamics of neural response rates in a
662 hierarchy, we started with the simplest possible model that might capture those dynamics: we
663 used a model architecture consisting of a hidden stage of processing containing two units that
664 linearly converged onto a single output unit. An external input was applied separately to each
665 hidden stage unit, which can be viewed as representing different features for downstream
666 integration. We varied the connections between the two hidden units within the hidden
667 processing stage (lateral connections) or between hidden and output stage units (feedforward
668 and feedback connections) to instantiate different model families. The details of the different
669 architectures specified by each model class can be visualized by their equivalent neural network
670 diagrams and connection matrices (**Supplementary Fig. 1**). Here, we provide a basic
671 description for each model tested. All models utilize a 2x2 feedforward identity matrix A that
672 simply transfers inputs \mathbf{u} (2x1) to hidden layer units \mathbf{x} (2x1) and a 1x2 feedforward matrix B that
673 integrates hidden layer activations \mathbf{x} into a single output unit y .

674

$$675 \quad A = k_a \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}, \quad B = k_b [1 \quad 1] \quad (1)$$

676

677 To generate dynamics in the simple networks below, we assumed that neurons act as leaky
678 integrators of their total synaptic input, a standard rate-based model of a neuron used in
679 previous work^{14,21}.

680

681 *Pure feedforward.* In the purely feedforward family, connections were exclusively from hidden to
682 output stages through feedforward matrices A and B .

683

684 $\dot{\mathbf{x}} = A\mathbf{u} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - y/\tau$ (2)
 685

686 where τ is the time constant of the leak current which can be seen as reflecting the biophysical
 687 limitations of neurons (a perfect integrator with large τ would have almost no leak and hence
 688 infinite memory).
 689

690 *Lateral inhibition.* Lateral connections (matrix with off-diagonal terms) are included and are
 691 inhibitory. The scalar k_l sets the relative strength of lateral inhibition versus bottom-up input.
 692

693 $\dot{\mathbf{x}} = A\mathbf{u} - \begin{bmatrix} 0 & k_l \\ k_l & 0 \end{bmatrix} \mathbf{x} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - y/\tau$ (3)
 694

695 *Normalization.* An inhibitory term that scales with the summed activity of units within a stage is
 696 included. The scalar k_s sets the relative strength of normalization versus bottom-up input.
 697

698 $\dot{\mathbf{x}} = A\mathbf{u} - k_s \sum x \cdot \mathbf{x} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - k_s y \cdot y - y/\tau$ (4)
 699

700 *Normalization (nonlinear)*²³. The summed activity of units within a stage is used to nonlinearly
 701 scale shunting inhibition.
 702

703 $\dot{\mathbf{x}} = A\mathbf{u} - \frac{\mathbf{x}}{\tau \sqrt{1 - k_s \sum x}} \quad \dot{y} = B\mathbf{x} - \frac{y}{\tau \sqrt{1 - k_s y}}$ (5)
 704

705 Since the normalization term in equation (5) is not continuously differentiable, we used the
 706 fourth-order Taylor approximation around zero in the simulations of equation (5).
 707

708 *Feedback (linear reconstruction).* The feedback-based model is derived using a normative
 709 framework that performs optimal inference in the linear case¹⁴ (unlike the networks in equations
 710 (2)-(5) which are motivated from a mechanistic perspective but do not directly optimize a
 711 squared error performance loss). The feedback network minimizes the cost C of reconstructing
 712 the inputs of each stage (i.e. mean squared error of layer n predicting layer $n-1$).
 713

714 $C = \frac{1}{2} (\mathbf{u} - A^T \mathbf{x})^2 + \frac{1}{2} (\mathbf{x} - B^T y)^2$ (6)
 715

716 Differentiating this coding cost with respect to the encoding variables in each layer \mathbf{x} , y yields:
 717

718 $\frac{\partial C}{\partial \mathbf{x}} = -A(\mathbf{u} - A^T \mathbf{x}) + (\mathbf{x} - B^T y) \quad \frac{\partial C}{\partial y} = -B(\mathbf{x} - B^T y)$ (7)
 719

720 The cost function C can be minimized by descending these gradients over time to optimize the
 721 values of \mathbf{x} and y .
 722

723 $\frac{d\mathbf{x}}{dt} = -\frac{\partial C}{\partial \mathbf{x}} = A(\mathbf{u} - A^T \mathbf{x}) - (\mathbf{x} - B^T y) - \mathbf{x}/\tau$ (8)
 724

$$\frac{d\mathbf{y}}{dt} = -\frac{\partial C}{\partial \mathbf{y}} = B(\mathbf{x} - B^T \mathbf{y}) - \mathbf{y}/\tau$$

725
726

727 The above dynamical equations are equivalent to a linear network with a connection matrix
728 containing symmetric feedforward (B) and feedback (B^T) weights between stages \mathbf{x} and \mathbf{y} as
729 well as within-stage pooling followed by recurrent inhibition ($-AA^T \mathbf{x}$ and $-BB^T \mathbf{y}$) that resembles
730 normalization. The property that symmetric connections minimize the cost function C
731 generalizes to a feedforward network of any size or number of hidden processing stages (i.e.
732 holds for arbitrary lower triangular connection matrix B). The final activation states (\mathbf{x}, \mathbf{y}) of the
733 hierarchical generative network are optimal in the sense that the bottom-up activations
734 (implemented through feedforward connections) are balanced by the top-down expectations
735 (implemented by feedback connections) which is equivalent to a Bayesian network combining
736 bottom-up likelihoods with top-down priors to compute the maximum *a posteriori* (MAP)
737 estimate. Here, the priors are embedded in the weight structure of the network. In simulations,
738 we include an additional scalar k_{td} that sets the relative weighting of bottom-up versus top-down
739 signals.

740

$$\dot{\mathbf{x}} = A(\mathbf{u} - A^T \mathbf{x}) - k_{td}(\mathbf{x} - B^T \mathbf{y}) - \mathbf{x}/\tau \quad (9)$$

741
742

743 *Error signals computed in the feedback model.* In equation (9), inference can be thought of as
744 proceeding through integration of inputs on the dendrites of neuron population \mathbf{x} . In this
745 scenario, all computations are implicit in dendritic integration. Alternatively, the computations in
746 equation (9) can be done in two steps where, in the first step, reconstruction errors are
747 computed (i.e. $\mathbf{e}_0 = \mathbf{u} - A^T \mathbf{x}$, $\mathbf{e}_1 = \mathbf{x} - B^T \mathbf{y}$) and explicitly represented in a separate error coding
748 population. These error signals can then be integrated to generate the requisite update to the
749 state signal of neuron population \mathbf{x} .

750

$$\dot{\mathbf{x}} = A\mathbf{e}_0 - k_{td}\mathbf{e}_1 - \mathbf{x}/\tau \quad \dot{\mathbf{y}} = B\mathbf{e}_1 - \mathbf{y}/\tau \quad (10)$$

751
752

753 An advantage of this strategy is that there are now only two input populations to a state unit,
754 and those inputs allow implementation of an efficient Hebbian rule for learning weight matrices²¹
755 -- the gradient rule for learning is simply a product of the state activation and the input error
756 activation (weight updates obtained by differentiating equation (6) with respect to weight
757 matrices A and B : $\Delta A = \mathbf{x} \cdot \mathbf{e}_0^T$, $\Delta A^T = \mathbf{e}_0 \cdot \mathbf{x}^T$, $\Delta B = \mathbf{y} \cdot \mathbf{e}_1^T$, and $\Delta B^T = \mathbf{e}_1 \cdot \mathbf{y}$). Thus, the reconstruction
758 errors serve as computational intermediates for both the gradients of online inference (dynamics
759 in state space, equation (10)) and gradients for offline learning (dynamics in weight space).

760

761 In order for the reconstruction errors at each layer to be scaled appropriately in the
762 feedback model, we invoke an additional downstream variable z to predict activity at the top
763 layer such that, instead of $\mathbf{e}_2 = \mathbf{y}$ which scales as a state variable, we have $\mathbf{e}_2 = \mathbf{y} - C^T z$
764 (**Supplementary Fig. 1a**). This overall model reflects a state and error coding model as
765 opposed to a state only model. A third, less plausible possibility is to only represent error signals
766 explicitly (a.k.a. pure predictive coding; **Fig. 7** (iii)). In other words, the linear dynamical system
767 in equation (8) can be rewritten through a linear change of variables to include error variables
768 only, and the state variables become implicitly represented during dendritic integration of these
769 errors.

770

771 *Feedback (three-stage).* For the simulations in **Figs. 5,6**, a three-stage version of the above

772 models was used. These deeper network were also wider such that they began with four input
 773 units (\mathbf{u}) instead of only two inputs in the two-stage models. These inputs converged through
 774 successive processing stages ($\mathbf{w}, \mathbf{x}, \mathbf{y}$) to one unit at the top node (\mathbf{z}) (**Supplementary Fig. 1b**).

775
 776 *Feedback (nonlinear reconstruction)*. We tested versions of feedback-based models that
 777 optimized different cost functions other than a linear reconstruction cost (**Supplementary Fig.**
 778 **2**). In nonlinear hierarchical inference, reconstruction is performed using a monotonic
 779 nonlinearity with a threshold (th) and bias (bi):

$$780$$

$$781$$

$$782 \quad C = \frac{1}{2} (\mathbf{u} - f(A^T \mathbf{x}))^2 + \frac{1}{2} (\mathbf{x} - f(B^T \mathbf{y}))^2, \quad \text{where } f(x) = \tanh(x - th) + bi \quad (11)$$

$$783$$

$$784 \quad \dot{\mathbf{x}} = A(\mathbf{u} - f(A^T \mathbf{x}))(1 - \tanh(A^T \mathbf{x} - th)^2) - k_{td}(\mathbf{x} - f(B^T \mathbf{y})) - \mathbf{x}/\tau$$

$$785 \quad (12)$$

$$786 \quad \dot{\mathbf{y}} = B(\mathbf{x} - f(B^T \mathbf{y}))(1 - \tanh(B^T \mathbf{y} - th)^2) - \mathbf{y}/\tau$$

$$787$$

788 *Feedback (linear construction)*. Instead of a reconstruction cost, we additionally simulated the
 789 states and errors in a feedback network minimizing a linear construction cost:

$$790$$

$$791 \quad C = \frac{1}{2} (A\mathbf{u} - \mathbf{x})^2 + \frac{1}{2} (B\mathbf{x} - \mathbf{y})^2 \quad (13)$$

$$792$$

$$793 \quad \dot{\mathbf{x}} = (A\mathbf{u} - \mathbf{x}) - k_{td}B^T(B\mathbf{x} - \mathbf{y}) - \mathbf{x}/\tau \quad \dot{\mathbf{y}} = (B\mathbf{x} - \mathbf{y}) - \mathbf{y}/\tau \quad (14)$$

$$794$$

795 **Model simulation.** To simulate the dynamical systems in equations (2)-(14), a step input \mathbf{u} was
 796 applied. This input was smoothed using a Gaussian kernel to approximate the lowpass nature of
 797 signal propagation in the series of processing stages from the retina to pIT:

$$798$$

$$799 \quad \mathbf{u}(t) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(t-t_0)^2}{2\sigma^2}} * \mathbf{h}(t) \Rightarrow \dot{\mathbf{u}} = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(t-t_0)^2}{2\sigma^2}} \cdot \mathbf{h} \quad (15)$$

$$800$$

801 where the elements of \mathbf{h} are scaled Heaviside step functions. The input is thus a sigmoidal ramp
 802 whose latency to half height is set by t_0 and rise time is set by σ . For simulation of two-stage
 803 models, there were ten basic parameters: latency of the input t_0 , standard deviation of the
 804 Gaussian ramp σ , system time constant τ , input connection strength A , feedforward connection
 805 strength B , the four input values across two stimulus conditions (i.e. $h_{11}, h_{12}, h_{21}, h_{22}$), and a
 806 factor sc for scaling the final output to the neural activity. In the deeper three-stage network,
 807 there were a total of fifteen parameters which included an additional feedforward connection
 808 strength C and additional input values since the three-stage model had four inputs instead of
 809 two. The lateral inhibition model class required one additional parameter k_l as did the
 810 normalization model family k_s , and for feedback model simulations, there was an additional
 811 feedback weight k_{td} to scale the relative contribution of the top-down errors in driving online
 812 inference. For the error coding variants of the feedback model, gain parameters C (two-stage)
 813 and D (three-stage) were included to scale the overall magnitude of the top level reconstruction
 814 error.

815

816 **Model parameter fits to neural data.** In fitting the models to the observed neural dynamics, we
817 mapped the summed activity in the hidden stage (\mathbf{x}) to population averaged activity in pIT, and
818 we mapped the summed activity in the output stage (\mathbf{y}) to population averaged signals
819 measured in aIT. To simulate error coding, we mapped the reconstruction errors $\mathbf{e}_1 = \mathbf{x} - B^T \mathbf{y}$ and
820 $\mathbf{e}_2 = \mathbf{y} - C^T \mathbf{z}$ to activity in pIT and aIT, respectively. We applied a squaring nonlinearity to the model
821 outputs as an approximation to rectification since recorded extracellular firing rates are non-
822 negative (and linear rectification is not continuously differentiable). Analytically solving this
823 system of dynamical equations (2)-(14) for a step input is precluded because of the higher order
824 interaction terms (the roots of the determinant and hence the eigenvalues/eigenvectors of a 3x3
825 matrix are not analytically determined, except for the purely feedforward model which only has
826 first-order interactions), and in the case of the normalization models, there is an additional
827 nonlinear dependence on the shunt term. Thus, we relied on computational methods
828 (constrained nonlinear optimization) to fit the parameters of the dynamical systems to the neural
829 data with a quadratic (sum of squares) loss function.

830
831 Parameter values were fit in a two step procedure. In the first step, we fit only the
832 difference in response between image classes (differential mode which is the selectivity profile
833 over time, see **Fig. 5b**, left panel), and in the second step, we refined fits to capture an equally
834 weighted average of the differential mode and the common mode (the common mode is the
835 average across images of the response time course of visual drive). This two-step procedure
836 was used to ensure that each model had the best chance of fitting the dynamics of selectivity
837 (differential mode) as these selectivity profiles were the main phenomena of interest but were
838 smaller in size (20% of response) compared to overall visual drive. In each step, fits were done
839 using a large-scale algorithm (interior-point) to optimize coarsely, and the resulting solution was
840 used as the initial condition for a medium-scale algorithm (sequential quadratic programming)
841 for additional refinement. The lower and upper parameter bounds tested were: $t_0 = [50 \ 70]$, $\sigma = [0.5$
842 $25]$, $\tau = [0.5 \ 1000]$, $k_l, k_s, k_{td} = [0 \ 1]$, $A, B, C, D = [0 \ 2]$, $h = [0 \ 20]$, $sc = [0 \ 100]$, $th = [-20 \ 20]$, and $bi = [-1 \ 1]$
843 which proved to be adequately liberal as parameter values converged to values that did not
844 generally approach these boundaries. To avoid local minima, the algorithm was initialized to a
845 number of randomly selected points ($n=50$), and after fitting the differential mode, we took the
846 top fits ($n=25$) for each model class and used these as initializations in subsequent steps. The
847 single best fitting instance of each model class is shown in the main figures.

848
849 **Model predictions.** For the predictions in **Fig. 6**, all architectural parameters obtained by the
850 fitting procedure above were held fixed; only the pattern of inputs to the network was varied. For
851 **Fig. 6a**, to test the input integration properties of a model, we used the top-performing model
852 (most optimal solution) and compared the response to all inputs presented simultaneously with
853 the sum of the responses to each input alone.

854
855 For **Fig. 6b**, we approximated novel versus familiar images as random patterns versus
856 structured input patterns that matched the learned weights of the network. Here, we used a
857 version of the model with two independent outputs reflecting detectors for two familiarized input
858 patterns (output 1 tuned to pattern A: u_1, u_2, u_3, u_4 active and output 2 tuned to pattern 2: $u_5, u_6,$
859 u_7, u_8 active) (**Fig. 6b**). Alternating between these two input patterns simulates alternation of two
860 familiarized (learned) images as compared to purely random patterns (u_{1-8} independent and
861 identically distributed). To parametrically vary the degree of correlation of inputs to the learned
862 weight patterns from random (correlation = 0) to deterministic (correlation = 1), we drew input
863 values from a joint distribution $P(u_1, u_2, u_3, u_4, u_5, u_6, u_7, u_8)$ where u_{1-4} were drawn from a high-

864 valued uniform distribution on the interval $[1-\varepsilon, 1]$ and u_{5-8} were drawn from a low-valued uniform
865 distribution $[0, \varepsilon]$ for stimulus pattern A and the opposite for pattern B (u_{5-8} high-valued and u_{1-4}
866 low-valued). The parameter ε determines the range of values that could be drawn from purely
867 deterministic (0 or 1) to randomly uniformly distributed (from 0 to 1). Thus, the correlation of the
868 inputs correspondingly varies according to $\rho(u_i, u_j) = \rho(u_k, u_l) = (\varepsilon-1)^2 / ((\varepsilon-1)^2 + \varepsilon^2/3)$ where $1 \leq i, j \leq 4$, $i \neq j$
869 and $5 \leq k, l \leq 8$, $k \neq l$ approaching correlation equal to 0 for a purely, random pattern ($\varepsilon=1$) that had a
870 low probability of matching the learned patterns A and B .

871

872 **Code availability.** All data analysis and computational modeling were done using custom
873 scripts written in Matlab. All code is available upon request.

874

875 **Statistics.** Error bars represent standard errors of the mean obtained by bootstrap resampling
876 ($n = 1000$). All statistical comparisons including those of means or correlation values were
877 based on 99% confidence intervals obtained by bootstrap resampling ($n = 1000$). All statistical
878 tests were two-sided unless otherwise specified. Spearman's rank correlation coefficient was
879 used.

880

881

882

883

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Figure 1

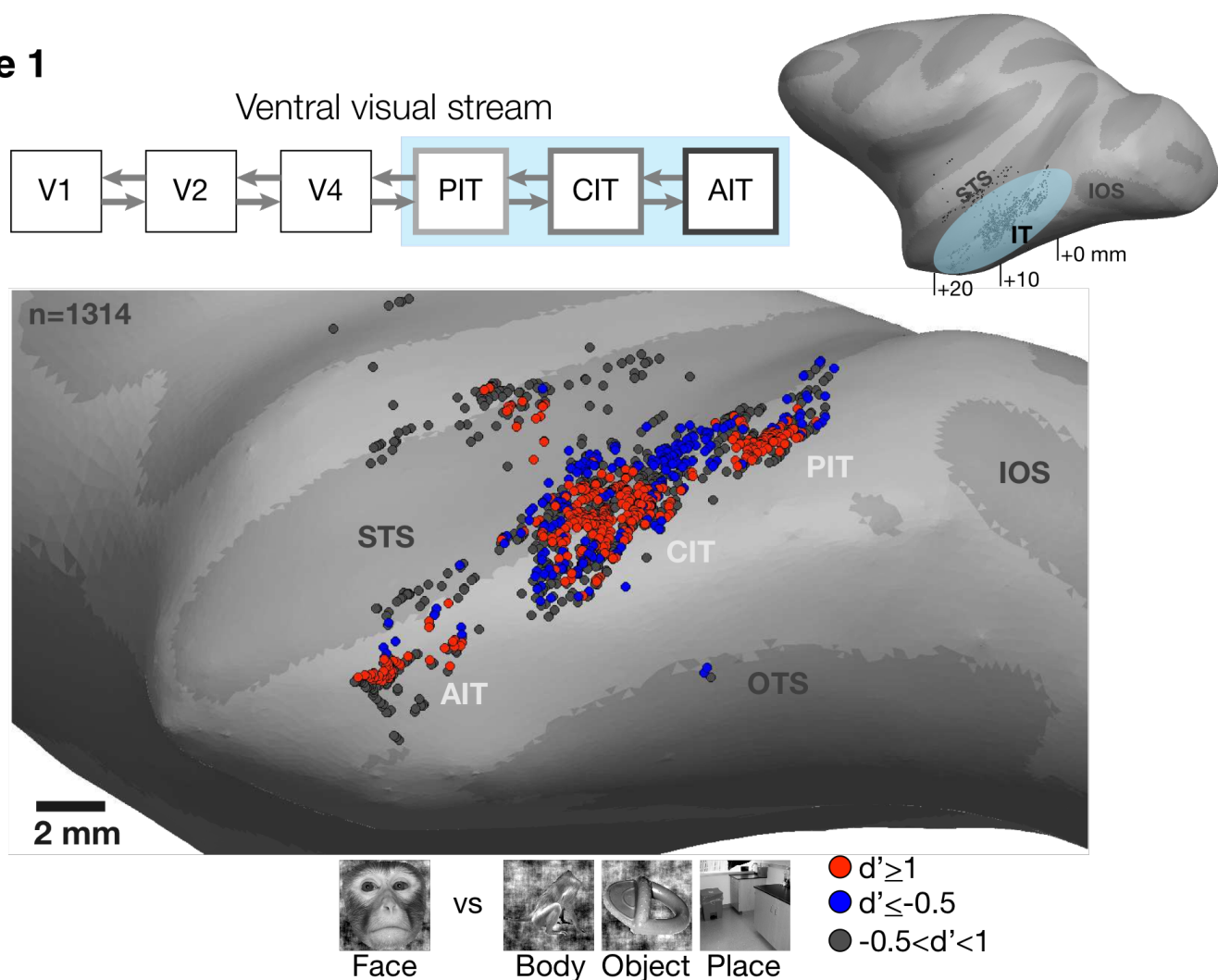


Figure 1 Neurophysiological recordings of face-selective subregions in the ventral visual stream. The ventral visual stream is a series of hierarchically connected areas (diagram in top row) and includes at least three IT processing stages (blue box). Neurons in these three stages were recorded along the lateral convexity of the inferior temporal lobe spanning the posterior to anterior extent of IT (+0 to +20 mm AP, Horsely-Clarke coordinates) in two monkeys (data from monkey 1 are shown). “Face neuron” sites (red) were operationally defined as those with a response preference for images of faces versus images of nonface objects (see **SI Methods**). While these were found throughout IT, they tended to be found in clusters that mapped to previously identified subdivisions of IT (posterior, central, and anterior IT) and also corresponded to face-selective areas identified under fMRI in the same subject (28)(48) (STS=superior temporal sulcus, IOS=inferior occipital sulcus, OTS=occipitotemporal sulcus).

Figure 2

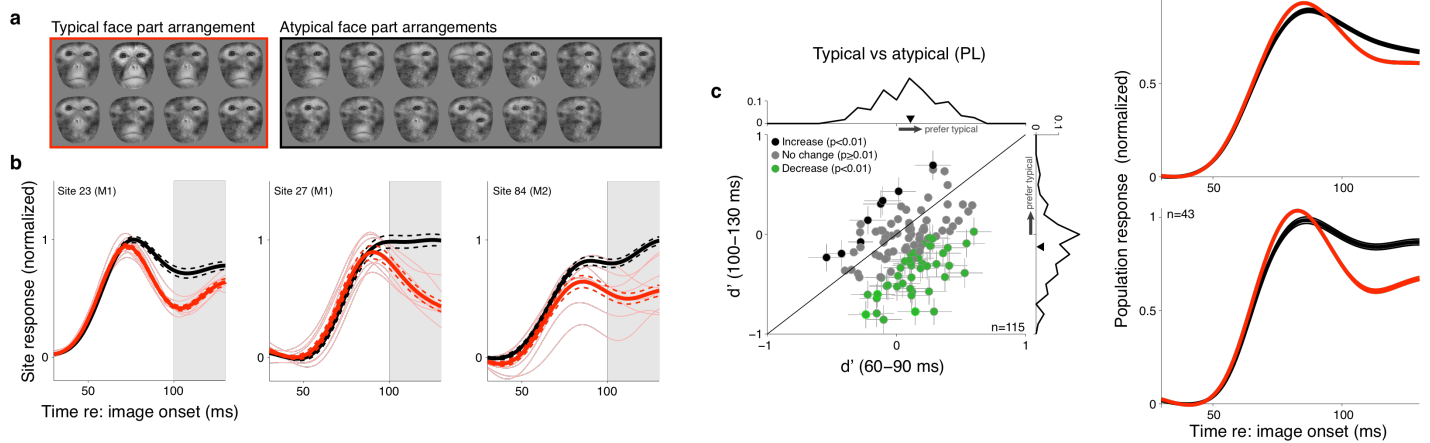


Figure 2 Responses in posterior IT to face-like images with typical and atypical face part arrangements. **(a)** Novel images were generated from an exemplar monkey face by positioning the face parts in different positions within the face outline. This procedure generated both typical (red) and atypical (black) arrangements of the face parts, and images that drove $\geq 90\%$ of the early (60-100 ms) response to the whole face at the population level are shown here (first image in red box is synthesized whole face; compare to the second image which is the original whole face). **(b)** Responses of three example sites. Preference for images with atypical arrangements (black line = mean responses of 13 images shown in **(a)**) over images with typical arrangements (red lines; 8 images) emerged late in the neural response (gray box). **(c)** Preference for typical versus atypical arrangements of individual sites in early (60-90 ms) and late (100-130 ms, same as gray box in **(b)**) time windows (sites showing significant changes in their preference ($p < 0.01$) are plotted with error bars and colored according to whether their preference (d') for typical vs. atypical face images increased (black), decreased (green), or did not change (gray); the three example sites from **(b)** all had decreasing preference and are highlighted by the green outlines; the marginal distributions are shown where inverted triangles are the median d' in early and late time windows). **(d)** Population average response in pIT across all images from **(a)** with typical or atypical arrangements for the whole recorded population (top) and for the sites showing significantly decreased preference for typical versus atypical arrangements (bottom) (same sites as green dots in **(c)**).

Figure 3

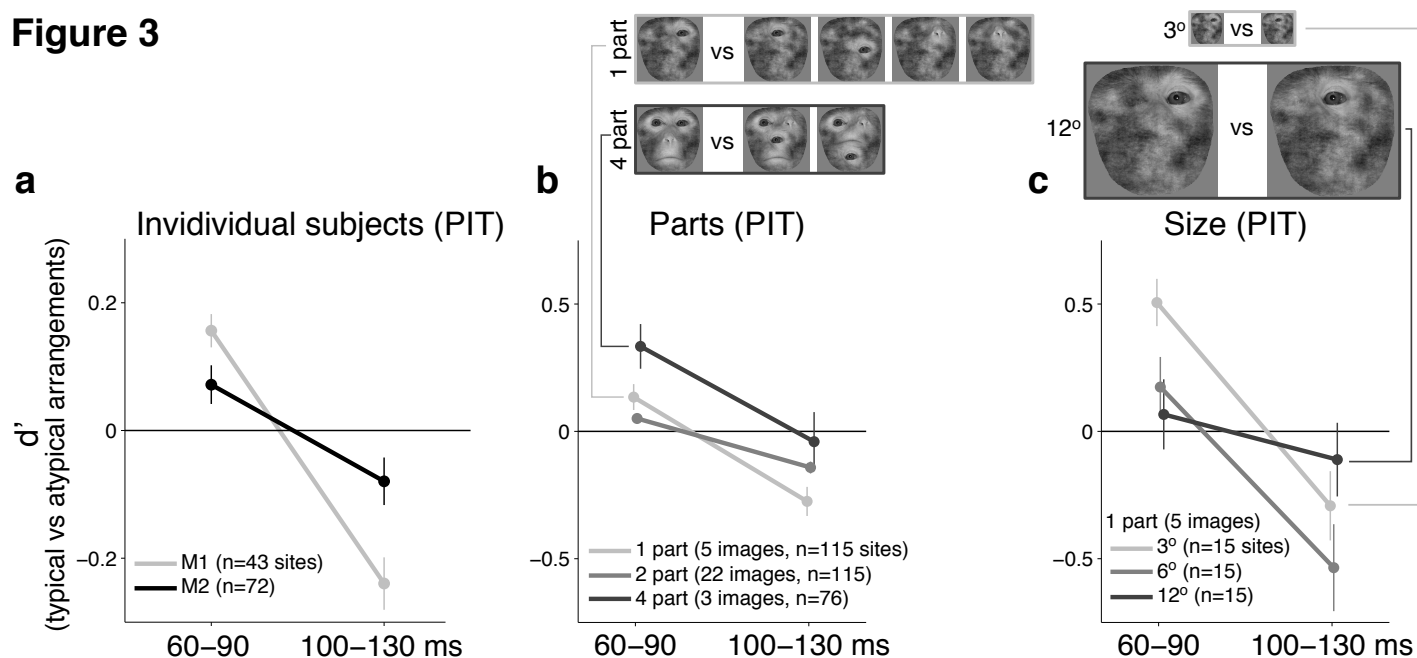


Figure 3 Individual monkey comparison and image controls for the reversal of rank-order selectivity in pIT. **(a)** Preference for images with typical versus atypical face part arrangements analyzed separately for each monkey. Median d' across sites in both early and late time windows is shown. **(b)** Preference for images with typical versus atypical face part arrangements was re-computed using image subsets containing the same number of parts in the outline (the five 1-part and the three 4-part image subsets shown at top; the larger 2-part subset contained 30 images total and is not shown). **(c)** The 1-part image subset was further tested at three different sizes (3° , 6° , and 12°). In all cases, pIT responses showed an initial preference for typically-arranged face parts, followed by a later preference for atypically arranged face parts.

Figure 4

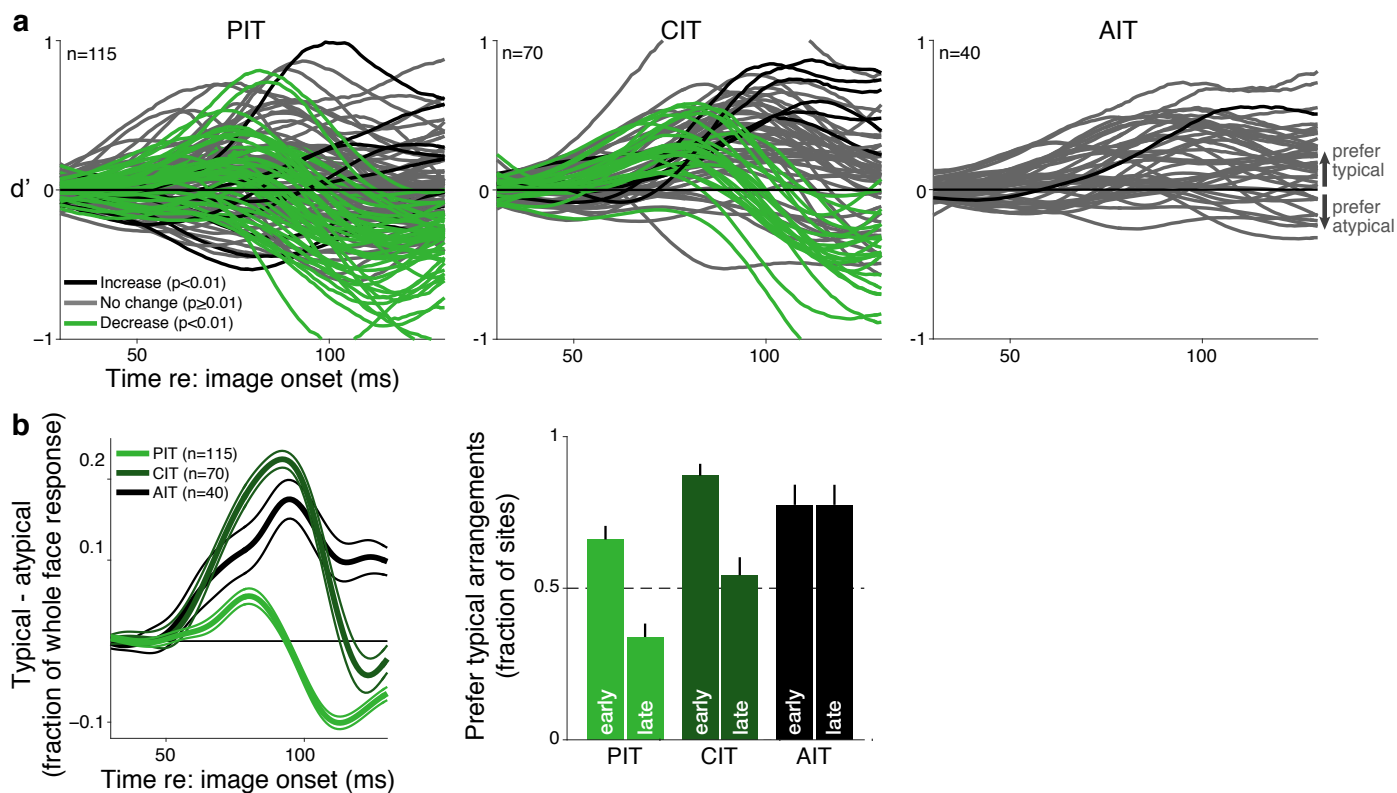


Figure 4 Time course of neural response preferences in posterior, central, and anterior IT for images with typical versus atypical face part arrangements. **(a)** The initial preference (d') in pIT for typical face part arrangements reversed in many sites following the first 30 ms of the response (left panel; sites in green showed significant decreases in selectivity ($p < 0.01$); same color scheme as **Fig. 2c**). In aIT, selectivity for face-like arrangements of parts generally increased over time (right panel) (sites plotted in order of absolute change in preference; sites with largest absolute change in preference are in foreground). **(b)** Left: Population average response to typical versus atypical arrangements (left, normalized by the mean response of the population to the whole face). Right: The fraction of sites whose responses showed a preference for images of typical, face-like arrangements of the face parts (right) in pIT (green), cIT (dark green) and aIT (black) in the early (60-90 ms) and late (100-130 ms) phase of the response. Note that, in the late phase of the response, most pIT neurons paradoxically show a preference for atypical arrangements of face parts.

Figure 5

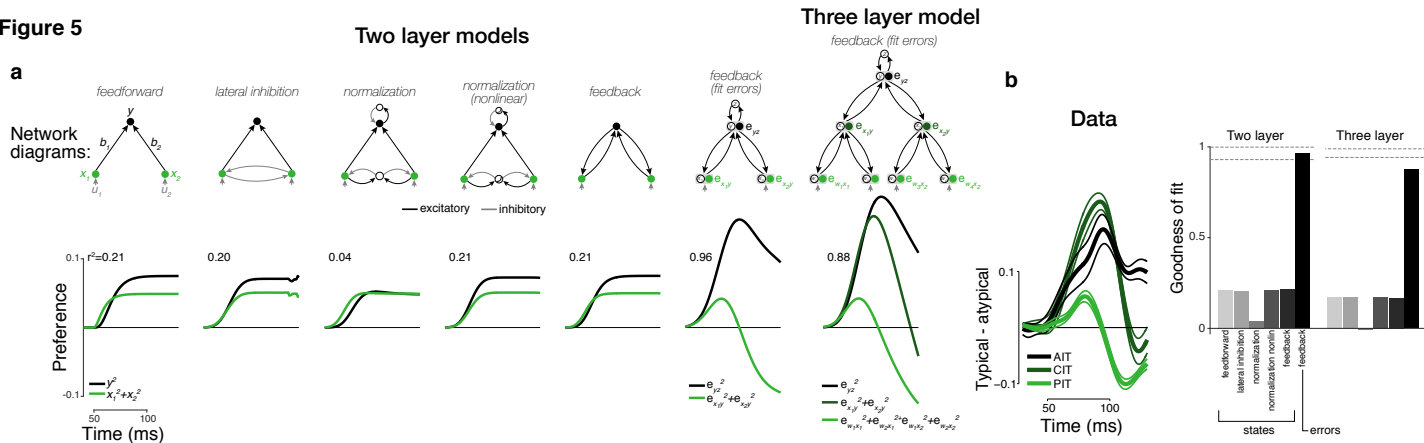


Figure 5 Computational modeling of neural dynamics in IT. **(a)** Simple two-layer neural networks (first six columns) with recurrent dynamics (network diagrams in top row; see **SI Methods** and **Supplementary Fig. 1 a,b**) were constructed to model neural signals measured in pIT and aIT corresponding to the first (green) and second (black) model processing stages. All models received two inputs (gray) into two hidden stage units (green) which sent feedforward projections that converged onto a single unit in the output stage (black). Besides this feedforward architecture, additional excitatory and inhibitory connections between units were used to implement recurrent dynamics (self connections reflecting leak currents are not shown here for clarity; see **Supplementary Fig. 1a** for detailed diagrams). In the five models on the left, the responses of the simulated neurons are assumed to code the current estimates of some set of features in the world (a.k.a states), as is standard in most such networks. The best fit to the population averaged neural data in **(b)** (same as left panel in **Fig. 4b**) of the states of each model class are shown (first five columns). These state coding models showed increasing selectivity over time from hidden to output layers and did not demonstrate the strong reversal of stimulus preference in their hidden processing stage (green lines) as observed in the pIT neural population. However, the neurons coding errors in a feedback-based hierarchical model did show a strong reversal of stimulus preference in the hidden processing stage (sixth column; reconstruction errors instead of the states were fit directly to the data). This model which codes the error signals (filled circles) also codes the states (open circles) (network diagram in sixth column of top row). A three-layer version of this model also produced reversal dynamics in the hidden layers (seventh column). **(b)** Population averaged neural selectivity profile shown (left, same as left panel in **Fig. 4b**). (right) Goodness of fit of all two-layer and three-layer models tested (dashed lines represent mean and standard error of reliability of neural data as estimated by bootstrap resampling).

Figure 6

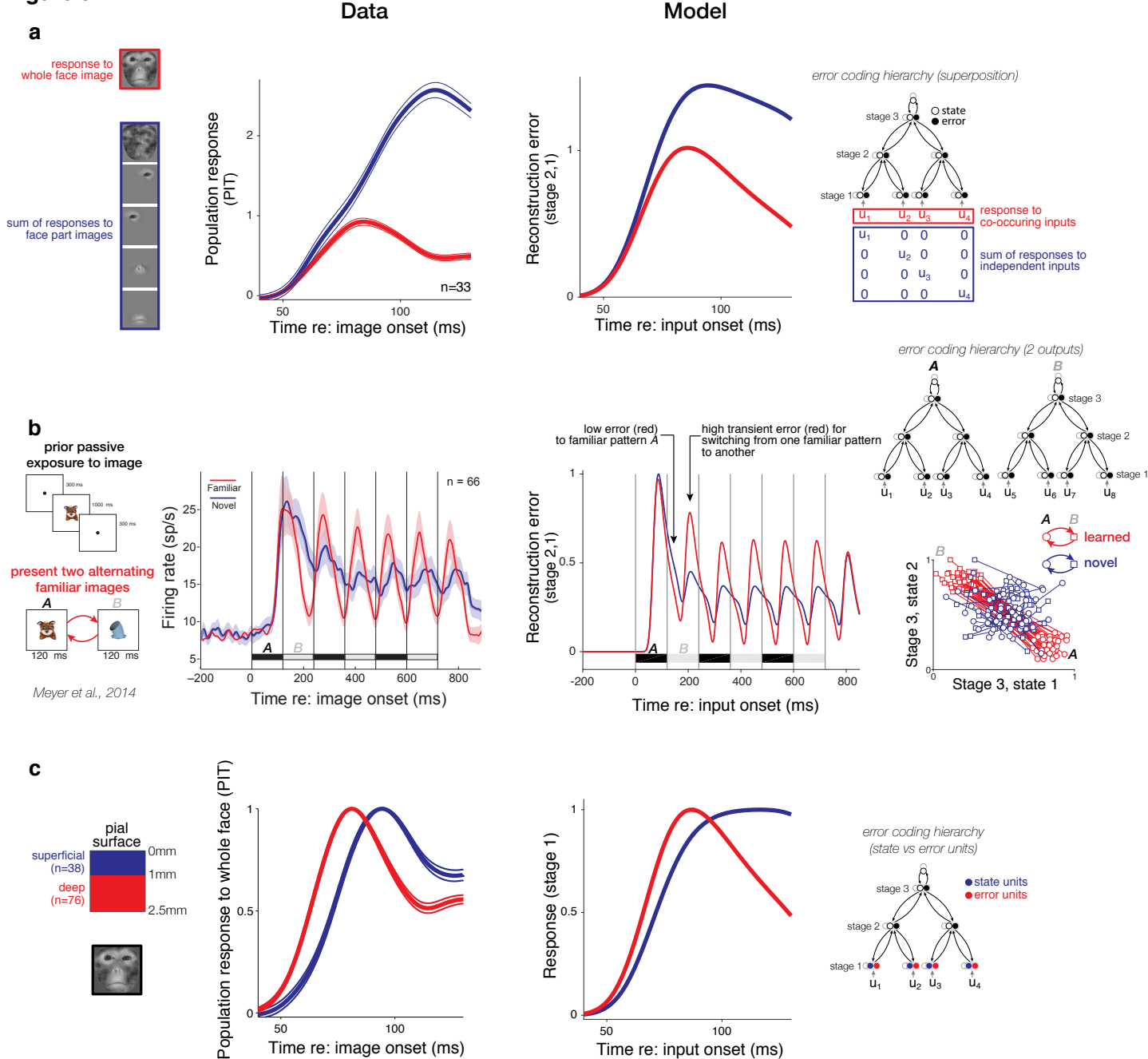


Figure 6 Comparison of neuronal findings and predictions of a feedback-based error coding model. To generate model predictions, the architectural parameters of the model (i.e. connection weights and time constants) were held fixed, and only the input patterns u were varied. **(a)** Neuronal (left panel): in pIT, we found that the sum of the neuronal response to the face parts presented individually (blue) exceeded the response to the same parts presented simultaneously (i.e. a whole face, red). Each line is the mean response of 33 pIT sites normalized by the peak response to the whole face. Model (right panel): The magnitude of errors between stage 1 and stage 2 of the model showed a similar degree of sublinear integration (responses are normalized by peak response to the simultaneous input condition). **(b)** We extended the model to include two units in the third, output stage that responded to two learned input patterns (see **SI Methods**), increasing separation of patterns A and B in this high-level feature space (red markers in far right panel; 50 draws were made from distributions for A and B and were compared to relatively arbitrary inputs, blue markers). When alternating the two learned or familiar patterns A and B, activations in the top layer of the networks experienced greater

changes than when two randomly selected patterns were alternated (compare distances traversed in high-level feature space by red lines versus blue lines, far right panel). Because of this large change in high-level activations, transient errors were generated back through the network for learned patterns. Strong oscillations could be observed in the error signals between stage 1 and stage 2 of the model for alternated familiar inputs *A* and *B* (120 ms period) (red curve, middle right panel). In contrast, alternating novel inputs with similar amplitude but with patterns not matching the learned weights led to small amplitude oscillations in upstream error signals (blue curve). Note that the average response strength of state signals to novel and familiar inputs was matched at all network levels by construction. The differing response dynamics of error signals under familiar versus novel patterns are qualitatively consistent with the IT findings for novel versus familiar images (left, reproduced with permission from *Figs. 1 & 2b* of Meyer et al., 2014). **(c)** The average response to the whole face for pIT sites recorded 0 to 1 mm below the pial surface (left panel; superficial recordings, blue curve) and for sites 1 to 2.5 mm beneath the pial surface (red curve). (right panel) Average response of state units (blue) and error units (red) in stage 1 of the model. Note the lagged response in state units which is similar to the lagged response of units in superficial recordings (red curve, left panel).

Figure 7

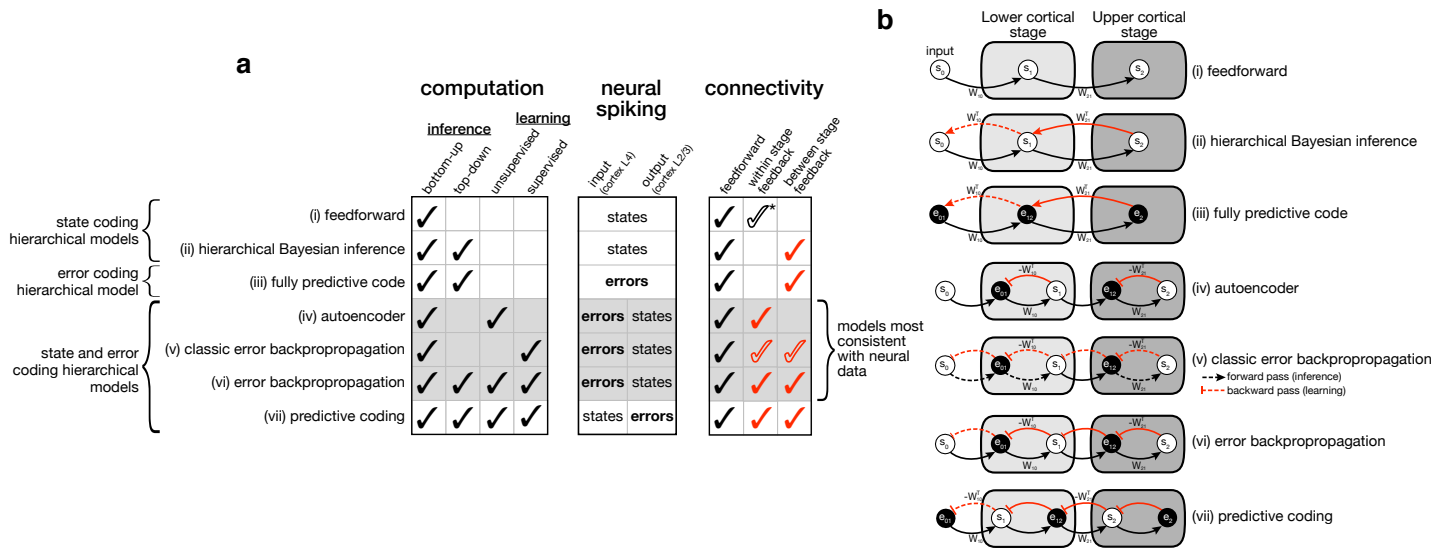
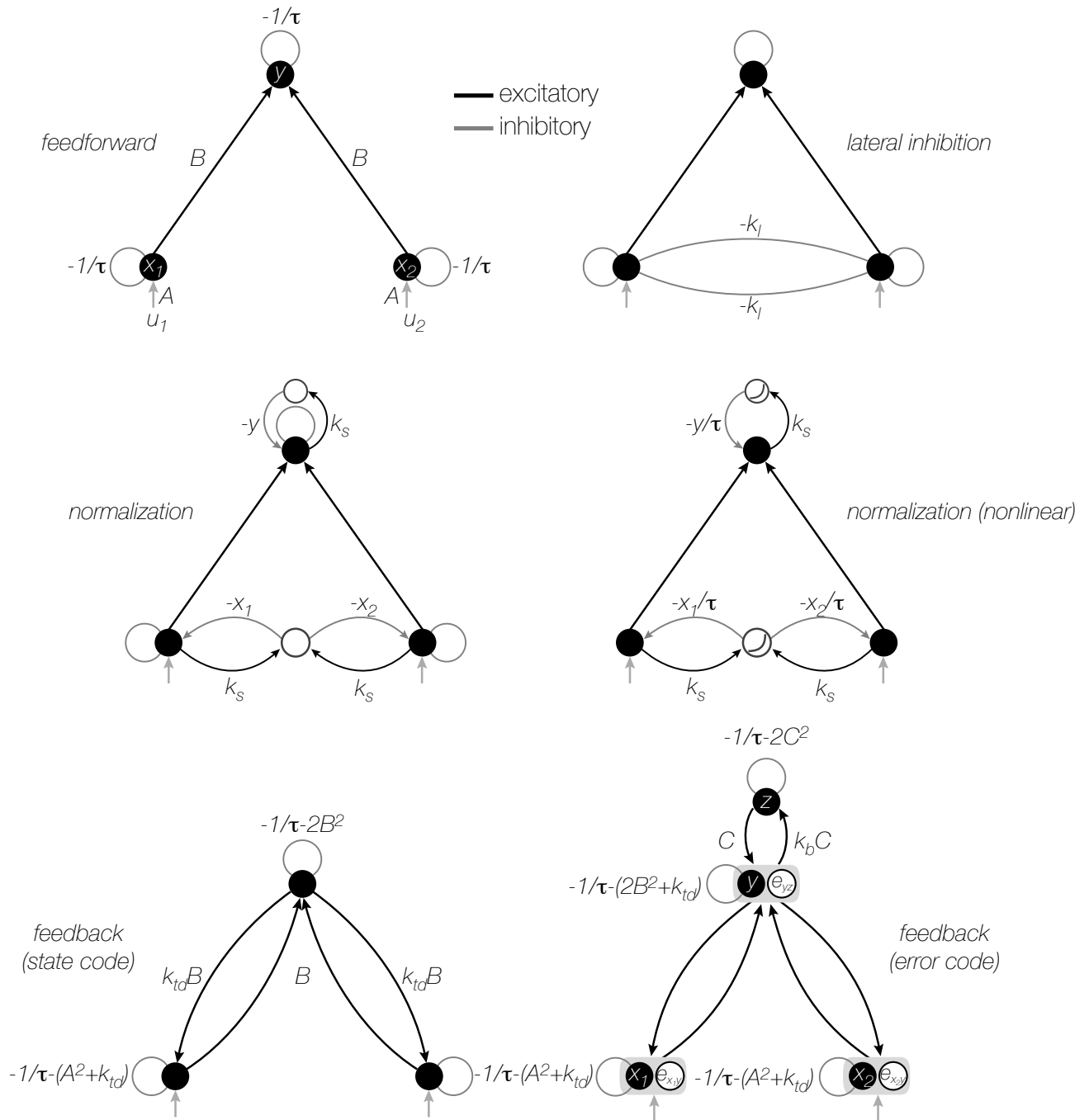


Figure 7 Summary of hierarchical model families. **(a)** Neural network models can be divided based on the computations that they perform (left table) and how they instantiate those computations through neurons and their connectivity (right two tables). (i): All model families shown build from the basic feedforward hierarchical model family to include either top-down inference, unsupervised learning, or supervised learning ((* indicates that feedforward models can in some cases include local, within-stage recurrent connections such as the normalization models tested in **Fig. 5**). (ii): In hierarchical Bayesian inference (14)(31), feature estimates (i.e. inferred states of variables in the world) are communicated between processing stages to combine bottom-up estimates and top-down predictions. (iii): In a fully predictive code, neural spiking encodes the residual errors instead of directly encoding the feature estimates. This coding strategy amounts to a simple change of variables (see **SI Methods**) and is energy efficient in that the transmitted error signals are smaller (minimized by virtue of minimizing reconstruction cost) than state signals. A fully predictive coding model is shown here only for completeness as one example of an error coding model that performs efficient inference but not efficient learning, but see (vii) for a variant of predictive coding that also performs efficient learning. (iv): A simple form of a stacked autoencoder (33)(42) uses reconstructive error signals to drive unsupervised learning within each processing stage. (v): In classic error backpropagation (5)(19), errors are passed backwards between network levels during supervised training offline. (vi) Contemporary backpropagation approaches (42)(43) utilize error signals during online inference (similar to hierarchical Bayesian inference) through between-stage connections as well as during offline learning, through the same between-stage connections. Placing error coding units in the feedforward path makes these operations possible through basic neural integration and Hebbian plasticity mechanisms. (vii): Predictive coding (21)(40) is computationally similar to (vi), but it differs at the implementation level by specifically positing that error information (rather than state information) is passed to the next higher cortical stage. **(b)** Between-stage and within-stage connectivity diagrams corresponding to the models in **(a)**. Between-stage errors (black circles), measuring reconstruction performance, are computed in a similar fashion across models and can drive efficient hierarchical learning when coupled with state signals (white circles) (bottom four networks). The state and error computing networks only differ in the details of how error signals and state signals interact during inference and learning. Our data provide evidence for this large family of error-computing networks and rule out pure state-estimating models and variants including normalization and lateral inhibition (**Fig. 5**). The present data do not distinguish between the autoencoder and error backpropagation classes when directly compared (**Supplementary Fig. 2**); however, the stronger presence of state-like signals in the superficial cortical layers (**Fig. 6c**) argues against the predictive coding models in (iii) and (vii).

Supplementary Figure 1

a Two-stage models (network diagrams)



b Two-stage models (connection matrices)

$$\dot{\mathbf{v}} = W\mathbf{v} + A\mathbf{u}, \quad \mathbf{v} = \begin{bmatrix} x_1 \\ x_2 \\ y \end{bmatrix}, \quad \mathbf{u} = \begin{bmatrix} u_1 \\ u_2 \\ 0 \end{bmatrix}$$

$$W_{\text{feedforward}} = \begin{pmatrix} -1/\tau & 0 & 0 \\ 0 & -1/\tau & 0 \\ B & B & -1/\tau \end{pmatrix}$$

$$W_{\text{lateral inhibition}} = \begin{pmatrix} -1/\tau & -k_l & 0 \\ -k_l & -1/\tau & 0 \\ B & B & -1/\tau \end{pmatrix}$$

$$W_{\text{normalization}} = \begin{pmatrix} -1/\tau - k_s \sum x & 0 & 0 \\ 0 & -1/\tau - k_s \sum x & 0 \\ B & B & -1/\tau - k_s y \end{pmatrix}$$

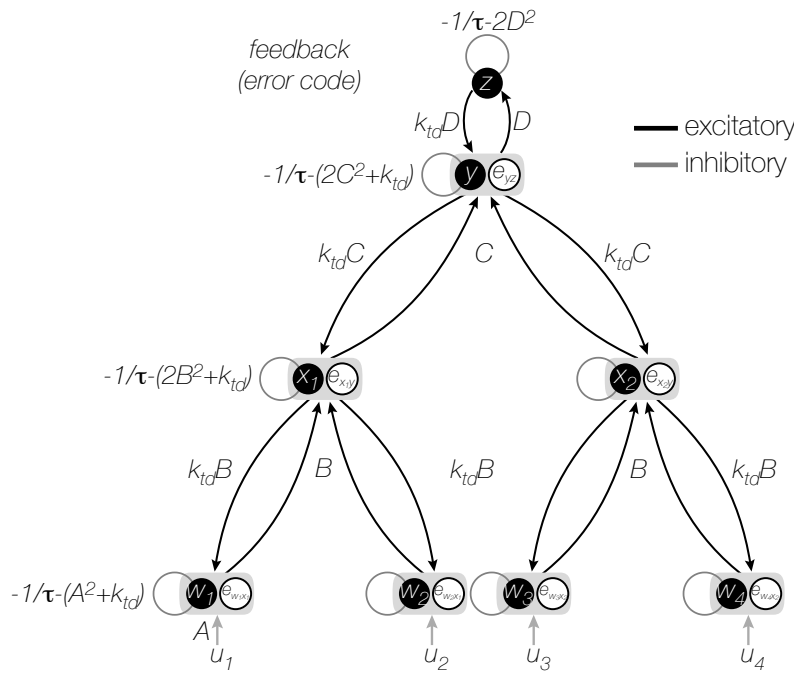
$$W_{\text{normalization (nonlinear)}} = \begin{pmatrix} -\frac{1}{\tau\sqrt{1 - k_s \sum x}} & 0 & 0 \\ 0 & -\frac{1}{\tau\sqrt{1 - k_s \sum x}} & 0 \\ B & B & -\frac{1}{\tau\sqrt{1 - k_s y}} \end{pmatrix}$$

$$W_{\text{hierarchical generative (state)}} = \begin{pmatrix} -1/\tau - (A^2 + k_b) & 0 & k_b B \\ 0 & -1/\tau - (A^2 + k_b) & k_b B \\ B & B & -1/\tau - 2B^2 \end{pmatrix}$$

$$W_{\text{hierarchical generative (error)}} = \begin{pmatrix} -1/\tau - (A^2 + k_b) & 0 & k_b B & 0 \\ 0 & -1/\tau - (A^2 + k_b) & k_b B & 0 \\ B & B & -1/\tau - (2B^2 + k_b) & k_b C \\ 0 & 0 & C & -1/\tau - 2C^2 \end{pmatrix}$$

C

Three-stage feedback model (network diagram)



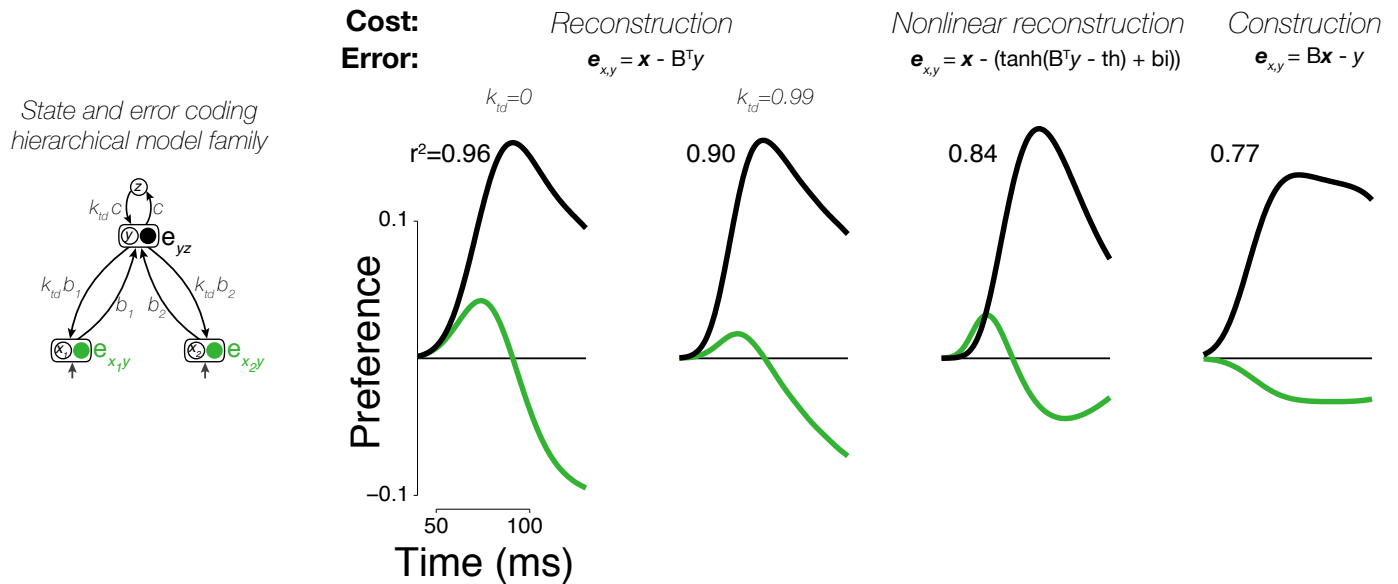
Three-stage feedback model (connection matrix)

$$W_{\text{hierarchical generative (error)}} = \begin{pmatrix} -1/\tau - (A^2 + k_b) & 0 & 0 & 0 & k_b B & 0 & 0 & 0 & 0 \\ 0 & -1/\tau - (A^2 + k_b) & 0 & 0 & k_b B & 0 & 0 & 0 & 0 \\ 0 & 0 & -1/\tau - (A^2 + k_b) & 0 & 0 & k_b B & 0 & 0 & 0 \\ 0 & 0 & 0 & -1/\tau - (A^2 + k_b) & 0 & k_b B & 0 & 0 & 0 \\ B & B & 0 & 0 & -1/\tau - (2B^2 + k_b) & 0 & k_b C & 0 & 0 \\ 0 & 0 & B & B & 0 & -1/\tau - (2B^2 + k_b) & k_b C & 0 & 0 \\ 0 & 0 & 0 & 0 & C & C & -1/\tau - (2C^2 + k_b) & k_b D & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & D & -1/\tau - 2D^2 \end{pmatrix}$$

Supplementary Figure 1 Model diagrams and equations. **(a)** Network diagrams of two-stage models. All models have two inputs (u_1, u_2), at least two hidden units (x_1, x_2) (a.k.a. layer 1 of the network), and at least one output unit (y). For simulations, the inputs u_1, u_2 are set independently to simulate each hidden node receiving different amounts of external drive, depending on the choice of the applied image relative to the unit's preferred image. The connection weights $B = [b_1, b_2]$ transforming the hidden stage activations to the output unit are modeled as the same ($b_1 = b_2$). All units have self-connections that determine the degree of leak current set by the time constant τ . In the normalization models, the leak term is additionally controlled (linearly or nonlinearly) by the total activity in each stage (second row). In the feedback-based model, the feedback connections are symmetric to the feedforward connections with weights $B^T = [b_1, b_2]^T$, a column vector (third row). The error coding feedback model (bottom, right) has an additional stage that contributes to computation of error in the second stage (see **Methods** for details). **(b)** Equivalent system of differential equations for the networks shown in **(a)**. The general linear matrix equation governing the dynamics of the models is shown in the top row followed by the weight matrices W for each model in subsequent rows. The weight matrices determine the connection architecture of the models. Each model class uses a similar number of parameters to specify a different arrangement of connections. Strictly speaking, the two normalization model classes cannot

be written as linear systems of differential equations since the coefficients on the diagonal depend on the the state variables themselves creating nonlinear (multiplicative) terms, and for all simulations, solutions were computed using a standard iterative differential equation solver. **(c)** Extension of the two-stage model architectures to three stages for the feedback model (compare to two-stage diagram in **(a)**). The three-stage model has an additional hidden processing stage compared to the two-stage model. An extra node is introduced at the top of the hierarchy to produce an error signal in the third stage. The connection matrix for this model is shown below its network diagram (compare to matrix in **(b)**).

Supplementary Figure 2



Supplementary Figure 2 Variants of error coding hierarchical models using different algorithms for online inference. Existing forms of error computing networks can be distinguished by the type of online inference algorithm that they use (**Figure 7**). In one case, inference does not utilize top-down information between stages (autoencoder, classic error backpropagation; model classes (iv) and (v) in **Figure 7**). On the other hand, between-stage feedback can be used such as in more general forms of error backpropagation and predictive coding (models (vi) and (vii) in **Figure 7**). We approximated these two extremes by including a parameter (k_{id} , see **Methods**) controlling the relative weighting of bottom-up (feedforward) and top-down (feedback) evidence during online inference (first and second panels). We found that top-down inference between stages was not necessary to produce the appropriate error signals, and k_{id} was equal to zero in our best fitting two-layer and three-layers models (first panel is same model as two-layer error coding model in main text **Figure 5**) although models with $k_{id} \sim 1$ also performed well (second panel). Models can also differ in their goal (cost function) which directly impacts the error signals required (top row). Under a nonlinear reconstruction goal (emulating the nonlinear nature of spiking output), the resulting error signals are still consistent with our data (third column). A simple sigmoidal nonlinearity, however, did lead to additional details present in our neural data such as a rapid return of stimulus preference to zero in the hidden layer. When we tested a discriminative, construction goal more consistent with a supervised learning setting (e.g. classic error backpropagation) where bottom-up responses simply have to match a downstream target signal in classification tasks, we found that the errors of construction did not match the data as well as reconstruction errors (compare fourth column to first three columns; note absence of a reversal of selectivity in the hidden processing stage in fourth column) although both types of error outperformed all state-based models and were overall very similar (compare to **Fig. 5**).