

1 **Neural dynamics at successive stages of the ventral visual stream are** 2 **consistent with hierarchical error signals**

3 Running title: Error signaling in the ventral stream

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14 **AUTHOR CONTRIBUTIONS**

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19

20 **ABSTRACT**

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22
23 Ventral visual stream neural responses are dynamic, even for static image presentations. However,
24 dynamical neural models of visual cortex are lacking as most progress has been made modeling static,
25 time-averaged responses. Here, we studied population neural dynamics during face detection across
26 three cortical processing stages. Remarkably, ~30 milliseconds after the initially evoked response, we
27 found that neurons in intermediate level areas decreased their preference for faces, becoming anti-face
28 preferring on average even while neurons in higher level areas achieved and maintained a face
29 preference. This pattern of hierarchical neural dynamics was inconsistent with extensions of standard
30 feedforward circuits that implemented recurrence within a cortical stage. Rather, recurrent models
31 computing errors between stages captured the observed temporal signatures. Without additional
32 parameter fitting, this model of neural dynamics, which simply augments the standard feedforward
33 model of online vision to encode errors, also explained seemingly disparate dynamical phenomena in
34 the ventral stream.
35

36 INTRODUCTION

37 The primate ventral visual stream is a hierarchically organized set of cortical areas beginning with the
38 primary visual cortex (V1) and culminating with distributed patterns of neural firing across the inferior
39 temporal cortex (IT) that explicitly encode objects (i.e. linearly decodable object identity) (Hung,
40 Kreiman, Poggio, & DiCarlo, 2005) and quantitatively account for core invariant object discrimination
41 behavior in primates (Majaj, Hong, Solomon, & DiCarlo, 2015). Formalizing object recognition as the
42 result of a series of feedforward computations yields models that achieve impressive performance in
43 object categorization (Krizhevsky, Sutskever, & Hinton, 2012)(Zeiler & Fergus, 2013) similar to the
44 absolute level of performance achieved by IT neural populations, and these models are the current best
45 predictors of neural responses in IT cortex and its primary input layer, V4 (Cadieu et al., 2014)(Yamins
46 et al., 2014). Thus, the feedforward inference perspective provides a simple but powerful, first-order
47 framework for the ventral stream and core invariant object recognition.

48

49 However, visual object recognition behavior may not be executed via a single feedforward neural
50 processing pass (a.k.a. feedforward inference) because IT neural responses are well-known to be
51 dynamic even in response to images without dynamic content (Brincat & Connor, 2006)(Sugase,
52 Yamane, Ueno, & Kawano, 1999)(Chen et al., 2014)(Meyer, Walker, Cho, & Olson, 2014), raising the
53 question of what computations those neural activity dynamics might reflect. Prior work has proposed
54 that such neuronal response dynamics could be the result of different types of circuits executing
55 different types of computation such as: 1) recurrent circuits within each ventral stream processing stage
56 implementing local normalization of the feedforward information as it passes through the stage
57 (Carandini, Heeger, & Movshon, 1997)(Schwartz & Simoncelli, 2001)(Carandini & Heeger, 2012), 2)
58 feedback circuits between each pair of ventral stream stages implementing the integration of top-down
59 with bottom-up information to improve the current (online) inference (Seung, 1997)(Lee, Yang, Romero,
60 & Mumford, 2002)(Zhang & Heydt, 2010)(Epshtein, Lifshitz, & Ullman, 2008), or 3) feedback circuits

61 between each pair of stages comparing top-down and bottom-up information to compute prediction
62 errors that guide changes in synaptic weights so that neurons are better tuned to features useful for
63 future feedforward behavior (learning) (Rao & Ballard, 1999). Thus, neural dynamics may reflect the
64 various adaptive computations (within-stage normalization, top-down Bayesian inference) or reflect the
65 underlying error intermediates that could be generated during those processes (e.g. predictive coding).
66

67 These computationally motivated ideas can each be implemented as neural circuits to ask which
68 idea best predicts response dynamics across the visual hierarchy. Here, our main goal was to look
69 beyond the initial, feedforward response edge to see if we could disambiguate among dynamics that
70 might result from stacked feedforward, lateral, and feedback operations. Rather than record from a
71 single processing level, we measured the dynamics of neural signals across three hierarchical levels
72 (pIT, cIT, aIT) within macaque IT. We focused on face processing subregions within each of these
73 levels for three reasons. First, prior evidence argues that these three subregions are tightly
74 anatomically and functionally connected and that the subregion in pIT is the dominant input to the
75 higher subregions (Grimaldi, Saleem, & Tsao, 2016)(Moeller, Freiwald, & Tsao, 2008). Second,
76 because prior work argues that a key behavioral function of these three subregions is to distinguish
77 faces from non-faces, this allowed us to focus our testing on a relatively small number of images
78 targeted to engage that processing function. Third, prior knowledge of pIT neural tuning properties (Issa
79 & DiCarlo, 2012) allowed us to design images that were quantitatively matched in their ability to drive
80 neurons in the pIT input subregion but that should ultimately be processed into two separate groups
81 (face vs. non-face). We reasoned that these images would force important computations for
82 disambiguation to occur somewhere between the pIT subregion and the higher level (cIT, aIT)
83 subregions. With this setup, our aim was to observe the dynamics at all three levels of the hierarchy in
84 response to that image processing challenge so that we might discover – or at least constrain -- which
85 type of computation is at work.

86

87 Consistent with the idea that the overall system performs face vs. non-face discrimination (i.e.
88 face detection), we found that in the highest face processing stage (aIT), neurons rapidly developed
89 and maintained a response preference for faces over non-faces even though our images were
90 designed to be challenging for frontal face detection. However, we found that many neurons in the early
91 (pIT) and intermediate (cIT) processing levels of IT had face selectivity that rapidly but paradoxically
92 *decreased*. That is, the responses of these neurons evolved to prefer images of non-faces over images
93 of faces within 30 milliseconds of their feedforward response. We found that standard feedforward
94 models that employ local recurrences such as adaptation, lateral inhibition, and normalization could not
95 capture this stage-wise pattern of image selectivity despite our best attempts. However, we found that
96 *decreasing* -- rather than increasing -- face preference in early and intermediate processing stages is a
97 natural dynamical signature of previously suggested “error coding” models (Rao & Ballard, 1999) in
98 which the neural spiking activity at each processing stage carries both an explicit representation of the
99 variables of interest (e.g. is a face present?) and an explicit encoding of errors computed between each
100 pair of stages in the hierarchy (e.g. a face was predicted, but a non-face was present leading to an
101 error).

102

103 **RESULTS**

104 We leveraged the hierarchically arranged face processing system in macaque ventral visual cortex to
105 study the dynamics of neural processing across a hierarchy (Tsao, Freiwald, Tootell, & Livingstone,
106 2006)(Tsao, Moeller, & Freiwald, 2008) (**Figure 1A**). The serially arranged posterior, central, and
107 anterior face-selective subregions of IT (pIT, cIT, and aIT) can be conceptualized as building increasing
108 selectivity for faces culminating in aIT representations (Freiwald & Tsao, 2010)(Chang & Tsao, 2017).
109 Using serial, single electrode recording, we sampled neural sites across the posterior to anterior extent
110 of the IT hierarchy in the left hemispheres of two monkeys to generate neurophysiological maps

111 **(Figure 1A**; example neurophysiological map in one monkey using a faces versus non-face objects
112 screen set) (Issa, Papanastassiou, & DiCarlo, 2013). We localized the recording locations *in vivo* and
113 co-registered across all penetrations using a stereo microfocal x-ray system (~400 micron *in vivo*
114 resolution) (Cox, Papanastassiou, Oreper, Andken, & DiCarlo, 2008)(Issa, Papanastassiou, Andken, &
115 DiCarlo, 2010) allowing accurate assignment of sites to different face processing stages (n = 633 out of
116 1891 total sites recorded were assigned as belonging to a face-selective subregion based on their
117 spatial location; see **Methods**). Results are reported here for sites that were spatially located in a face-
118 selective subregion, that showed visual drive to any category in the screen set (see **Methods**), and that
119 were subsequently tested with our face versus non-face challenge set (**Figure 1B**, left panel) (n = 115
120 pIT, 70 cIT, and 40 aIT sites).

121

122 Our experimental design was intended to test previously proposed computational hypotheses of
123 hierarchical neural dynamics during visual face processing (**Figure 1B**). Briefly, these hypotheses
124 predict how stimulus preference (in this instance, for faces versus non-faces) might change over time in
125 a neural population (**Figure 1B**, middle panel): (1) simple spike-rate adaptation predicts that initial rank-
126 order selectivity (i.e. relative stimulus preference) will be largely preserved (**Figure 1B**, dashed line)
127 while neurons adapt their absolute response strength over time, (2) local normalization predicts that
128 stronger responses are in some cases normalized to match weaker responses based on population
129 activity to specific dimensions (Carandini et al., 1997); importantly, normalization is strongest for
130 nuisance (non-coding) dimensions (e.g. low versus high stimulus contrast) and in its idealized form
131 would not alter selectivity along coding dimensions (e.g. face versus non-face) (**Figure 1B**, dashed
132 line), (3) evidence accumulation through temporal integration, winner-take-all through recurrent
133 inhibition, or Bayesian inference through top-down feedback mechanisms all predict increasing
134 selectivity for faces over time (Lee & Mumford, 2003) (**Figure 1B**, light gray line), and (4) predictive
135 coding posits that, for neurons that are coding error, their responses would show increasing activity for

136 non-faces (images whose properties are inconsistent with predictions of a face) and thus decreasing
137 face selectivity over time (Rao & Ballard, 1999) (**Figure 1B**, black line). Note, that error signaling is a
138 qualitatively different computation than normalization, as error coding predicts a decreased response
139 along the coding dimension (face versus non-face) whereas normalization would ideally not affect
140 selectivity for faces versus non-faces and only affect variation along orthogonal, nuisance dimensions.
141 Properly testing these predictions (no change in face selectivity, increased face selectivity, decreased
142 selectivity to be anti-face preferring) requires measurements from the intermediate stages of the
143 hierarchy as all of these models operate under the premise that the system builds and maintains a
144 preference for faces at the top of the hierarchy (**Figure 1B**, right, and see Introduction). Thus, the
145 intermediate stages (here pIT, see **Figure 1B**) are most likely to be susceptible to face/non-face
146 confusion and thus be influenced by, for example, the top-down mechanisms posited in Bayesian
147 inference and predictive coding where higher areas encode the face predictions that directly influence
148 the responses of lower areas (Lee & Mumford, 2003)(Rao & Ballard, 1999).

149

150 **Face and non-face images driving similar initial responses in pIT**

151 Here, we chose to focus our key, controlled tests on pIT – an intermediate stage in the ventral stream
152 hierarchy, but the first stage within IT where neural specialization for face detection (i.e. face vs. non-
153 face) has been reported (Grimaldi et al., 2016). Consistent with its intermediate position in the ventral
154 visual system, we had previously found that pIT face-selective neurons are not truly selective for whole
155 faces but respond to local face features, specifically those in the eye region (Issa & DiCarlo, 2012).
156 Taking advantage of this prior result, we created both face and non-face stimuli that challenged the
157 face processing system by strongly driving pIT responses, thus forcing the higher IT stages to complete
158 the discrimination between face and challenging non-face images. To generate ambiguous face-like
159 images, we systematically varied the positions of parts, in particular the eye, within the face (Issa &
160 DiCarlo, 2012) (see **Methods**). This set included images that contained face parts in positions

161 consistent with a frontal view of a face or images that only differed in the relative spatial configuration of
162 the face parts within the face outline (**Figure 1B**, left). Of the 82 images screened, we identified 21 part
163 configurations that each drove the pIT population response to $\geq 90\%$ of its response to a correctly
164 configured whole face. Of those 21 images, 13 images were inconsistent with the part configuration of a
165 frontal face (**Figure 1B**, black box). For the majority of the results that follow, we focus on comparing
166 the neural responses to these 13 pIT-matched images that could *not* have arisen from frontal faces
167 (referred to hereafter as “non-face images”) with the 8 images that could have arisen from frontal faces
168 (referred to hereafter as “face images”). Again, we stress that these two groups of images were
169 selected to evoke closely matched initial pIT population activity.

170

171 Importantly, the pIT-matched images used here presented a more stringent test of face vs. non-
172 face discrimination than prior work. Specifically, most prior work used images of faces and non-face
173 objects (“classic images”) that contain differences across multiple dimensions including local contrast,
174 spatial frequency, and types of features (Tsao et al., 2006)(Afraz, Kiani, & Esteky, 2006)(Moeller,
175 Crapse, Chang, & Tsao, 2017)(Sadagopan, Zarco, & Freiwald, 2017). Consistent with this, we found
176 that the population decoding classification accuracy of our recorded neural populations using these
177 classic images (faces versus non-face objects) is near perfect ($>99\%$ in pIT, cIT, and aIT, $n=30$ sites
178 per region). However, we found that population decoding classification accuracy for the pIT-matched
179 face vs. non-face images we used here was near chance level (50%) in pIT (**Figure 1C**, blue bar; by
180 comparison, classification accuracy for face versus non-face objects classification was 99.6% using the
181 same pIT sites). Further downstream in regions cIT and aIT, we found that the linear population
182 decoding classification of these pIT-matched face vs. non-face images was well above chance,
183 suggesting that our pIT-matched face detection challenge is largely solved somewhere between pIT
184 and aIT (**Figure 1C**).

185

186 **Time course of responses in pIT for images with face versus non-face arrangements of parts**

187 We next closely examined the pIT neural response dynamics. To do this, we defined a face preference
188 value (d' ; see Methods) that measured each site's average selectivity for the face images relative to the
189 non-face images, and we asked how a given site's preference evolved over time (see alternative
190 hypotheses in **Figure 1B**). First, we present three example sites which were chosen based on having
191 the largest selectivity (absolute d') in the late phase (100-130 ms post image onset). In particular, most
192 standard interpretations of face processing would predict a late phase preference for faces ($d' > 0$).
193 However, all three sites with the largest absolute d' had evolved a strong late phase preference for the
194 non-face images ($d' < 0$) despite having had very similar rising edge responses to the face and non-
195 face stimulus classes (response in early phase from 60-90 ms) (**Figure 2**, left column). A late, non-face
196 preference was not restricted to the example sites as a majority of pIT sites (66%) preferred non-faces
197 over faces in the late response phase (prefer frontal face arrangement: 60-90 ms = 66% vs. 100-130
198 ms = 34%; $p = 0.000$, $n = 115$) (**Figure 3B**, blue bars).

199 Next, we examined the dynamics of face selectivity across the pIT population as this is key to
200 disambiguating among the competing models outlined earlier (**Figure 1B**). In the adaptation and
201 normalization models, we would expect no change in the average population face selectivity, and the
202 evidence accumulation, winner-take-all, or Bayesian inference models predict an increase in face
203 selectivity over the population over time. Instead, we found that many sites significantly decreased their
204 face preference over time similar to the three example sites. Of the 51 sites in our pIT sample that
205 showed a significantly changing preference over time ($p < 0.01$ criterion for significant change in d'),
206 84% of these sites showed a decreasing preference ($n = 43$ of 51 sites, $p < 10^{-6}$, binomial test, $n =$
207 115 total sites) (**Figure 3A**, left column, light gray vs. black lines). This surprising trend -- decreasing
208 face preference -- was strong enough that it erased any small, initial preference for images of frontally
209 arranged face parts over the population (median d' : 60-90 ms = 0.11 ± 0.02 vs. 100-130 ms = $-0.12 \pm$
210 0.03, $p = 0.000$, $n=115$ sites), and this trend was observed in both monkeys when analyzed separately

211 ($p_{M1} = 0.000$, $p_{M2} = 0.002$, $n_{M1} = 43$, $n_{M2} = 72$ sites; **Figure 4A**). This decreasing face selectivity over
212 time was driven by decreasing firing rates to the face images containing normally arranged face parts.
213 Responses to these images were weaker by 18% on average in the late phase of the response
214 compared to the early phase (Δrate (60-90 vs 100-130 ms) = $-18\% \pm 4\%$, $p = 0.000$; $n = 7$ images)
215 while responses to the non-face images with atypical spatial arrangements of face parts -- also capable
216 of driving high early phase responses -- did not experience any firing rate reduction in the late phase of
217 the response (Δrate (60-90 vs 100-130 ms) = $2 \pm 1\%$, $p = 0.467$; $n = 13$ images).

218

219 The above observation of decreasing face preference over the pIT population seemed most
220 consistent with the prediction of error coding models, but one potential confound was that initial
221 responses to faces and challenging non-faces were not perfectly matched across the population (recall
222 that we only required face and non-face images to drive a response $\geq 90\%$ of the whole face response).
223 As a result, initial selectivity was non-zero ($d' = 0.11$, $n=115$ sites). This residual face preference may
224 be small, but if this residual face selectivity is driven by nuisance dimensions, for example excess
225 stimulus contrast in the face class relative to the non-face class, then the face class may have
226 experienced stronger activity dependent adaptation or normalization resulting in a decreasing face
227 preference over time. To more adequately limit general activity dependent mechanisms leading to
228 decreasing face responses, we performed control analyses where initial activity was tightly matched per
229 site or where the number of parts were matched across images.

230

231 **Controls in pIT for firing rate and low-level image variation**

232 To strictly control for the possibility that simple initial firing rate differences could predict the observed
233 phenomenon, we re-computed selectivity after first matching initial responses site-by-site. For this
234 analysis, images were selected on a per site basis to evoke similar initial firing rates (images driving

235 initial response within 25% of synthetic whole face response for that site, at least 5 images required per
236 class). This image selection procedure virtually eliminated any differences in initial responses between
237 the face and non-face image classes and hence any firing rate difference driven by differences in
238 nuisance parameters between faces and challenge non-faces (**Figure 3C**, 60-90 ms), yet we still
239 observed a significant drop in preference for images with typical frontal face part arrangements versus
240 atypical face part arrangements in pIT ($\Delta d' = -0.10 \pm 0.03$, $p = 0.001$, $n = 77$) (**Figure 3C**, blue line).
241 Thus, the remaining dependence of firing rate dynamics on the image class and not on initial response
242 strength argued against an exclusively activity based explanation to account for decreasing neural
243 responses to faces over time. Further arguing against this hypothesis, we found that the pattern of late
244 phase population firing rates in pIT across images could not be significantly predicted from early phase
245 pIT firing rates for each image ($\rho_{\text{pIT early, pIT late}} = 0.07 \pm 0.17$, $p = 0.347$; $n = 20$ images).

246 Thus far, we have performed analyses where images from the face and non-face class were
247 similar in their initially evoked response which equated images at the level of neural activity but
248 produced images varying in the number of parts. An alternative is to match the number of face parts
249 between the face and non-face classes as another means of limiting the differences in nuisance
250 dimensions such as the contrast, spatial frequency and retinal position of energy across images (see
251 examples in **Figure 4B**). When we recomputed selectivity across subsets of images containing a
252 matched number of one, two, or four parts ($n=5$, 30, and 3 images, respectively), we still observed that
253 pIT face selectivity decreased. For all three image subsets controlling the number of face parts, d'
254 began positive on average in the sampled pIT population (i.e. preferred frontal face part arrangements
255 in 60-90 ms post-image onset) (median d' for 60-90 ms = 0.13 ± 0.05 , 0.05 ± 0.02 , 0.33 ± 0.09 for one,
256 two, and four parts) and significantly decreased in the next phase of the response (100-130 ms post-
257 image onset) becoming negative on average (median d' for 100-130 ms: -0.27 ± 0.06 , -0.14 ± 0.02 , -
258 0.04 ± 0.12 ; one, two, four parts: $p = 0.000$, 0.000 , 0.004 , for d' comparisons between 60-90 ms and

259 100-130 ms, $n = 115$, 115, 76 sites) (**Figure 4B**). A similar decreasing face selectivity profile was
260 observed when we re-tested single part images at smaller (3°) and larger (12°) image sizes suggesting
261 a dependence on the relative configuration of the parts and not on their absolute retinal location or
262 absolute retinal size (median d' for 60-90 ms vs. 100-130 ms: three degrees = 0.51 ± 0.09 vs. $-0.29 \pm$
263 0.14 , twelve degrees = 0.07 ± 0.14 vs. -0.11 ± 0.14 ; $n = 15$; $p = 0.000, 0.025, 0.07$) (**Figure 4C**). Thus,
264 we suggest that the decreasing population face selectivity dynamic in pIT is a fairly robust phenomenon
265 specific to the face versus non-face dimension as this dynamic persists even when limiting potential
266 variation across nuisance dimensions. This phenomenon can be distinguished from normalization
267 mechanisms that would operate most strongly to reduce selectivity along nuisance dimensions rather
268 than along dimensions that directly solve the face versus non-face discrimination challenge.

269

270 **Time course of responses in aIT and cIT for images with face versus non-face arrangements of** 271 **parts**

272 Under the possibility that decreasing face selectivity in intermediate stage pIT is a signature of error
273 signaling, we next asked whether the source of the prediction signal could be observed in higher
274 cortical areas. In the anterior face-selective regions of IT which are furthest downstream of pIT and
275 reflect additional stages of feedforward processing (see block diagram in **Figure 1B**), we did not
276 observe the strong decreases in the selectivity profile seen in pIT. Indeed, the three sites with the
277 greatest selectivity (absolute d') in the late response phase (100-130 ms) in our aIT sample all
278 displayed a preference for frontal face part arrangements ($d' > 0$) (**Figure 2**, right column). Also, in
279 contrast to the dynamic selectivity profiles observed in many pIT sites, 98% of aIT sites (39 of 40) did
280 not significantly change their relative preference for face vs. non-face arrangements of the parts ($p <$
281 0.01 criterion for significant change at the site level) (**Figure 3A**, right column, bottom row, dark gray
282 sites). Rather, we observed a stable selectivity profile over time in aIT (median d' : 60-90 ms = $0.13 \pm$
283 0.03 vs. 100-130 ms = 0.17 ± 0.03 , $p = 0.34$, $n=40$ sites). As a result, the majority of anterior sites

284 preferred images with typical frontal arrangements of the face parts in the late phase of the response
285 (prefer face: 60-90 ms = 78% of sites vs. 100-130 ms = 78% of sites; $p = 0.451$, $n = 40$ sites; **Figure**
286 **3B**, red bars) despite only a minority (34%) of upstream sites in pIT preferring these images in their late
287 response. Thus, spiking responses of individual aIT sites were as expected from a computational
288 system whose purpose is to detect faces, as previously suggested (Freiwald & Tsao, 2010), and aIT
289 serves as one candidate for the putative prediction signal underlying face prediction errors.

290

291 In cIT whose anatomical location is intermediate to pIT and aIT, we observed many sites with
292 decreasing selectivity (**Figure 2C & 3A**, middle columns), a dynamic that persisted even when we
293 tightly matched initial responses on a site by site basis similar to pIT (**Figure 2C**, green line). The
294 overall stimulus preference in cIT was intermediate to that of pIT and aIT (**Figure 3B**) consistent with
295 the intermediate position of cIT in the IT hierarchy. Interestingly, we found that the patterns of
296 responses across images in the early phases of cIT and aIT activity were significant predictors of late
297 phase activity in pIT ($\rho_{\text{cIT early, pIT late}} = -0.52 \pm 0.11$, $p = 0.000$; $\rho_{\text{aIT early, pIT late}} = -0.36 \pm 0.14$, $p = 0.012$;
298 $n_{\text{pIT}}=115$, $n_{\text{cIT}}=70$, $n_{\text{aIT}}=40$ sites; $n = 20$ images), even better predictors than early phase activity in pIT
299 itself ($\rho_{\text{pIT early, pIT late}} = 0.07 \pm 0.17$, $p = 0.347$). That is, for images that produced high early phase
300 responses in cIT and aIT, the following later phase responses of units in the lower level area (pIT)
301 tended to be low, consistent with error coding models which posit that feedback from higher areas (in
302 the form of predictions) would contribute to the decreasing selectivity observed in lower areas.

303

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

305 We next proceeded to formalize the conceptual ideas introduced in **Figure 1B** and build neurally
306 mechanistic, dynamical models of gradually increasing complexity to determine the minimal set of
307 assumptions that could capture our empirical findings of non-trivial, dynamic selectivity changes during

308 face detection across face-selective subregions in IT. Previous functional and anatomical data show
309 that the face-selective subregions in IT are connected forming an anterior to posterior hierarchy and
310 show that pIT serves as the primary input into this hierarchy (Moeller et al., 2008)(Freiwald & Tsao,
311 2010)(Grimaldi et al., 2016). Thus, we evaluated dynamics in different hierarchical architectures using a
312 linear dynamical systems modeling framework where pIT, cIT, and aIT act as sequential stages of
313 processing (**Figure 5** and see **Methods**). A core principle of feedforward ventral stream models is that
314 object selectivity is built by stage-wise feature integration in a manner that leads to relatively low
315 dimensional representations at the top of the hierarchy abstracted from the high-dimensional input
316 layer. We were interested in how signals temporally evolve across a similar architectural layout. We
317 used the simplest feature integration architecture where a unit in a downstream area linearly sums the
318 input from units in an upstream area, and we stacked this computation to form three layer networks
319 (**Figure 5B**). This simple, generic feedforward encoding model conceptualizes the idea that different
320 types of evidence, local and global (i.e. information about the parts and the relative spatial arrangement
321 of parts), have to converge and be integrated to separate face from non-face images in our image set.
322 We used linear networks as monotonic nonlinearities can be readily accommodated in our framework
323 (Seung, 1997)(Rao & Ballard, 1999)(also see **Figure 7C**). Importantly, we used a simple encoding
324 scheme as our goal was not to build full-scale deep neural network encoding models of image
325 representation (Yamins et al., 2014) but to bring focus to an important biological property that is often
326 not considered in deep nets, neural dynamics.

327 We implemented a range of ideas previously proposed in the literature. The functional
328 implications of these ideas were highlighted in **Figure 1B**, but at a mechanistic level, these functional
329 properties can be directly realized via different recurrent processing motifs between neurons (**Figure 5**,
330 base feedforward model (first column) is augmented with recurrent connections to form new models
331 (remaining columns)). For example, self-connections can be viewed as implementing spike rate
332 adaptation in a feedforward architecture (**Figure 5A**, top row), lateral connections support winner-take-

333 all or normalization mechanisms (Carandini et al., 1997) (**Figure 5A**, second, third, and fourth
334 columns), and top-down connections implement Bayesian inference (Seung, 1997) (**Figure 5A**, fifth
335 and sixth columns). Here, normalization is implemented by a leak term that scales adaptation (the
336 degree of decay in the response) and is controlled recursively by the summed activity of the network
337 (Carandini et al., 1997), ultimately scaling down responses to strong driving stimuli over time. To
338 constrain our choice of a feedback-based model, we took a normative approach minimizing a quadratic
339 reconstruction cost between stages as the classical reconstruction cost is at the core of an array of
340 hierarchical generative models including hierarchical Bayesian inference (Lee & Mumford, 2003),
341 Boltzmann machines (Ackley, Hinton, & Sejnowski, 1985), analysis-by-synthesis networks (Seung,
342 1997), sparse coding (Olshausen & Field, 1996), predictive coding (Rao & Ballard, 1999), and
343 autoencoders in general (Rifai, Vincent, Muller, Glorot, & Bengio, 2011). Optimizing a quadratic loss
344 results in feedforward and feedback connections that are symmetric -- reducing the number of free
345 parameters -- such that inference on the represented variables at any intermediate stage is influenced
346 by both bottom-up sensory evidence and current top-down interpretations. Critically, a common feature
347 of this large model family is the computation of between-stage error signals via feedback, which is
348 distinct from state-estimating model classes (i.e. feedforward models) that do not compute or propagate
349 errors. A dynamical implementation of such a network uses leaky integration of error signals which, as
350 shared computational intermediates, guide gradient descent of the values of the represented variables
351 to a previously learned target value (Δ activity of each neuron => online inference) or descend the
352 connection weights to values that give the best future behavior (Δ synaptic strengths => offline learning),
353 here defined as an unsupervised reconstruction goal (similar results were found using other goals and
354 networks such as supervised discriminative networks; see **Figure 7C**).

355 When we fit each of the models to our neural data, they generally produced an increase in
356 selectivity from the first stage of the network to the later stages of the network. This increase is not
357 surprising because the models had built-in converging feedforward connections from the first to second

358 to third stages (**Figure 6A**, first five columns, compare blue to red curves). However, discrepancies
359 between models emerged in the lower stages where we found that neither the lateral inhibition model,
360 nor the normalization model, could capture the decreasing selectivity phenomenon observed in pIT.
361 Instead, the selectivity of these models simply increased to a saturation level set by the leak term
362 (shunting inhibition) in the system (**Figure 6A**, first five columns). Similar behavior was present when
363 we tried a nonlinear implementation of the normalization model that more powerfully modulated
364 shunting inhibition (Carandini et al., 1997). That normalization proved insufficient to generate the
365 observed neural dynamics can be explained by the fact that the normalized response to a stimulus
366 cannot easily fall below the response to a stimulus that was initially similar in strength. Thus, a
367 decreasing average preference for a stimulus across a population of cells (i.e. **Figures 2-4**, pIT data)
368 for similar levels of average input is difficult when only using a basic normalization model mediated by
369 surround (within-stage) suppression.

370

371 In contrast to the above models, we found that the feedback model capable of computing
372 hierarchical error signals naturally displayed a strong decrease of selectivity in a sub-component of its
373 first processing stage -- qualitatively similar behavior to the selectivity decrease that we observed in
374 many pIT and cIT neural sites. Specifically, this model displayed these dynamics in the magnitude of its
375 reconstruction error signals but not in its state signals (the feature values) (**Figure 6A**, compare fifth
376 and sixth columns). These error signals integrate converging state signals from two stages -- one
377 above (prediction) and one below (sensory evidence). The term "error" is thus meaningful in the hidden
378 processing stages where state signals from two stages can converge. The top nodes of a hierarchy
379 receive little descending input and hence do not carry additional errors with respect to the desired
380 computation; rather, top nodes convey the face predictions that influence errors in bottom nodes. This
381 behavior in the higher processing stages is consistent with our observation of explicit representation of
382 faces in aIT in all phases of the response (**Figures 2-3**) and with similar observations of decodable

383 identity signals by others in all phases of aIT responses for faces (Meyers, Borzello, Freiwald, & Tsao,
384 2015) and objects (Hung et al., 2005)(Majaj et al., 2015). We also found similar error dynamics when
385 using a simpler two-layer network as opposed to three layers suggesting that these error signal
386 dynamics along with prediction signals emerge even in the simplest cascaded architecture (**Figure 7A**).

387

388 **Predictions of an error coding hierarchical model**

389 While our neural observations at multiple stages of the IT hierarchy led us to the error coding
390 hierarchical model above, a stronger test of the idea of error signaling is whether it predicts other IT
391 neural phenomena. To identify stimulus regimes that would lead to insightful predictions, we asked in
392 what way would the behavior of error-estimating hierarchical models differ most from the behavior of
393 generic feedforward state-estimating models. Because our feedback-based model uses feedforward
394 inference at its core, it behaves similarly to a state-estimating hierarchical feedforward model when the
395 statistics of inputs match the learned feedforward weight pattern of the network (i.e. 'natural' images
396 drawn from everyday objects and scenes) since for these inputs, feedforward inferences derived from
397 the sensory data are aligned with top-down expectations. Thus, predictions of feedback-based models
398 that could distinguish them from feedforward-only models are produced when the natural statistics of
399 images are altered so that they differ from the feedforward patterns previously learned by the network
400 and hence differ from the predictions generated in the network. We have (above) considered one such
401 type of alteration: images where local face features are present but altered from their naturally
402 occurring (i.e. statistically most likely) arrangement in frontal faces. Next, we tested two other image
403 manipulations from recent physiology studies which yielded novel neural phenomena that lacked a
404 principled, model-based explanation (Freiwald, Tsao, & Livingstone, 2009)(Meyer et al., 2014). To test
405 whether the error coding hierarchical model family displays these behaviors, we fixed the architectural
406 parameters derived from our fitting procedure in **Figure 6** and simply varied the input to this network,
407 specifically the correlation between the inputs and the network's feedforward weight pattern, in order to

408 match the nature of the image manipulations performed in prior experiments.

409

410 *Sublinear integration of the face features.* In the face-selective subregion of cIT, the sum of the
411 responses to the face parts exceeds the response to the whole face when averaging firing rates over a
412 200 ms window (Freiwald et al., 2009; see their Figure 2C), and we observed a similar phenomenon in
413 the face-selective subregion in pIT (Issa & DiCarlo, 2012). Interestingly, examining the dynamics of part
414 integration more closely within the first 100ms of the response reveals that the response to the whole
415 face does begin at a level similar to the sum of the responses to the face parts but becomes sublinear
416 in the late phase (whole face response relatively low compared to linear prediction from parts
417 responses) (ratio of sum of responses to parts vs. response to whole: 60-90 ms = 1.5 ± 0.1 , 100-130
418 ms = 4.6 ± 0.3 ; $p = 0.000$, $n = 33$ sites) (**Figure 8A**, left panel). This result runs counter to what would
419 be expected in a model where selectivity for the whole face is built from the conjunction of the parts. In
420 such a model, the population response to the whole face would be at least as large if not greater
421 (superlinear) than the summed responses to the individual features. To test whether an error coding
422 model exhibited the phenomenon of sublinear feature integration at the population level, we compared
423 the response with all inputs active (co-occurring features) to the sum of the responses when each input
424 was activated independently (individual features). The reconstruction errors in our feedback-based
425 model showed a strong degree of sublinear integration of the inputs such that the response to the
426 simultaneous inputs (whole) was much smaller than what would be predicted by a linear sum of the
427 responses to each input alone (parts), and the model's sublinear integration behavior qualitatively
428 replicated the time course observed in pIT without any additional fitting of parameters (**Figure 8A**, right
429 panel). Although we certainly expect that similar sublinear integration may also be observed in a
430 normalization model, this would require a particularly strong form of normalization since population
431 activity to the whole face would have to be normalized to nearly the same level as that for an individual
432 part in the late response phase (ratio of response to single part vs. response to whole = 0.92 ± 0.06 ,

433 100-130 ms, $n = 33$ sites) despite being three times larger during the early response phase (ratio of
434 response to single part vs. response to whole = 0.3 ± 0.02 , 60-90 ms, $n = 33$ sites). Furthermore, the
435 stacked normalization model could not account for the main dynamical phenomenon that we observed
436 (**Figure 6**); therefore, an error coding perspective may provide a more parsimonious account of the
437 observed set of dynamical phenomena.

438

439 *Evolution of neural signals across time.* Neural responses to familiar images are known to rapidly
440 attenuate in IT when compared to responses to novel images (Freedman, Riesenhuber, Poggio, &
441 Miller, 2006)(Woloszyn & Sheinberg, 2012)(Meyer & Olson, 2011). This observation seems to
442 contradict what would be predicted by simple Hebbian potentiation for the more exposed stimuli.
443 Furthermore, familiar image responses show much sharper temporal dynamics than responses to novel
444 images when presented repeatedly (Meyer et al., 2014). These qualitatively different dynamics for
445 familiar versus novel images are surprising given that stimuli are drawn from the same distribution of
446 natural images and are thus matched in their average image-level statistical properties (color, spatial
447 frequency, contrast). To test whether our network displayed these different dynamical behaviors, we
448 simulated familiar inputs as those that match the learned weight pattern of a high-level detector and
449 novel inputs as those with the same overall input level but with weak correlation to the learned network
450 weights (here, we have extended the network to include two units in the output stage corresponding to
451 storage of the two familiarized input patterns to be alternated; conceptually, we consider these familiar
452 pattern detectors as existing downstream of IT in a region such as perirhinal cortex which has been
453 shown to code familiarized image statistics and memory-based object signals (Murray, Bussey, &
454 Saksida, 2007)). We repeatedly alternated two familiar inputs or two novel inputs and found that error
455 coding model responses in the hidden processing stage were temporally sharper for familiar inputs that
456 matched the network's feedforward weight patterns compared to novel patterns of input (pseudo-
457 randomly drawn; see **Methods**), consistent with the previously observed phenomenon (**Figure 8B**; data

458 reproduced with permission from Meyer et al., 2014). Model responses reproduced additional details of
459 the neural dynamics including a large initial peak followed by smaller peaks for responses to novel
460 inputs and a phase delay in the oscillations of responses to novel inputs compared to familiar inputs.
461 Intuitively, these dynamics are composed of two phases after the initial response transient to the onset
462 of the image sequence. In the first phase, familiar patterns lead to lower errors and hence lower neural
463 responses than random patterns (**Figure 8B**, red curve drops below the black curve after the onset
464 response), similar to the observed weaker response to more familiar face-like images present in our
465 data (**Figure 2**, red curves drop below black curves in pIT example sites). When the familiar pattern *A*
466 is switched to another familiar pattern *B*, this induces a short-term error in adjusting to the new pattern
467 (**Figure 8B**, red curve briefly goes above the black curve during pattern switch and then decreases). In
468 contrast, two unfamiliar patterns are closer together in the high-level encoding space than two learned
469 patterns (**Figure 8B**, inset at right), and the switch between two unlearned patterns introduces relatively
470 less shift in top-down signals and hence a smaller dynamical change in error signals. This result
471 demonstrates that our model, derived from fitting only the first 70 ms (60-130 ms post image onset) of
472 IT responses to face images, can extend to much longer timescales and may generalize to studies of
473 images besides face images.

474

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

476 In the large family of state-error coding hierarchical networks, a number of different cortical circuits are
477 possible. A key distinction of two such circuit mapping hypotheses (predictive coding versus error
478 backpropagation) is the expected laminar location of state coding neurons transmitting information
479 about features in the image. In typical neural network implementations, the feedforward projecting
480 neurons in superficial lamina are presumed to encode estimates about states of the visual world (e.g.
481 presence of a face). In contrast, predictive coding posits that superficial layers contain error units and
482 that errors are projected forward to the next cortical level (Rao & Ballard, 1999)(Friston & Kiebel,

483 2009)(Hyvärinen, Hurri, & Hoyer, 2009). State and error signals can be distinguished by their dynamical
484 signatures in our leading model, which was fit on error signals but produces predictions of the
485 corresponding state signals underlying the generation of errors. Since state units are integrators (see
486 **Methods**), they have slower dynamics than error units leading to longer response latencies and a
487 milder decay in responses (**Figure 8C**, red curve in right panel). To test this prediction, we localized our
488 recordings relative to the cortical mantle by co-registering the x-ray determined locations of our
489 electrode (~400 micron *in vivo* accuracy; (Issa et al., 2010)) to structural MRI data (see **Methods**).
490 When we separated units into those at superficial depths closer to the pial surface (1/3 of our sites;
491 corresponds to approximately 0 to 1 mm in depth) versus those in the deeper layers (remaining 2/3 of
492 sites, ~1 to 2.5 mm in depth), we found a longer latency and less response decay in superficial units
493 consistent with the expected profile of state units (**Figure 8C**, left panel). Thus, the trend toward state-
494 like signals in superficial layers is more consistent with typical error backpropagation models (states fed
495 forward, errors fed backward) than with predictive coding proposals. In fact, the latency difference
496 between cortical lamina in pIT (deep vs superficial: 66.0 ± 1.7 vs 76.0 ± 1.8 ms, $p = 0.002$) was greater
497 than the conduction delay from pIT to cIT (i.e. from superficial layers of pIT to the deeper layers of cIT)
498 (superficial pIT vs deep cIT: 76.0 ± 1.8 vs 75.5 ± 2.0 ms, $p = 0.15$) even though laminar distances
499 within pIT are smaller than the distance traveled between cortical stages pIT and cIT. Thus, instead of a
500 simple conduction delay accounting for latency differences across lamina, our model suggests that
501 temporal integration of inputs in superficial lamina, more consistent with the behavior of state units as
502 opposed to error units, may drive the lagged dynamical properties of neurons in superficial lamina.

503

504 **DISCUSSION**

505 We have measured neural responses during a difficult frontal face detection task across the IT
506 hierarchy and demonstrated that the population preference for faces in the intermediate (a.k.a hidden)
507 processing stages decreases over time – that is population responses at lower levels of the hierarchy

508 (pIT and cIT) rapidly evolved toward preferring unnatural, non-face part arrangements whereas the top
509 level (aIT) rapidly developed and then maintained a preference for natural, frontal face part
510 arrangements. The relative speed of selectivity changes in pIT (~30 ms) makes high-level explanations
511 based on fixational eye movements or shifts in attention (e.g. from behavioral surprise to unnatural
512 arrangements of face parts) unlikely as saccades and attention shifts occur on slower timescales
513 (hundreds of milliseconds) (Egeth & Yantis, 1997). The presence of stronger responses to face images
514 than to non-face images in aIT further argues against general arousal effects as these would have been
515 expected to cause stronger responses to challenging non-face images in aIT. Rather, the rapid
516 propagation of neural signals over tens of milliseconds suggested intracortical processing within the
517 ventral visual stream in a manner that was not entirely consistent with a pure feedforward model, even
518 when we included strong nonlinearities in these models such as normalization and even when we
519 stacked these operations to form more complex three stage models. However, augmenting the
520 feedforward model so that it represented the errors generated during hierarchical processing produced
521 the observed neural dynamics and hierarchical signal propagation (**Figures 6-7**). This view argues that
522 many IT neurons code error signals. Using this new modeling perspective, we went on to generate
523 predictions of previously observed IT neural phenomena (**Figure 8**).

524

525 **Comparison to previous neurophysiology studies in IT**

526 Our suggestion that many IT neurons code errors is consistent with the observation of strong
527 responses to extremes in face space (Leopold, Bondar, & Giese, 2006) providing an alternative
528 interpretation to the prior suggestion that cIT neurons are not tuned for typical faces but are instead
529 tuned for atypical face features (i.e. extreme feature tuning) (Freiwald et al., 2009). In that prior work,
530 the response preference of each neuron was determined by averaging over a long time window (~200
531 ms). By looking more closely at the fine time scale dynamics of the IT response, we suggest that this
532 same extreme coding phenomenon can instead be interpreted as a natural consequence of networks

533 that have an actual tuning preference for typical faces (as evidenced by an initial response preference
534 for typical frontal faces in pIT, cIT, and aIT; **Figure 3B**) but that also compute error signals with respect
535 to that preference. Under the present hypothesis, some IT neurons are preferentially tuned to typical
536 spatial arrangements of face features, and other IT neurons are involved in coding errors with respect
537 to those typical arrangements. We speculate that these intermixed state estimating and error coding
538 neuron populations are both sampled in standard neural recordings of IT, even though only state
539 estimating neurons are truly reflective of the tuning preferences of that IT processing stage. This
540 intermixed view of IT neural signaling is further supported by recent studies demonstrating correlates of
541 temporal prediction errors to image sequences in IT (Meyer & Olson, 2011) including the face-selective
542 subregion of cIT (Schwiedrzik & Freiwald, 2017).

543

544 The precise fractional contribution of errors to total neural activity is difficult to estimate from our
545 data. Under the primary image condition tested, not all sites significantly decreased their selectivity
546 (~60% did not change their selectivity). We currently interpret these sites as coding state (feature)
547 estimates (**Figure 3A**, light and dark gray lines in top and bottom rows, respectively), and we did
548 observe evidence of emergence of state-like signals in our superficial neural recordings (**Figure 8C**).
549 Alternatively, at least some of the non-reversing sites might be found to code errors under other image
550 conditions than the one that we tested. Furthermore, while in our primary image condition selectivity
551 decreases only accounted for ~15% of the overall spiking modulation (**Figure 6A**, data panel), larger
552 modulations in late phase neural firing (50-100%) are possible under other image conditions (**Figure**
553 **8A,B**). At a computational level, the absolute contribution of error signals to spiking may not be the
554 critical factor as even a small relative contribution may have important consequences in the network.

555

556 **Comparison across dynamical models of neural processing**

557 Our goal was to test a range of existing recurrent models by recording neural dynamics across multiple

558 cortical stages which provided stronger constraints on computational models than fitting neural
559 responses from only one area as in prior work (Carandini et al., 1997)(Rao & Ballard, 1999). Crucially,
560 we found that the multi-stage neural dynamics observed in our data could not be adequately fit by only
561 using recurrences within a stage such as adaptation, lateral inhibition, and standard forms of
562 normalization (**Figure 6**). These results did not change when we made our simple networks more
563 complex by adding more stages (compare **Figure 6** versus **Figure 7**) or by using more realistic model
564 units with monotonic nonlinearities similar to a spiking nonlinearity (data not shown). Indeed, we
565 specifically chose our stimuli to evoke similar levels of within stage neural activity to limit the effects of
566 known mechanisms that depend on activity levels within an area (e.g. adaptation, normalization), and
567 we fully expect that these activity dependent mechanisms would operate in parallel to top-down,
568 recurrent processes during general visual processing. We emphasize that we only tested the standard
569 form of normalization as originally proposed, using within stage pooling and divisive mechanisms
570 (Carandini et al., 1997). Since that original mechanistic formulation, normalization has evolved to
571 become a term that broadly encapsulates many forms of suppression phenomena and can include both
572 lateral interactions within an area and feedback interactions from other areas (Nassi, Gómez-Laberge,
573 Kreiman, & Born, 2014)(Coen-Cagli, Kohn, & Schwartz, 2015). Thus, while our results do not follow
574 from the original mechanistic form of normalization, they may yet fall under normalization more broadly
575 construed as a term for suppression phenomena (error coding would require a similar suppressive
576 component). Here, we have provided a normative model for how top-down suppression would follow
577 from the well-defined computational goals of many hierarchical neural network models.

578

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

580 The notion of error is inherent to many existing models in the literature that go beyond the basic
581 feedforward, feature estimation class. Example models use errors for guiding top-down inference by
582 computing errors implicitly (hierarchical Bayesian inference (Seung, 1997)(Lee & Mumford, 2003)) or by

583 representing errors explicitly (predictive coding (Rao & Ballard, 1999)). In addition, errors can be used
584 specifically for driving unsupervised learning (autoencoder (Rifai et al., 2011)) or supervised learning
585 (classic error backpropagation (Williams & Hinton, 1986)). Recently, models have incorporated aspects
586 of both inference and learning (Salakhutdinov & Hinton, 2012)(Patel, Nguyen, & Baraniuk, 2015). A
587 key, unifying feature across inference and learning models is the need to compute an error signal
588 between processing stages. This error signal can be in the form of a generative, reconstruction cost
589 (stage n predicting stage $n-1$) or a discriminative, construction cost (stage $n-1$ predicting stage n).
590 Across-stage “performance” error terms are used in all model cost functions, are typically the only term
591 combining signals from different model stages, and are distinct from within-stage “regularization” terms
592 (i.e. sparseness or weight decay) in driving network behavior (Marblestone, Wayne, & Kording, 2016).
593 The present study provides evidence that such errors are not only computed, but that they are explicitly
594 encoded in spiking rates. We emphasize that this result at the level of population dynamics was robust
595 across choices of cost function such as those used in the literature; we tested models with different
596 unsupervised and supervised performance errors (reconstruction, nonlinear reconstruction, and
597 discriminative) and found similar population level error signals across these networks in the basic two-
598 layer implementation (**Figure 7C**). Thus, errors as generally instantiated in the state-error coding
599 hierarchical model family provide a good approximation to IT population neural dynamics.

600

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

602 In error-computing networks, errors provide control signals for guiding learning giving these networks
603 additional adaptive power over basic feature estimation networks. This property helps augment the
604 classical, feature coding view of neurons which, with only feature activations and Hebbian operations,
605 does not lead to efficient learning in the manner produced by gradient descent using error
606 backpropagation (Williams & Hinton, 1986). Observation of error signals may provide insight into how
607 more intelligent unsupervised and supervised learning algorithms such as backpropagation could be

608 plausibly implemented in the brain. A potentially important contribution of this work is the suggestion
609 that gradient descent algorithms are facilitated by using an error code so that efficient learning is
610 reduced to a simple Hebbian operation at synapses and efficient inference is simply integration of
611 inputs at the cell body (see *eqn 10* and text in **Methods**). This representational choice, to code the
612 computational primitives of gradient descent in spiking activity, would simply leverage the existing
613 biophysical machinery of neurons for inference and learning.

614

615 **MATERIALS & METHODS**

616 **Animals and surgery.** All surgery, behavioral training, imaging, and neurophysiological techniques are
617 identical to those described in detail in previous work (Issa & DiCarlo, 2012). Two rhesus macaque
618 monkeys (*Macaca mulatta*) weighing 6 kg (Monkey 1, female) and 7 kg (Monkey 2, male) were used. A
619 surgery using sterile technique was performed to implant a plastic fMRI compatible headpost prior to
620 behavioral training and scanning. Following scanning, a second surgery was performed to implant a
621 plastic chamber positioned to allow targeting of physiological recordings to posterior, middle, and
622 anterior face patches in both animals. All procedures were performed in compliance with National
623 Institutes of Health guidelines and the standards of the MIT Committee on Animal Care and the
624 American Physiological Society.

625

626 **Behavioral training and image presentation.** Subjects were trained to fixate a central white fixation
627 dot during serial visual presentation of images at a natural saccade-driven rate (one image every 200
628 ms). Although a 4° fixation window was enforced, subjects generally fixated a much smaller region of
629 the image ($<1^\circ$) (Issa & DiCarlo, 2012). Images were presented at a size of 6° except for control tests at
630 3° and 12° sizes (**Figure 4C**), and all images were presented for 100 ms duration with 100 ms gap
631 (background gray screen) between each image. Up to 15 images were presented during a single
632 fixation trial, and the first image presentation in each trial was discarded from later analyses. Five

633 repetitions of each image in the general screen set were presented, and ten repetitions of each image
634 were collected for all other image sets. The screen set consisted of a total of 40 images drawn from
635 four categories (faces, bodies, objects, and places; 10 exemplars each) and was used to derive a
636 measure of face versus non-face object selectivity. Following the screen set testing, some sites were
637 tested using an image set containing images of face parts presented in different combinations and
638 positions (**Figure 1B**, left panel). We first segmented the face parts (eye, nose, mouth) from a monkey
639 face image. These parts were then blended using a Gaussian window, and the face outline was filled
640 with pink noise to create a continuous background texture. A face part could appear on the outline at
641 any one of nine positions on an evenly spaced 3x3 grid. Although the number of possible images is
642 large ($4^9 = 262,144$ images), we chose a subset of these images for testing neural sites ($n = 82$
643 images). Specifically, we tested the following images: the original whole face image, the noise-filled
644 outline, the whole face reconstructed by blending the four face parts with the outline, all possible single
645 part images where the eye, nose, or mouth could be at one of nine positions on the outline ($n = 3 \times 9 =$
646 27 images), all two part images containing a nose, mouth, left eye, or right eye at the correct outline-
647 centered position and an eye tested at all remaining positions ($n = 4 \times 8 - 1 = 31$ images), all two part
648 images containing a correctly positioned contralateral eye while placing the nose or mouth at all other
649 positions ($n = 2 \times 8 - 2 = 14$ images), and all correctly configured faces but with one or two parts missing
650 besides those already counted above ($n = 4 + 3 = 7$ images). The particular two-part combinations
651 tested were motivated by prior work demonstrating the importance of the eye in early face processing
652 (Issa & DiCarlo, 2012), and we sought to determine how the position of the eye relative to the outline
653 and other face parts was encoded in neural responses. The three and four part combinations were
654 designed to manipulate the presence or absence of a face part for testing the integration of face parts,
655 and in these images, we did not vary the positions of the parts from those in a naturally occurring face.
656 In a follow-up test on a subset of sites, we permuted the position of the four face parts under the
657 constraint that they still formed the configuration of a naturally occurring face (i.e. preserve the 'T'

658 configuration, $n = 10$ images; **Figure 4B**). We tested single part images at 3° and 12° sizes in a subset
659 of sites ($n = 27$ images at each size; **Figure 4C**). Finally, we measured the responses to the individual
660 face parts in the absence of the outline ($n = 4$ images; **Figure 8A**).

661

662 **MR Imaging and neurophysiological recordings.** Both structural and functional MRI scans were
663 collected in each monkey. Putative face patches were identified in fMRI maps of face versus non-face
664 object selectivity in each subject. A stereo microfocal x-ray system (Cox et al., 2008) was used to guide
665 electrode penetrations in and around the fMRI defined face-selective subregions of IT. X-ray based
666 electrode localization was critical for making laminar assignments since electrode penetrations are
667 often not perpendicular to the cortical lamina when taking a dorsal-ventral approach to IT face patches.
668 Laminar assignments of recordings were made by co-registering x-ray determined electrode
669 coordinates to MRI where the pial-to-gray matter border and the gray-to-white matter border were
670 defined. Based on our prior work estimating sources of error (e.g. error from electrode tip localization
671 and brain movement), registration of electrode tip locations to MRI brain volumes has a total of <400
672 micron error which is sufficient to distinguish deep from superficial layers (Issa et al., 2013). Multi-unit
673 activity (MUA) was systematically recorded at 300 micron intervals starting from penetration of the
674 superior temporal sulcus such that all sites at these regular intervals were tested with a screen set
675 containing both faces and non-face objects, and a subset of sites that were visually driven were further
676 tested with our main image set manipulating the position of face parts. Although we did not record
677 single-unit activity, our previous work showed similar responses between single-units and multi-units on
678 images of the type presented here (Issa & DiCarlo, 2012), and our results are consistent with
679 observations in previous single-unit work in IT (Freiwald et al., 2009). Recordings were made from PL,
680 ML, and AM in the left hemisphere of monkeys 1 and 2 and additionally from AL in monkey 2. AM and
681 AL are pooled together in our analyses forming the aIT sample while PL and ML correspond to the pIT
682 and cIT samples, respectively.

683

684 **Neural data analysis.** The face patches were physiologically defined in the same manner as in our
685 previous study (Issa & DiCarlo, 2012). Briefly, we fit a graded 3D sphere model (linear profile of
686 selectivity that rises from a baseline value toward the maximum at the center of the sphere) to the
687 spatial profile of face versus non-face object selectivity across our sites. We tested spherical regions
688 with radii from 1.5 to 10 mm and center positions within a 5 mm radius of the fMRI-based centers of the
689 face patches. The resulting physiologically defined regions were 1.5 to 3 mm in diameter. Sites which
690 passed a visual response screen (mean response in a 60-160 ms window $>2 \times \text{SEM}$ above baseline for
691 at least one of the four categories in the screen set) were included in further analysis. All firing rates
692 were baseline subtracted using the activity in a 25-50 ms window following image onset averaged
693 across all repetitions of an image. Finally, given that the visual response latencies in monkey 2 were on
694 average 13 ms slower than those in monkey 1 for corresponding face-selective regions, we applied a
695 single latency correction (13 ms shift to align monkey 1 and monkey 2's data) prior to averaging across
696 monkeys. This was done so as not to wash out any fine timescale dynamics by averaging. Similar
697 results were obtained without using this latency correction as dynamics occurred at longer timescales
698 (~ 30 ms). This single absolute adjustment was more straightforward than the site-by-site adjustment
699 used in our previous work (Issa & DiCarlo, 2012) (though similar results were obtained using this
700 alternative latency correction); even when each monkey was analyzed separately, we still observed pIT
701 selectivity dynamics (**Figure 4A**). Images that produced an average population response ≥ 0.9 of the
702 initial response (60-100 ms) to a face image with all face parts arranged in their typical positions in a
703 frontal face were analyzed further (**Figures 2 and 3**). Stimulus selection was intended to limit
704 potentially confounding differences in visual drive between image classes. In a control test, we also
705 repeated our analysis by selecting images on a site-by-site basis where images with frontal face and
706 non-face arrangements of parts were chosen to be within 0.75x to 1.25x of the initial response to the
707 complete face image (minimum of five face and five non-face images in this response range for

708 inclusion of site in analysis). In follow-up analyses of population responses, we specifically limited
709 comparison to images with the same number of parts (**Figure 4B,C**). For example, for single part
710 images, we used the image with the eye in the upper, contralateral region of the outline as a reference
711 requiring a response ≥ 0.9 of the initial population response to this reference for inclusion of the images
712 in this analysis. We found that four other images of the 27 single-part images elicited a response at
713 least as large as 90% of the response to this standard image. For images containing all four face parts,
714 we used the complete, frontal face as the standard and found non-face arrangements of the four face
715 parts that drove at least 90% of the early response to the whole face (2 images out of 10 tested). To
716 compute individual site d' for each of these stimulus partitions (e.g. typical versus atypical
717 arrangements of 1 face part), we combined all presentations of images with frontal face arrangements
718 and compared these responses to responses from all presentations of images with non-face
719 arrangements using $d' = (u_1 - u_2) / ((\text{var}_1 + \text{var}_2) / 2)^{1/2}$ where variance was computed across all trials for that
720 image class (e.g. all presentations of all typical face images); this was identical to the d' measure used
721 in previous work for computing selectivity for faces versus non-face objects (Aparicio, Issa, & DiCarlo,
722 2016; Ohayon, Freiwald, & Tsao, 2012). For example, for the main image set (**Figure 2A**), we
723 compared all presentations of frontal face arrangements (8 images x 10 presentations/image = 80 total
724 presentations) to all presentations of non-face arrangements (13 images x 10 presentations/image =
725 130 total presentations) to compute the d' values for each site in two time windows (60-90 ms and 100-
726 130 ms) as shown in **Figure 3A**. A positive d' implies a stronger response to more naturally occurring
727 frontal face arrangements of face parts while a negative d' indicates a preference for unnatural non-
728 face arrangements of the face parts.

729

730 **Dynamical models**

731 *Modeling framework and equations.* To model the dynamics of neural response rates in a hierarchy, we
732 start with the simplest possible model that might capture those dynamics: a model architecture

733 consisting of a hidden stage of processing containing two units that linearly converge onto a single
734 output unit. We use this two-stage cascade for illustration of the basic concepts which can be easily
735 extended to longer cascades with additional stages, and we ultimately used a three-stage version of the
736 model to fit our neural data collected from three cortical stages (**Figures 5B & 6**).

737 An external input is applied separately to each hidden stage unit, which can be viewed as
738 representing different features for downstream integration. We vary the connections between the two
739 hidden units within the hidden processing stage (lateral connections) or between hidden and output
740 stage units (feedforward and feedback connections) to instantiate different model families. The details
741 of the different architectures specified by each model class can be visualized by their equivalent neural
742 network diagrams (**Figure 5**). Here, we provide a basic description for each model tested. All two-stage
743 models utilized a 2x2 feedforward identity matrix A that simply transfers inputs \mathbf{u} (2x1) to hidden layer
744 units \mathbf{x} (2x1) and a 1x2 feedforward vector B that integrates hidden layer activations \mathbf{x} into a single
745 output unit y .

746

$$747 \quad A = aI, \quad B = b[1, 1] \quad (1)$$

748

749 By simply substituting in the appropriate unit vector and weight matrix transforming inputs from one
750 layer to the next for the desired network architecture, this simple two-stage architecture can be
751 extended to larger networks (e.g. see three-stage network diagrams in **Figure 5B**). To generate
752 dynamics in the simple networks below, we assumed that neurons act as leaky integrators of their total
753 synaptic input, a standard rate-based model of a neuron used in previous work (Seung, 1997) (Rao &
754 Ballard, 1999).

755

756 *Pure feedforward.* In the purely feedforward family, connections are exclusively from hidden to output

757 stages through feedforward matrices A and B .

758

$$759 \quad \dot{\mathbf{x}} = A\mathbf{u} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - y/\tau \quad (2)$$

760

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

764

765 *Lateral inhibition.* Lateral connections (matrix with off-diagonal terms) are included and are inhibitory.

766 The scalar k_l sets the relative strength of lateral inhibition versus bottom-up input.

767

$$768 \quad \dot{\mathbf{x}} = A\mathbf{u} - \begin{bmatrix} 0 & k_l \\ k_l & 0 \end{bmatrix} \mathbf{x} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - y/\tau \quad (3)$$

769

770 *Normalization.* An inhibitory term that scales with the summed activity of units within a stage is

771 included. The scalar k_s sets the relative strength of normalization versus bottom-up input.

772

$$773 \quad \dot{\mathbf{x}} = A\mathbf{u} - k_s \sum x \cdot \mathbf{x} - \mathbf{x}/\tau \quad \dot{y} = B\mathbf{x} - k_s y \cdot y - y/\tau \quad (4)$$

774

775 *Normalization (nonlinear)* (Carandini et al., 1997). The summed activity of units within a stage is used to
776 nonlinearly scale shunting inhibition.

777

$$778 \quad \dot{\mathbf{x}} = A\mathbf{u} - \frac{\mathbf{x}}{\tau \sqrt{1 - k_s \sum x}} \quad \dot{y} = B\mathbf{x} - \frac{y}{\tau \sqrt{1 - k_s y}} \quad (5)$$

779

780 Note that this is technically a nonlinear dynamical system, and since the normalization term in equation
781 (5) is not continuously differentiable, we used the fourth-order Taylor approximation around zero in the
782 simulations of equation (5).

783

784 *Feedback (linear reconstruction)*. The feedback-based model is derived using a normative framework
785 that performs optimal inference in the linear case (Seung, 1997) (unlike the networks in equations (2)-
786 (5) which are motivated from a mechanistic perspective but do not directly optimize a squared error
787 performance loss). The feedback network minimizes the cost C of reconstructing the inputs of each
788 stage (i.e. mean squared error of layer n predicting layer $n-1$).

789

$$790 \quad C = \frac{1}{2} (\mathbf{u} - A^T \mathbf{x})^2 + \frac{1}{2} (\mathbf{x} - B^T \mathbf{y})^2 \quad (6)$$

791

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

793

$$794 \quad \frac{\partial C}{\partial \mathbf{x}} = -A(\mathbf{u} - A^T \mathbf{x}) + (\mathbf{x} - B^T \mathbf{y}) \quad \frac{\partial C}{\partial \mathbf{y}} = -B(\mathbf{x} - B^T \mathbf{y}) \quad (7)$$

795

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

798

$$799 \quad \frac{d\mathbf{x}}{dt} = -\frac{\partial C}{\partial \mathbf{x}} = A(\mathbf{u} - A^T \mathbf{x}) - (\mathbf{x} - B^T \mathbf{y}) - \mathbf{x}/\tau \quad (8)$$

800

801
$$\frac{d\mathbf{y}}{dt} = -\frac{\partial C}{\partial \mathbf{y}} = B(\mathbf{x} - B^T \mathbf{y}) - \mathbf{y}/\tau$$

802

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

814

815
$$\dot{\mathbf{x}} = A(\mathbf{u} - A^T \mathbf{x}) - k_{td}(\mathbf{x} - B^T \mathbf{y}) - \mathbf{x}/\tau \quad (9)$$

816

817 *Error signals computed in the feedback model.* In equation (9), inference can be thought of as
818 proceeding through integration of inputs on the dendrites of neuron population \mathbf{x} . In this scenario, all
819 computations are implicit in dendritic integration. Alternatively, the computations in equation (9) can be
820 done in two steps where, in the first step, reconstruction errors are computed (i.e. $\mathbf{e}_0 = \mathbf{u} - A^T \mathbf{x}$, $\mathbf{e}_1 = \mathbf{x} -$
821 $B^T \mathbf{y}$) and explicitly represented in a separate error coding population. These error signals can then be
822 integrated by their downstream target population to generate the requisite update to the state signal of
823 neuron population \mathbf{x} .

824

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

826

827 An advantage of this strategy is that the a state unit now directly receives errors as inputs, and those
828 inputs allow implementation of an efficient Hebbian rule for learning weight matrices (Rao & Ballard,
829 1999) -- the gradient rule for learning is simply a product of the state activation and the input error
830 activation (weight updates obtained by differentiating equation (6) with respect to weight matrices A and
831 B : $\Delta A = \mathbf{x} \cdot \mathbf{e}_0^T$, $\Delta A^T = \mathbf{e}_0 \cdot \mathbf{x}^T$, $\Delta B = y \cdot \mathbf{e}_1^T$, and $\Delta B^T = \mathbf{e}_1 \cdot y$). Thus, the reconstruction errors serve as
832 computational intermediates for both the gradients of online inference mediated by dendritic integration
833 (dynamics in state space, equation (10)) and gradients for offline learning mediated by Hebbian
834 plasticity (dynamics in weight space).

835

836 In order for the reconstruction errors at each layer to be scaled appropriately in the feedback
837 model, we invoke an additional downstream variable z to predict activity at the top stage such that,
838 instead of $\mathbf{e}_2 = y$ which scales as a state variable, we have $\mathbf{e}_2 = y \cdot C^T z$ (**Figure 5A**). This overall model
839 reflects a state and error coding model as opposed to a state only model.

840

841 *Feedback (three-stage)*. For the simulations in **Figures 6 and 8**, three-stage versions of the above
842 equations were used. These deeper networks were also wider such that they began with four input
843 units (\mathbf{u}) instead of only two inputs in the two-stage models. These inputs converged through
844 successive processing stages ($\mathbf{w}, \mathbf{x}, \mathbf{y}$) to one unit at the top node (\mathbf{z}) (**Figure 5B**).

845

846 *Feedback (nonlinear reconstruction)*. To test the generality of our findings beyond a linear
847 reconstruction cost, we simulated feedback-based models which optimized different candidate cost
848 functions proposed for the ventral stream (**Figure 7C**). In nonlinear hierarchical inference,

849 reconstruction is performed using a monotonic nonlinearity with a threshold (th) and bias (bi):

850

$$851 \quad C = \frac{1}{2} (\mathbf{u} - f(A^T \mathbf{x}))^2 + \frac{1}{2} (\mathbf{x} - f(B^T \mathbf{y}))^2, \quad \text{where } f(x) = \tanh(x - th) + bi \quad (11)$$

852

$$853 \quad \dot{\mathbf{x}} = A(\mathbf{u} - f(A^T \mathbf{x}))(1 - \tanh(A^T \mathbf{x} - th)^2) - k_{td}(\mathbf{x} - f(B^T \mathbf{y})) - \mathbf{x}/\tau$$

854 (12)

$$855 \quad \dot{\mathbf{y}} = B(\mathbf{x} - f(B^T \mathbf{y}))(1 - \tanh(B^T \mathbf{y} - th)^2) - \mathbf{y}/\tau$$

856

857 *Feedback (linear construction).* Instead of a reconstruction cost where responses match the input (i.e.
 858 generative model) as in unsupervised learning, we additionally simulated the states and errors in a
 859 feedback network minimizing a linear construction cost where the network is producing responses to
 860 match a given output (i.e. discriminative model) similar to supervised learning:

861

$$862 \quad C = \frac{1}{2} (A\mathbf{u} - \mathbf{x})^2 + \frac{1}{2} (B\mathbf{x} - \mathbf{y})^2 \quad (13)$$

863

$$864 \quad \dot{\mathbf{x}} = (A\mathbf{u} - \mathbf{x}) - k_{td}B^T(B\mathbf{x} - \mathbf{y}) - \mathbf{x}/\tau \quad \dot{\mathbf{y}} = (B\mathbf{x} - \mathbf{y}) - \mathbf{y}/\tau \quad (14)$$

865

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

869

$$870 \quad \mathbf{u}(t) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(t-t_0)^2}{2\sigma^2}} * \mathbf{h}(t) \Rightarrow \dot{\mathbf{u}} = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(t-t_0)^2}{2\sigma^2}} \cdot \mathbf{h} \quad (15)$$

871

872 where the elements of \mathbf{h} are scaled Heaviside step functions. The input is thus a sigmoidal ramp whose
873 latency to half height is set by t_0 and rise time is set by σ . For simulation of two-stage models, there
874 were ten basic parameters: latency of the input t_0 , standard deviation of the Gaussian ramp σ , system
875 time constant τ , input connection strength a , feedforward connection strength b , the four input values
876 across two stimulus conditions (i.e. h_{11} , h_{12} , h_{21} , h_{22}), and a factor sc for scaling the final output to the
877 neural activity. In the deeper three-stage network, there were a total of fifteen parameters which
878 included an additional feedforward connection strength c and additional input values since the three-
879 stage model had four inputs instead of two. The lateral inhibition model class required one additional
880 parameter k_l as did the normalization model family k_s , and for feedback model simulations, there was
881 an additional feedback weight k_{fd} to scale the relative contribution of the top-down errors in driving
882 online inference. For the error coding variants of the feedback model, gain parameters c (two-stage)
883 and d (three-stage) were included to scale the overall magnitude of the top level reconstruction error
884 (also see **Figure 5** for locations of parameters in network diagrams).

885

886 **Model parameter fits to neural data.** In fitting the models to the observed neural dynamics, we
887 mapped the summed activity in the hidden stage (\mathbf{w}) to population averaged activity in pIT, and we
888 mapped the summed activity in the output stage (\mathbf{y}) to population averaged signals measured in aIT. To
889 simulate error coding, we mapped the reconstruction errors $\mathbf{e}_1 = \mathbf{w} - \mathbf{B}^T \mathbf{x}$ and $\mathbf{e}_3 = \mathbf{y} - \mathbf{C}^T \mathbf{z}$ to activity in pIT
890 and aIT, respectively. We applied a squaring nonlinearity to the model outputs as an approximation to
891 rectification since recorded extracellular firing rates are non-negative (and linear rectification is not
892 continuously differentiable). Analytically solving this system of dynamical equations (2)-(14) for a step
893 input is precluded because of the higher order interaction terms (the roots of the determinant and hence
894 the eigenvalues/eigenvectors of a 3x3 or larger matrix are not analytically determined, except for the
895 purely feedforward model which only has first-order interactions), and in the case of the normalization

896 models, there is an additional nonlinear dependence on the shunt term. Thus, we relied on
897 computational methods (constrained nonlinear optimization) to fit the parameters of the dynamical
898 systems to the neural data with a quadratic (sum of squares) loss function.

899

900 Parameter values were fit in a two-step procedure. In the first step, we fit only the difference in
901 response between image classes (differential mode which is the selectivity profile over time, see **Figure**
902 **6A**, right data panel), and in the second step, we refined fits to capture an equally weighted average of
903 the differential mode and the common mode (the common mode is the average across images of the
904 response time course of visual drive). This two-step procedure was used to ensure that each model
905 had the best chance of fitting the dynamics of selectivity (differential mode) as these selectivity profiles
906 were the main phenomena of interest but were smaller in size (20% of response) compared to overall
907 visual drive. In each step, fits were done using a large-scale algorithm (interior-point) to optimize
908 coarsely, and the resulting solution was used as the initial condition for a medium-scale algorithm
909 (sequential quadratic programming) for additional refinement. The lower and upper parameter bounds
910 tested were: $t_0=[50\ 70]$, $\sigma=[0.5\ 25]$, $\tau=[0.5\ 1000]$, $k_i, k_s, k_{\sigma}=[0\ 1]$, $a, b, c, d=[0\ 2]$, $h=[0\ 20]$, $sc=[0\ 100]$, $th=[-$
911 $20\ 20]$, and $bi=[-1\ 1]$ which proved to be adequately liberal as parameter values converged to values
912 that did not generally approach these boundaries. To avoid local minima, the algorithm was initialized to
913 a number of randomly selected points ($n = 50$), and after fitting the differential mode, we took the top
914 fits ($n = 25$) for each model class and used these as initializations in subsequent steps. The single best
915 fitting instance of each model class is shown in the main figures.

916

917 **Model predictions.** For the predictions in **Figure 8**, all architectural parameters obtained by the fitting
918 procedure above were held fixed; only the pattern of inputs to the network was varied. For **Figure 8A**,
919 to test the input integration properties of a model, we used the top-performing model and compared the
920 response to all inputs presented simultaneously with the sum of the responses to each input alone.

921

922 For **Figure 8B**, we approximated novel versus familiar images by parametrically varying the
923 degree to which input patterns were random (inputs drawn from i.i.d. uniform distributions on the
924 interval [0,1]) versus structured in a way that matched the weight pattern of the network. Here, we used
925 a version of the model with two independent outputs reflecting detectors for two familiarized input
926 patterns (output 1 tuned to pattern A: u_1, u_2, u_3, u_4 active and output 2 tuned to pattern B: u_5, u_6, u_7, u_8
927 active) (**Figure 8B**). Alternating between these two input patterns simulates alternation of two
928 familiarized (learned) images as compared to purely random patterns (u_{1-8} independent and identically
929 distributed). The first-to-second layer weights were [1,1,1,1,0,0,0,0] for pattern A and [0,0,0,0,1,1,1,1]
930 for pattern B, so to parametrically vary the degree of correlation of inputs to this weight pattern from
931 random (correlation = 0) to deterministic (correlation = 1), we drew input values from a joint distribution
932 $P(u_1, u_2, u_3, u_4, u_5, u_6, u_7, u_8)$ where u_{1-4} were drawn from a high-valued uniform distribution on the interval
933 $[1-\varepsilon, 1]$ and u_{5-8} were drawn from a low-valued uniform distribution $[0, \varepsilon]$ for stimulus pattern A and the
934 opposite for pattern B (u_{5-8} high-valued and u_{1-4} low-valued). The parameter ε determines the range of
935 values that could be drawn from purely deterministic (0 or 1, $\varepsilon = 0$) to randomly uniformly distributed
936 (from 0 to 1, $\varepsilon = 1$). Thus, the correlation of the inputs correspondingly varies according to $\rho(u_i, u_j)$
937 $= \rho(u_k, u_l) = (\varepsilon-1)^2 / ((\varepsilon-1)^2 + \varepsilon^2 / 3)$ where $1 \leq i, j \leq 4, i \neq j$ and $5 \leq k, l \leq 8, k \neq l$ approaching correlation equal to 0 for a
938 purely, random pattern ($\varepsilon = 1$) that had a low probability of matching the learned patterns A and B.

939

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

942

943 **Statistics.** Error bars represent standard errors of the mean obtained by bootstrap resampling ($n =$
944 1000). All statistical comparisons including those of means or correlation values were obtained by

945 bootstrap resampling ($n = 1000$) producing p-values at a resolution of 0.001 so that the lowest p-value
946 that can be reported is $p = 0.000$ given the resolution of this statistical analysis. All statistical tests were
947 two-sided unless otherwise specified. Spearman's rank correlation coefficient was used.

948

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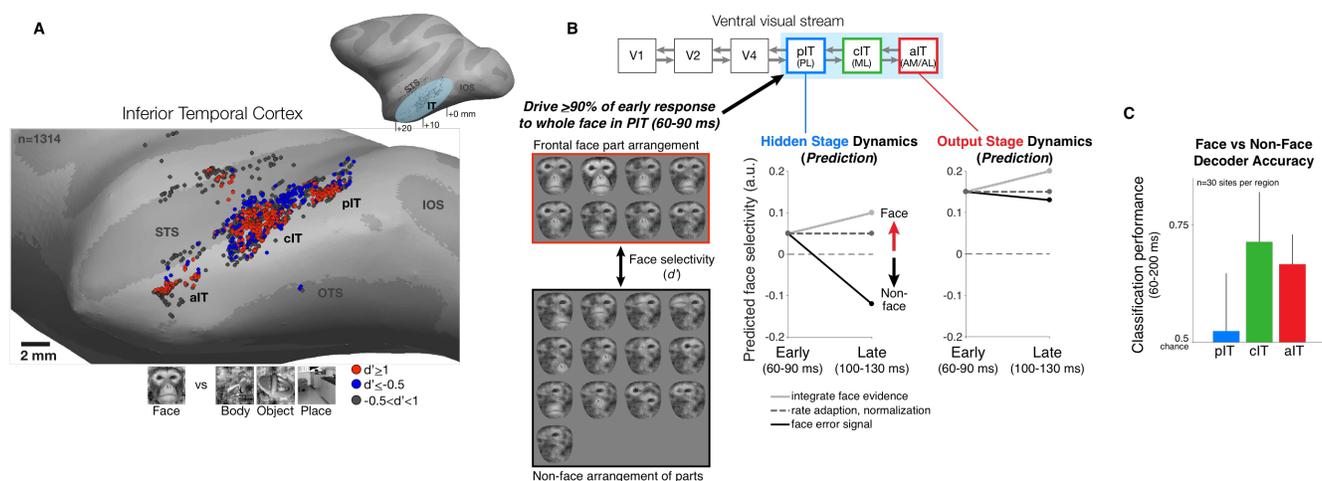
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Figure 1

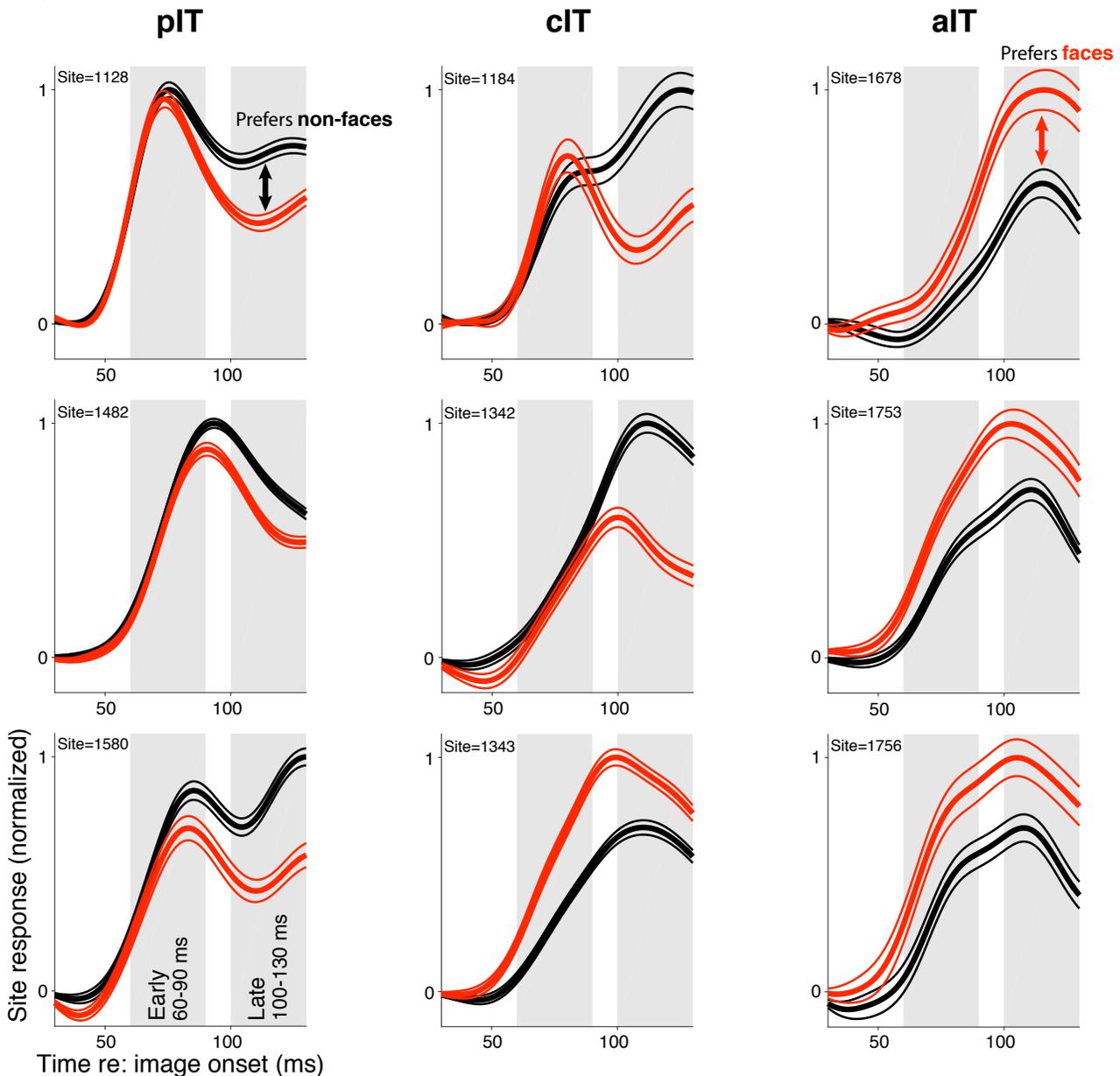


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Figure 1. Neural recordings and experimental design in face-selective subregions of the ventral visual stream. (A) Neurons were recorded along the lateral convexity of the inferior temporal lobe spanning the posterior to anterior extent of IT (+0 to +20 mm AP, Horsely-Clarke coordinates) in two monkeys (data from monkey 1 are shown). Based on prior work, face-selective sites (red) were operationally defined as those with a response preference for images of frontal faces versus images of non-face objects (see **Methods**). While these neurons 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 corresponded to face-selective areas identified under fMRI in the same subjects (Issa & DiCarlo, 2012)(Issa et al., 2013) (STS = superior temporal sulcus, IOS = inferior occipital sulcus, OTS = occipitotemporal sulcus). (B) (top diagram) The three visual processing stages in IT lie downstream of early visual areas V1, V2, and V4 in the ventral visual stream. (left) We designed our stimuli to focus on the intermediate stage pIT by seeking images of faces and images of non-faces that would, on average, drive equally strong initial responses in pIT. 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 frontal face and non-face arrangements of the face parts, and we identified 21 images (red and black boxes) that drove the mean, early (60-100 ms) pIT population response to $\geq 90\%$ of its response to the intact face (first image in red box is synthesized whole face; compare to the second image which is the original whole face), and of these 21 images, 13 images contained non-face arrangements of the face parts. For example, images with an eye centered in the outline (black box, 3rd and 4th rows) as opposed to the lateralized position of the eye in a frontal face (red box) have a global interpretation (“cyclops”) that is not consistent with a frontal face but still evoked strong pIT responses. Selectivity of neural sites (see **Figs. 3 & 4**) for face versus non-face images was quantified using a d' measure. (middle) Computational hypotheses of cortical dynamics make differing predictions about how neural selectivity in pIT may evolve following an initial, weak preference signal for images of frontal faces. (right) Predictions of how aIT would behave as an output stage building selectivity for images of frontal faces through multiple stages of processing. (C) A population decoder, trained on average firing rates (60-200 ms post image onset, linear SVM classifier) for frontal face versus non-face arrangements of the face parts in this image subset, performed poorly in pIT on held-out trials of the same images (trial splits used so that the same images were shown in classifier training and testing). However, the particular class (face vs non-face) could be determined at above chance levels when reading the clT and aIT population responses.

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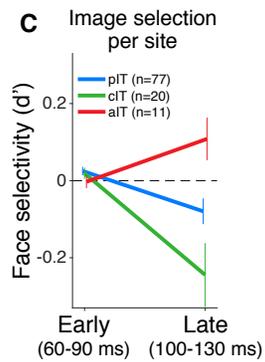
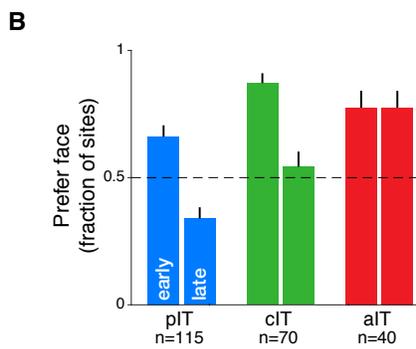
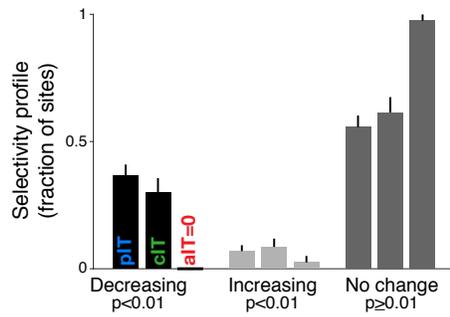
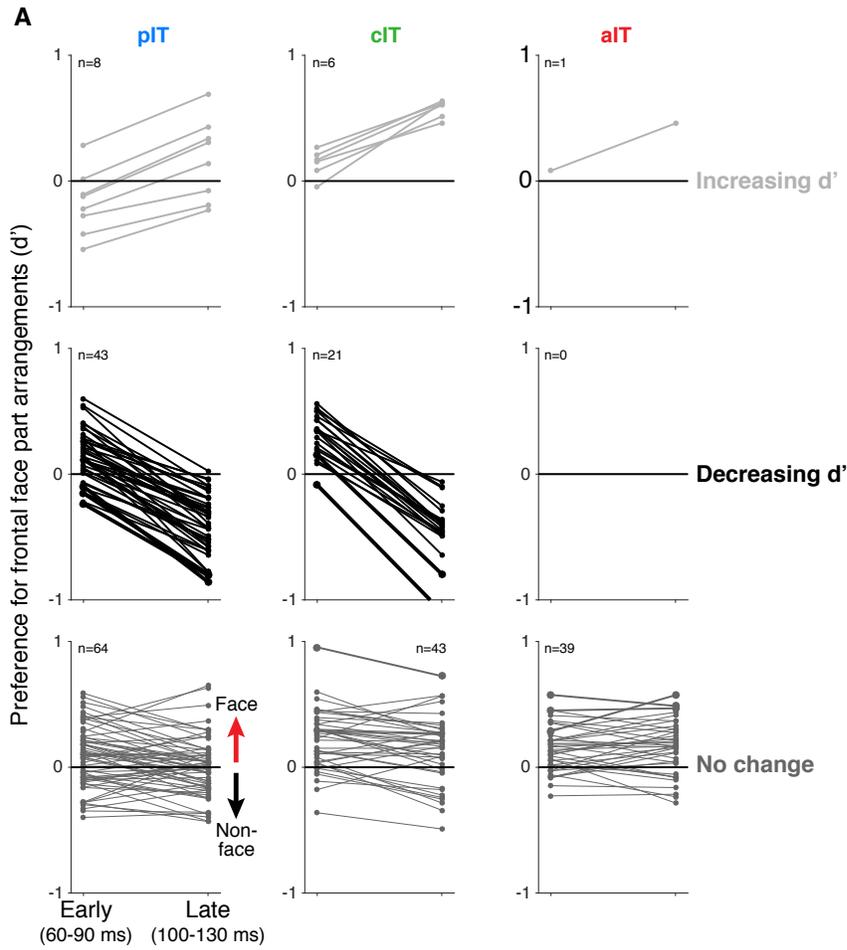
Figure 2



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Figure 2. Responses in example sites to face-like images with typical and atypical face part arrangements. The three sites with the highest selectivity in the late response phase in each region are shown (pIT, cIT, and aIT; left, middle, and right columns, respectively) (d' selectivity measured in a 100-130 ms window, gray shaded region shown in bottom, left panel). While the three aIT sites (right column) demonstrated late phase selectivity for face images, the three pIT sites evolved the opposite preference in their late phase (100-130 ms) responses (red line = mean response of 8 images shown in **Figure 1B** red box, and black line = mean response of 13 images shown in **Figure 1B** black box).

Figure 3



1199 **Figure 3. Time course of neural response preferences in pIT, cIT, and aIT for images with face**
1200 **versus non-face arrangements of the parts. (A)** Preferences for frontal face versus non-face part
1201 arrangements for each site are plotted in both early (60-90 ms post image onset) and late (100-130 ms)
1202 time windows. Sites are grouped based on region (pIT, cIT, aIT) and whether they showed a significant
1203 change in selectivity from early to late time windows (light gray = increased preference, black =
1204 decreased preference, and dark gray = no change in preference for face versus non-face images,
1205 significance tested at $p < 0.01$ level; example sites from **Figure 2** are plotted using thicker, darker
1206 lines). Many sites in pIT and cIT showed a decreasing preference for frontal face versus non-face
1207 images over time (black lines, middle row, left and center panels). In contrast, no sites in aIT had this
1208 dynamic (middle row, right panel). **(B)** The fraction of sites whose responses showed a preference for
1209 images of typical, face-like arrangements of the face parts in pIT (blue), cIT (green) and aIT (red) in the
1210 early (60-90 ms) and late (100-130 ms) phase of the response. Note that, in the late phase of the
1211 response, most pIT neurons paradoxically showed a preference for non-face arrangements of face
1212 parts. **(C)** Selectivity measured for images driving similar responses within a site. This procedure
1213 ensured matched initial responses on a site-by-site basis rather than using a fixed set of images based
1214 on the overall population response (i.e. the fixed image set of **Figure 1B**; here, the initial d' for 60-90
1215 ms is close to zero when images are selected site by site). Although initial response differences were
1216 near zero when using site based image selection, a late phase preference for non-face images still
1217 emerged in pIT and cIT but not in aIT similar to the decreasing selectivity profile observed when using a
1218 fixed image set for all sites.
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Figure 4

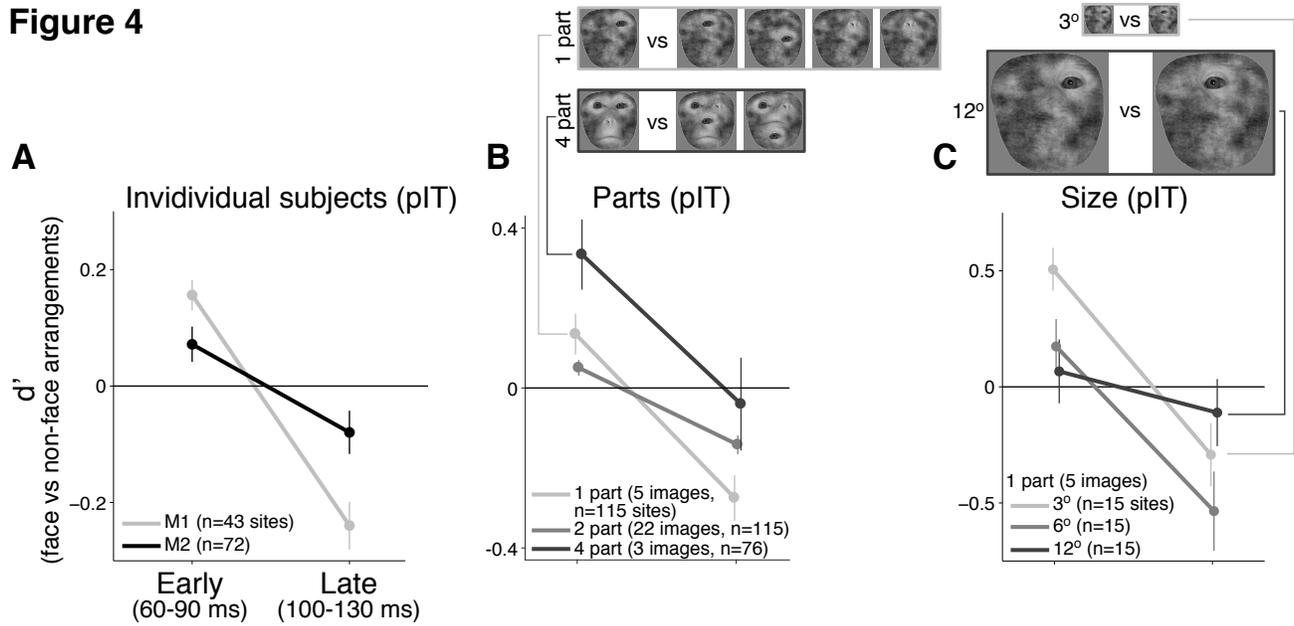


Figure 4. Individual monkey comparison and image controls for the decreasing selectivity profile in pIT. (A) Preference for images with frontal face part arrangements analyzed separately for each monkey. Median d' of pIT sites in both early and late time windows is shown. (B) Preference for images with face versus non-face arrangements of the parts 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 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 a decreasing preference over time for typically-arranged face parts leading to a preference for atypically arranged face parts in the later time window (100-130 ms).

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Figure 5

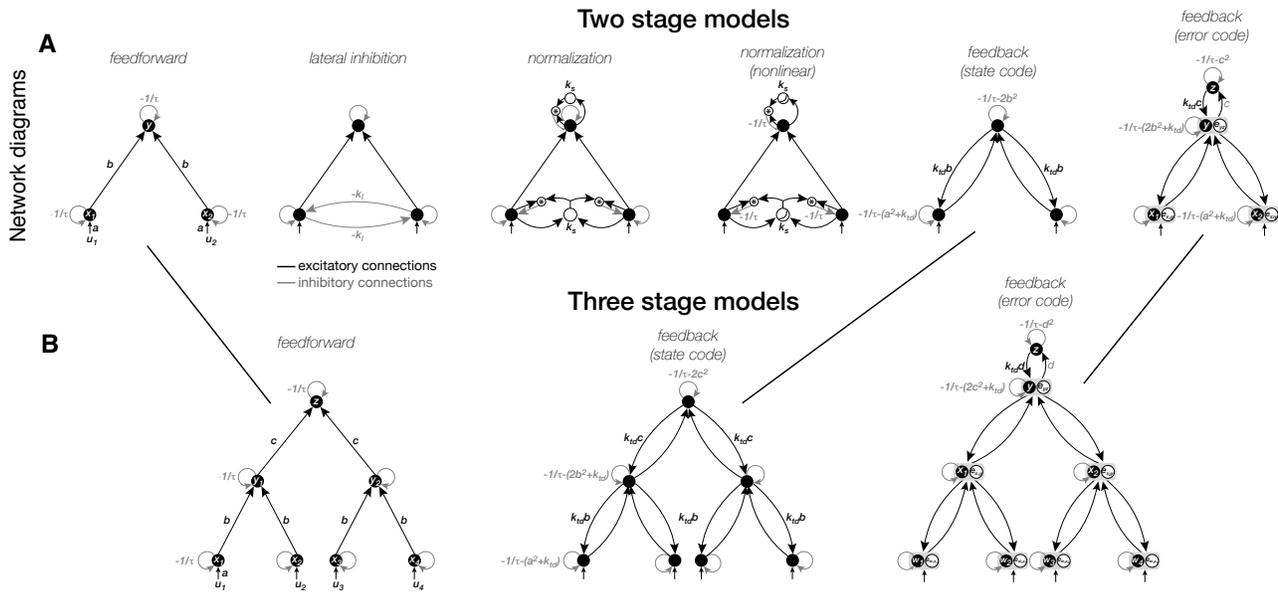
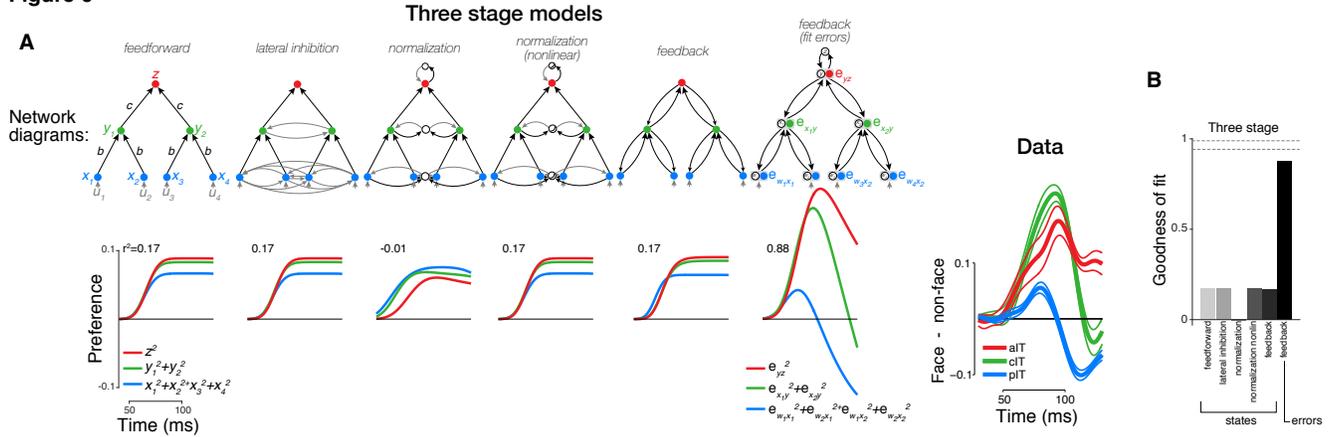


Figure 5. Model diagrams. (A) Network diagrams of two-stage models used in **Figure 7A,B**. The base feedforward architecture and parameters are shown in the first column while recurrent models with added connections and parameters are shown in the remaining columns. All models have two inputs (u_1, 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 (third and fourth columns). 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 (fifth and sixth columns). The error coding feedback model (sixth column) has an additional stage that contributes to computation of error in the second stage (see **Methods** for details). **(B)** Extensions of the two-stage model architectures to three stages are shown only for the feedforward and feedback models (compare to two-stage diagrams 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 of the error coding model.

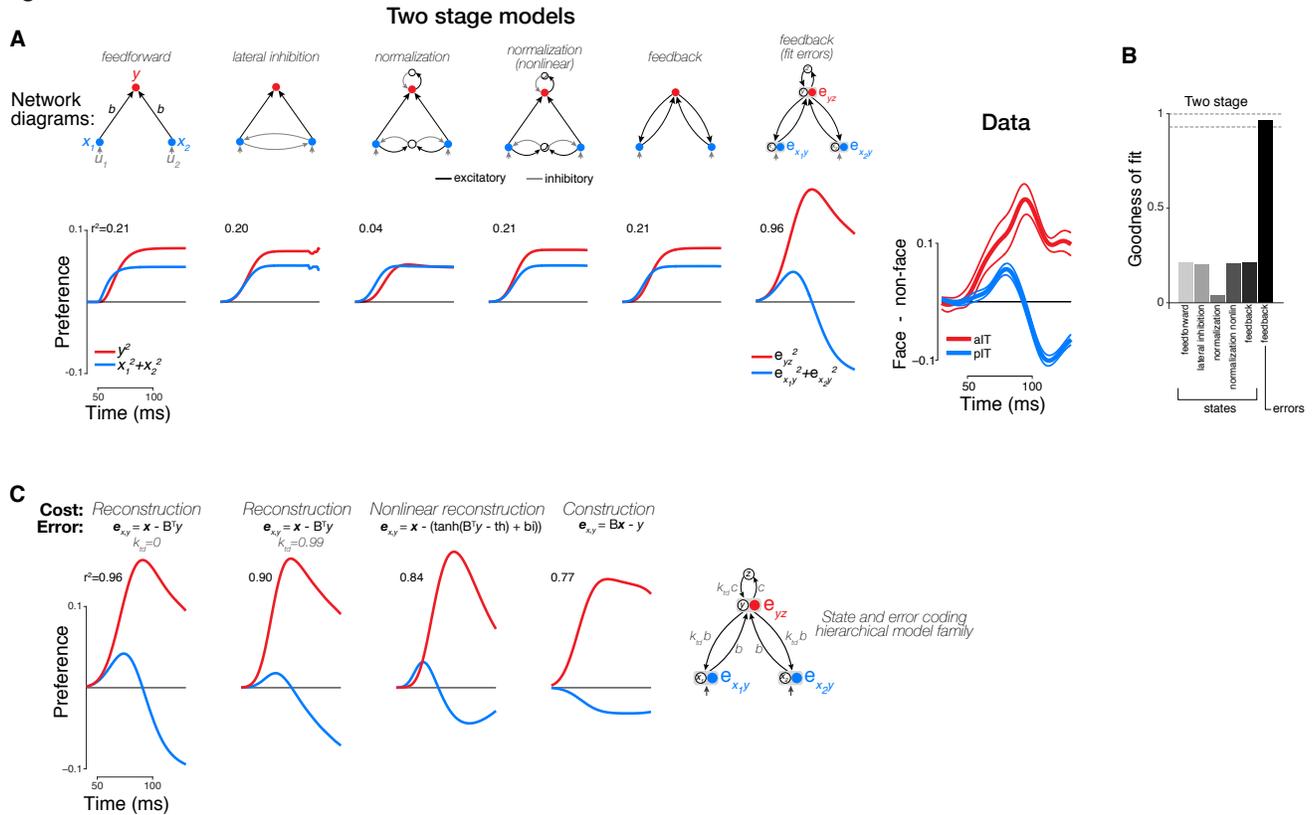
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Figure 6



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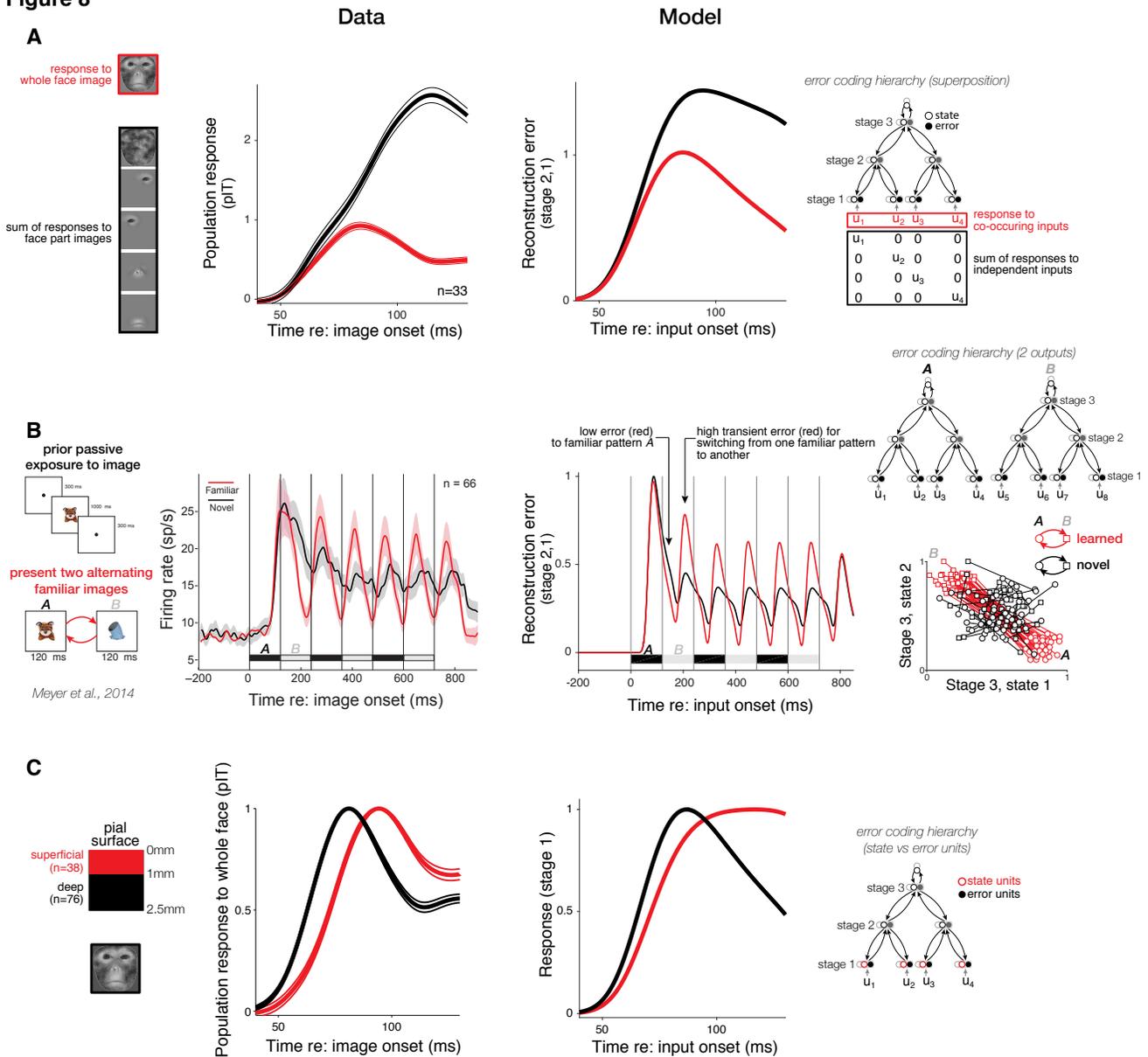
Figure 7



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1279 **Figure 7. Two stage model fits to neural dynamics and comparison of variants of error coding**
1280 **hierarchical models that use different algorithms for online inference. (A)** We tested how well two
1281 stage models could reproduce decreasing selectivity observed in the hidden layer (same format as
1282 **Figure 6A**). The two stage error coding feedback model displayed similar dynamical phenomena as the
1283 data suggesting that three stages of processing were not necessary to obtain error signal dynamics. **(B)**
1284 Goodness of fit to population neural data (right plot in **(A)**) for all two stage models. **(C)** Existing forms
1285 of error computing networks can be distinguished by the type of online inference algorithm that they
1286 use. In one case, inference does not utilize top-down information between stages (classic error
1287 backpropagation; between-stage feedback connections shown are not used in these networks during
1288 runtime). On the other hand, between-stage feedback can be used such as in more general forms of
1289 error backpropagation and predictive coding. We approximated these two extremes by including a
1290 parameter (k_{td} , see **Methods**) controlling the relative weighting of bottom-up (feedforward) and top-
1291 down (feedback) evidence during online inference (first and second panels). We found that top-down
1292 inference between stages was not necessary to produce the appropriate error signal dynamics, and k_{td}
1293 was equal to zero in our best fitting two-layer and three-layers models (first panel is same model as
1294 two-layer error coding model in **(A)**) although models with $k_{td} \sim 1$ also performed well (second panel).
1295 Models can also differ in their goal (cost function) which directly impacts the error signals required
1296 (equations in top row). Under a nonlinear reconstruction goal (emulating the nonlinear nature of spiking
1297 output), the resulting error signals were still consistent with our data (third column). A simple sigmoidal
1298 nonlinearity, however, did lead to additional details present in our neural data such as a rapid return of
1299 stimulus preference to zero in the hidden layer. When we tested a discriminative, construction goal
1300 more consistent with a supervised learning setting (e.g. classic error backpropagation) where bottom-
1301 up responses simply have to match a downstream target signal in classification tasks, we found that the

1302 errors of construction did not match the data as well as reconstruction errors (compare fourth column to
1303 first three columns) although both types of error outperformed all state-based models and were overall
1304 very similar (compare to **(A)**).
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Figure 8



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1308 **Figure 8. Comparison of neuronal phenomena to predictions of a feedback-based error coding**
 1309 **model.** To generate model predictions, the architectural parameters of the model (i.e. connection
 1310 weights and time constants) were held fixed, and only the input patterns u were varied. **(A)** Neuronal
 1311 (left panel): in pIT, we found that the sum of the neuronal response to the face parts presented
 1312 individually (black) exceeded the response to the same parts presented simultaneously (i.e. a whole
 1313 face, red). Each line is the mean response of 33 pIT sites normalized by the peak response to the
 1314 whole face. Model (right panel): The magnitude of errors between stage 1 and stage 2 of the model
 1315 showed a similar degree of sublinear integration (responses were normalized by peak response to the
 1316 simultaneous input condition). **(B)** We extended the model to include two units in the third, output stage
 1317 that responded to two learned input patterns (see **Methods**) with increased separation of patterns A
 1318 and B in this high-level feature space (red markers in far right panel; 50 draws were made from

1319 distributions for *A* and *B* and were compared to pseudo-randomly drawn inputs, black markers). When
1320 alternating the two learned or familiar patterns *A* and *B*, activations in the top layer of the networks
1321 experienced greater changes than when two randomly selected patterns were alternated (compare
1322 distances traversed in high-level feature space by red lines versus black lines, far right panel). Because
1323 of this large change in high-level activations, transient errors were generated back through the network
1324 for learned patterns. Strong oscillations could be observed in the error signals between stage 1 and
1325 stage 2 of the model for alternated familiar inputs *A* and *B* (120 ms period) (red curve, middle right
1326 panel). In contrast, alternating novel inputs with similar amplitude but with patterns not matching the
1327 learned weights led to small amplitude oscillations in upstream error signals (black curve). Note that the
1328 average response strength of state signals to novel and familiar inputs was matched at all network
1329 levels by construction (mean-matched inputs). The differing response dynamics of error signals under
1330 familiar versus novel patterns are qualitatively consistent with the IT findings for novel versus familiar
1331 images (left, reproduced with permission from *Figs. 1 & 2b* of Meyer et al., 2014). **(C)** The average
1332 response to the whole face for pIT sites recorded 0 to 1 mm below the pial surface (left panel;
1333 superficial recordings, red curve) and for sites 1 to 2.5 mm beneath the pial surface (black curve). (right
1334 panel) Average response of state units (red) and error units (black) in stage 1 of the model. Note the
1335 lagged response in state units which is similar to the lagged response of units in superficial recordings
1336 (red curve, left panel).