

1 **Title: A novel analytical decoder of BOLD signals for dissociating latent**
2 **neurobehavioral processes**

3
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35 **Abstract:**

36 Brain, as a complex cognitive system, often processes multiple dimension
37 information synchronously and integrate them to adapt dynamic environments and
38 make effective decisions.¹⁻³ How to retrieve latent neurobehavioral processes from
39 complex human neurobiological signals is an important yet previously unresolved
40 challenge.^{4,5} For instance, the previous literature has proposed two fundamental yet
41 mutually confounded processes during the decision making and affective processing,
42 i.e. valence and arousal.^{6,7} Here, we develop a novel analytical approach, orthogonal-
43 Decoding multi-Cognitive Processes (DeCoP), with which we dissociate neural
44 responses in processing valence and arousal information during tests of motivational
45 and emotional function. During reward/punishment anticipation, we decode brain-wide
46 responses into spatially overlapping, yet functionally independent, evaluation and
47 readiness networks, i.e., motivational valence and arousal processing, which are
48 modulated differentially by the meso-limbic vs nigro-striatal dopamine systems.
49 Similarly, during emotional reactivity, we decompose amygdala response into
50 independent emotional valence and facial arousal processing features. We demonstrate
51 that DeCoP can resolve paradoxically unexpected brain ‘inactivation’, and be applied
52 more generally to decode multiple latent neurobehavioral processes. Furthermore, we
53 anticipate our approach to advance both the design and hypothesis testing of cognitive
54 experimental task paradigms.

55

56 **Main**

57 The brain frequently engages parallel processing involving different latent
58 behavioural processes mediated by functionally distinct, though spatially over-lapping,
59 neural networks. Previously, human functional neuroimaging studies have been
60 unsuccessful in unravelling these processes from basal compound physiological signals.
61 It is therefore challenging to build process-specific and mechanistic models of the brain
62 or develop robust biomarkers for dysfunctional processes in psychiatric disorders.
63 Recent overarching frameworks propose two different cognitive processes engaged in
64 parallel during rewarded or punished behaviour, namely evaluation (i.e. valence
65 processing, scaling signal values from reward to punishment) and response readiness
66 (subsuming arousal and attentional salience, contributing to response preparatory
67 processes) ^{3,6}. The evaluation process is essential for guiding upcoming action
68 selections ⁸, for which the brain has evolved dedicated regions/circuits for evaluating
69 the value of actions ⁹⁻¹². Complementary to evaluation, both reward and punishment, as
70 highly salient events, attract higher attention than neutral stimuli, also engaging greater
71 levels of motor preparation and emotional arousal ¹³⁻¹⁵, hence contributing to response
72 (including motor) readiness. Therefore, evaluation and readiness signals are inevitably
73 confounded with each other during reward/punishment processing. Unfortunately,
74 decomposing this compound signal, for example in human fMRI studies, has proven
75 challenging because independent components cannot be identified in many
76 experimental test paradigms, although previous attempts have made to overcome this
77 problem ^{4,5,16}. However, existing approaches have failed to disentangle signals in brain
78 regions known to encode both evaluation and readiness signals, for example, in the
79 striatum and the ventromedial prefrontal cortex (vmPFC) ¹⁵, and have not provided
80 convincing evidence to either prove or refute the assumption of independence of the
81 signals. To resolve this complex theoretical issue, we developed a novel approach,
82 orthogonal-Decoding multi-Cognitive Processes (DeCoP), which, for the first time,
83 allows for a brain-wide decomposition of process-specific neurobiological
84 representations of complex neurobehavioral processes. This technique has wide

85 application for decomposing compound neuroimaging signals, such as the BOLD
86 response.

87 **Experiment and model designs of DeCoP**

88 A monetary incentive delay (MID) task (Fig. 1a), one of the classical and widely
89 used fMRI paradigms for reward processing, was conducted in 1939 children aged 9-
90 10 from the ABCD study (Extended Data Table. 1)¹⁷ to assess reward/punishment
91 processing with gain/loss conditions for small or large amounts of money and a neutral
92 condition (i.e., -5.0 \$, -0.2 \$, 0, 0.2 \$ or 5.0 \$). Specifically, in the second-level analyses
93 of the BOLD signal, based on these amounts, we defined plausible orthogonal contrasts
94 [-2, -1, 0, 1, 2] and [2, 1, 0, 1, 2] that reflected putative independent hypothetical
95 processes of evaluation and readiness, respectively (i.e. a large reward and a large
96 punishment are assumed to be equally salient). Their complementary orthogonal
97 contrasts (i.e., [-1, 2, 0, -2, 1] and [1, -2, 2, -2, 1]) could explain information not
98 accounted for by the above hypothetical models (Fig. 1b). We were thus able to retrieve
99 the decomposed signal components of those confounded processes (Extended Data Fig.
100 1a, see Methods for more details).

101 **Decompose neural signals with DeCoP**

102 The vmPFC (Brodmann area [BA] 10-11; Peak MNI: [-9, 49, -9], $t_{1938} = 15.69$,
103 Cohen's $D = 0.36$, $p_{\text{uncorrected}} = 2.55\text{E-}52$) and ventral striatum (VS, Peak MNI: [7, 25, -
104 3], $t_{1938} = 14.21$, Cohen's $D = 0.32$, $p_{\text{uncorrected}} = 1.14\text{E-}43$) were the most prominent
105 regions identified in the evaluation model (Fig. 1c upper left&1d), thus being highly
106 sensitive for tracking the entire dimension from punishment to reward. These areas
107 coincide with the terminal regions of the dopamine neuron projections from the ventral
108 tegmental area (VTA), i.e. the meso-corticolimbic dopamine system^{8,18,19}. For the
109 readiness model, however, the signals were more widely dispersed across cortical and
110 subcortical areas, including motor-somatosensory, salience and attention networks, and
111 regions such as the dorsal striatum (DS, Peak MNI: [0, 10, 4], $t_{1938} = 32.80$, Cohen's D
112 $= 0.75$, $p_{\text{uncorrected}} = 4.56\text{E-}188$) and thalamus (THA, Peak MNI: [13, -6, 16], $t_{1938} =$
113 24.66 , Cohen's $D = 0.56$, $p_{\text{uncorrected}} = 5.23\text{E-}117$) (Fig. 1c upper right &1d), consistent

114 with their engagement in processing both reward and punishment²⁰.

115 **Neural circuits for decomposed signals**

116 We then investigated whether the neural representations of evaluation and
117 readiness signals were underpinned by different neural circuits, in particular those
118 modulated putatively by the midbrain dopaminergic projections originating from either
119 the substantia nigra pars compacta (SNc) or the VTA, which plays a central role in
120 reward prediction and approach^{18,19}. We found regions of the evaluation model with
121 higher functional connectivity (FC) to VTA than to SNc (*paired t*-test: $t_{183} = 14.84$,
122 Cohen's $D = 1.10$, $p < 10E-32$, Fig. 2a), and regions of the readiness model with higher
123 FC to the SNc than to the VTA (*paired t*-test: $t_{183} = 3.63$, Cohen's $D = 0.27$, $p = 0.0004$,
124 Fig. 2a) based on 7T high-resolution resting-state fMRI data from the Human
125 Connectome Project (HCP)²¹. Further, we extracted the *t*-maps of the difference
126 between the seed-based FC from VTA and SNc (i.e. 'VTA > SNc') (Extended Data Fig.
127 2a-c), which was exhibited high similarities, although in opposite directions, with the
128 *t*-maps of both evaluation ($r = 0.22$, $p_{adj} < 10E-20$) and readiness ($r = -0.12$, $p_{adj} < 10E-$
129 12, Fig. 2b) models. Thus, the separate VTA and SNc dopamine projections could be
130 the putative source of evaluation and readiness signals, respectively.

131 **Independence of decomposed neural signals**

132 We further demonstrated that the distinct underlying neural pathways of
133 decomposed cognitive processes were indeed independent, which could be directly
134 inferred from uncorrelated signal components in the proposed decoding procedure (see
135 Methods for detailed proof). Based on our simulation results, only if the compound
136 signals were indeed a combination of independent signals, the decomposed signals from
137 DeCoP could be uncorrelated ($|r_{\text{mean}}| < 0.001$, the 'Independent' model in Extended Data
138 Fig. 1b & Table 2). Otherwise, the decomposed signals were highly correlated and
139 hence inseparable (the 'One Signal' model: $r_{\text{mean}} = 0.54$; the 'Push and Pull' model:
140 $r_{\text{mean}} = -0.45$; Extended Data Fig. 1b & Table 2). Thus, DeCoP also serves as an
141 analytical tool for the statistical inference on the independence of decomposed signals.
142 In this study, we found that the signals attributed to evaluation and N-shape models

143 (dependent signals: $r_{\text{mean}} = -0.093$, $p_{\text{bootstrap}} < 0.0001$ based on 10000 bootstrap; 32.06%
144 voxels with $r < -0.1$) together described the sensitivity of evaluation from punishment
145 to reward. This was independent ($|r_{\text{mean}}| < 0.01$, $p_{\text{bootstrap}} > 0.3$; > 99% voxels with $r \in [-$
146 $0.1, 0.1]$) of those attributed to the readiness and W-shape models (dependent signals:
147 $r_{\text{mean}} = -0.159$, $p_{\text{bootstrap}} < 0.0001$; 93.29% voxels with $r < -0.1$) that together described
148 the differentiated engagement of readiness from the neutral condition to
149 reward/punishment conditions (Fig. 2a&c; see Supplementary Information for more
150 details). Hence, the observed unbalanced sensitivity towards reward and punishment
151 could be parsed into two independent and balanced signal components, i.e. a point-
152 symmetric evaluation process and a line-symmetric readiness process. Moreover, the
153 paradoxical ‘inactive’ vmPFC during the large-win vs neutral contrast could now be
154 understood as a product of a trade-off between two independent processes: activation
155 by reward stimuli (i.e. of the evaluation process) and deactivation as part of the default
156 mode network (i.e. of the readiness process).

157 **Complementary signal components from DeCoP**

158 On the other hand, the N-shape and W-shape models (Fig. 1c) account for the
159 deviation from (or equivalently, they adjust) the shapes of the proposed evaluation and
160 readiness models, particularly the response to small reward/punishment conditions (Fig.
161 3c). This modification of the additional complementary orthogonal vectors also
162 provided an effective measurement of distance between the latent pure signal and the
163 proposed model (see Methods and Supplementary Information for more details).
164 Therefore, converging evidence indicated that most brain regions, including VS and
165 vmPFC, distinguish reward from punishment signals with their relative scales. In
166 another word, the brain simplifies the scales of the different values when processing
167 reward/punishment information, hence processing highly abstract information only.
168 However, the bilateral anterior insula (aINS) and dorsal anterior cingulate cortex
169 (dACC), commonly referred to as the salience network, were only sensitive to the large
170 reward/punishment conditions, while the small reward/punishment and neutral
171 conditions were undifferentiated, hence most likely tracking the parametric nature of

172 the experimental design (i.e. [-5, -0.2, 0, 0.2, 5] Extended Data Fig. 3&4 and Extended
173 Data Table 3; see Supplementary Information for detailed analyses). These results were
174 consistent with the role of aINS and dACC in updating and maintaining subjective value
175 information.^{22,23}

176 **Signal components affect task performance**

177 We further implemented a weighted voxel co-activation network analysis
178 (WVCNA) on the ABCD MID task to capture the most informative brain-wide signal
179 clusters²⁴ (see Supplementary Information for details) and identified 55 and 194
180 clusters for the evaluation and readiness models, respectively (Extended Data Fig. 5-
181 6& Supplementary Table 1). Again, we observed brain-wide low correlations between
182 evaluation and readiness clusters ($r_{\text{mean}} = 0.01$, $p = 0.1621$, > 99% pairwise $r \in [-0.1, 0.1]$;
183 Fig. 2b), hence further supporting their neural independence. We then investigated the
184 effects of decomposed signals on task performance using canonical correlation analysis
185 (CCA), and found associations between variations in the neural signal and task
186 performance across three condition categories (i.e. reward, neutral and punishment) for
187 both evaluation and readiness: accuracy (evaluation: adjusted η^2 (adj- η^2) = 0.025, p_{perm}
188 < 0.001; readiness: adj- η^2 = 0.079, p_{perm} < 0.001) and reaction time (evaluation: adj- η^2
189 = 0.020, p_{perm} < 0.001; readiness: adj- η^2 = 0.075, p_{perm} < 0.001) (Supplementary Table
190 2&3). For the evaluation clusters, higher accuracy and faster reactions for both reward
191 and punishment conditions (but not the neutral condition) were mainly associated with
192 reduced sensitivity in left ventral striatum (VS), bilateral inferior temporal-occipital
193 junctions (iTOJ), and in both sub- and pre-genual anterior cingulate cortices (sgACC
194 and pgACC) (presented by the first components; $r < -0.04$, Supplementary Table 2&3,
195 Extended Data Fig. 7a&b). For the readiness clusters, the executive regulation related
196 to accuracy might function differently in the presence or absence of valence signals: in
197 reward/punishment trials (represented by the first component), higher accuracy was
198 associated with greater engagement of task-preparation regions, i.e. including the
199 somatosensory-motor cortices and the dorsal attention network ($r > 0.04$) but
200 suppression of the salience network, i.e. the dorsal aINS and the dorsal ACC (dACC)

201 ($r < -0.04$), whereas on neutral trials (represented by the second component) both
202 relationships were reversed, with higher accuracy now being associated with
203 suppressed task-preparation areas ($r < -0.04$) and greater engagement of the salience
204 network, i.e. the ventral aINS and dACC ($r > 0.04$, Supplementary Table 2&Extended
205 Data Fig. 7c). Nevertheless, faster reactions were uniformly associated with greater
206 engagement of task-preparation areas ($r > 0.04$) and lesser engagement bilaterally of
207 the anterior putamen (aPUT), left VS, dACC and the right frontoparietal network (FPN,
208 i.e. dorsal lateral prefrontal cortex (DLPFC) and intra-parietal sulcus (IPS)) ($r < -0.04$)
209 across all three conditions (represented by the first component, Supplementary Table
210 3&Extended Data Fig. 7d; see Supplementary Information for more information).

211 **Decompose emotion processes with DeCoP**

212 To further demonstrate the broader applicability of this unified strategy for signal
213 decomposition, we decoded another complex cognitive process, emotional face
214 processing. Similarly to reward/punishment processing, emotional processing also
215 involves compound signals of distinct behavioural processes such as affective valence,
216 emotional arousal, and attention, especially in the amygdala²⁵. To address this issue
217 using the DeCoP method we analysed fMRI data for a modified emotional faces task
218 performed in 1091 19-year-old adolescents from the IMAGEN project²⁶. While the
219 right amygdala was activated by all types of facial stimuli in the emotional faces task
220 (Fig. 4a, $t_{1090} > 13$, Cohen's $D > 0.39$), there was no significant difference in activations
221 between the angry and neutral conditions (Fig. 4b upper right), consistent with previous
222 studies^{27,28}. Given the multidimensional role of the amygdala in several of the
223 underlying processes it is conceivable that the emotional-decoding process-specific
224 neural responses have been obscured by other processes, such as general processing of
225 facial features. Again, we hypothesized that two dissociable components are involved
226 in emotional processing, i.e. affective valence, corresponding to the neural
227 representation of the affective facial emotion information (represent as [1, 0, -1] from
228 angry to happy, Fig. 4c) and facial arousal, which was defined as the degree of
229 excitement or motivational reaction when a person experienced and recognized an

230 emotional face (represent as [1, 0, 1], Fig. 4c). By applying DeCoP, we found an
231 independent positive reaction to affective valence and negative reaction to facial arousal
232 (Fig. 4c, $r_{\text{mean}} = -0.041$, $p_{\text{bootstrap}} = 0.1083$; 97.78% voxels with $|r| < 0.1$). Hence, the
233 right amygdala linearly tracked emotional valence on the happy through neutral to
234 angry dimension, while the neutral faces attracted more attention (i.e. higher activation)
235 than emotional ones. Therefore, both processes lead to an undifferentiated contrast for
236 angry vs neutral faces.

237 **Conclusion**

238 In conclusion, we have developed and evaluated a universally applicable, novel
239 signal decomposition strategy, 'DeCoP', to disentangle behavioural processes that
240 confound the observation of functional neuroimaging signals. Through DeCoP we
241 demonstrated the independence of evaluation and readiness processing in the brain,
242 putatively modulated differentially by neural circuits targeting VTA and SNc; we also
243 demonstrated that most brain regions, including the ventral striatum, coded signals
244 based on abstract information instead of the observed exact values, except for the
245 salience network, i.e. pgACC/dACC and aINS. Most importantly, we demonstrated that
246 DeCoP could help to resolve common paradoxical observations in fMRI tasks which
247 involve multiple interferential latent behavioural or cognitive processes, for example,
248 the unexpectedly 'inactive' vmPFC in the contrast of large reward vs no reward, and
249 the lack of differential amygdala activation in the contrast of angry vs neutral faces. We
250 expect DeCoP can be applied usefully in many other comparably ambiguous data-sets.

251

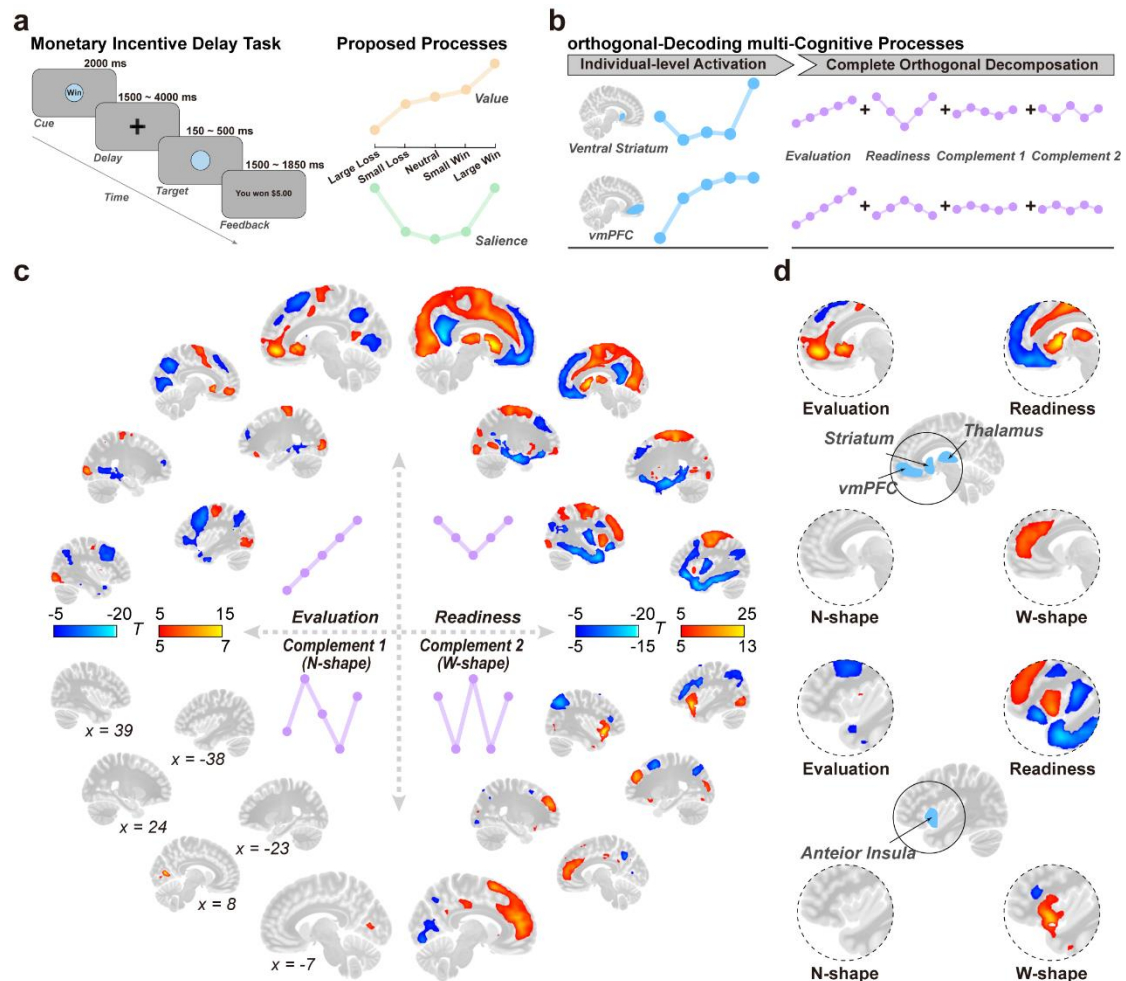
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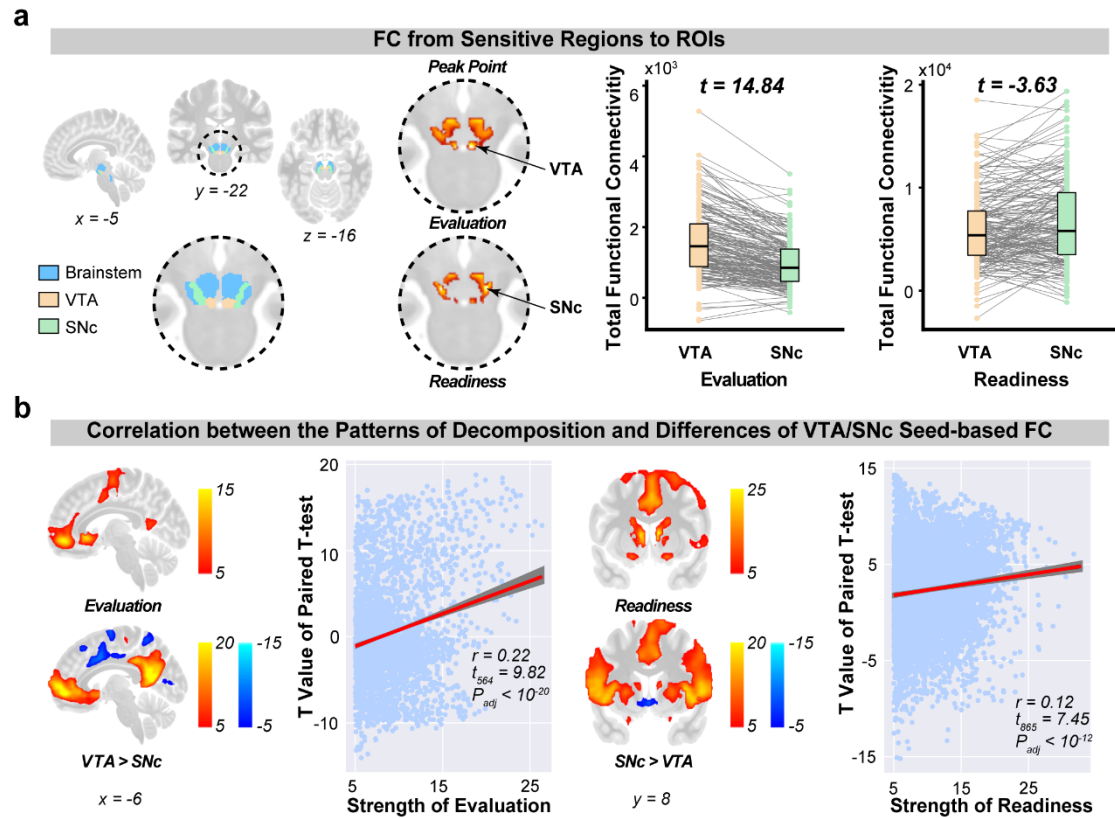
336 **Figures**



337

338 **Fig. 1. Experimental design and neural representations of the orthogonal**
 339 **decomposition.** **a.** Procedure of the monetary incentive delay task (MID) and the
 340 proposed evaluation and readiness cognitive processes; **b.** An illustration of
 341 Orthogonally Decoding multi-Cognitive Processes (DeCoP); **c.** Brain-wide T-maps of
 342 decomposed signals for orthogonal contrasts. Brain-wide significance was set as $|T| > 5$.
 343 The MNI coordinates of brain slices were inserted at the lower left; **d.** Decomposed
 344 signals in highlighted brain regions.

345



346

347 **Fig. 2 The decomposed evaluation and readiness processing targets VTA and SNc**

348 **neural circuits respectively. a.** Left: the masks of Brainstem, VTA and SNc were from

349 AAL3 atlas. Middle: strength of functional connectivities (FCs) to VTA and SNc from

350 the evaluation and readiness regions identified in Fig. 1. Right: Paired t-tests between

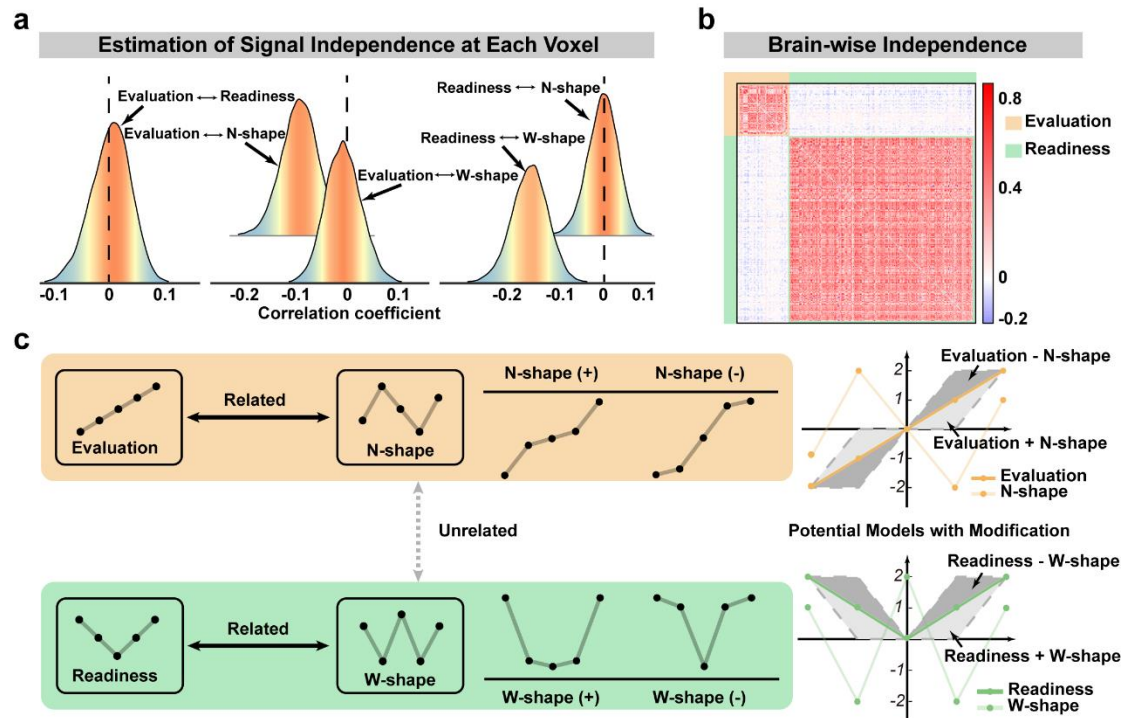
351 FCs to VTA and SNc from evaluation and readiness; **b.** Brain-wide pattern correlations

352 between the strength of decomposition signals (left: evaluation; right: readiness) and

353 the differences of seed-based FCs from VTA and SNc. Brain-wide significance was set

354 as $|T| > 5$.

355



356

357

358 **Fig. 3 Inference for independence among orthogonally decomposed signals during**

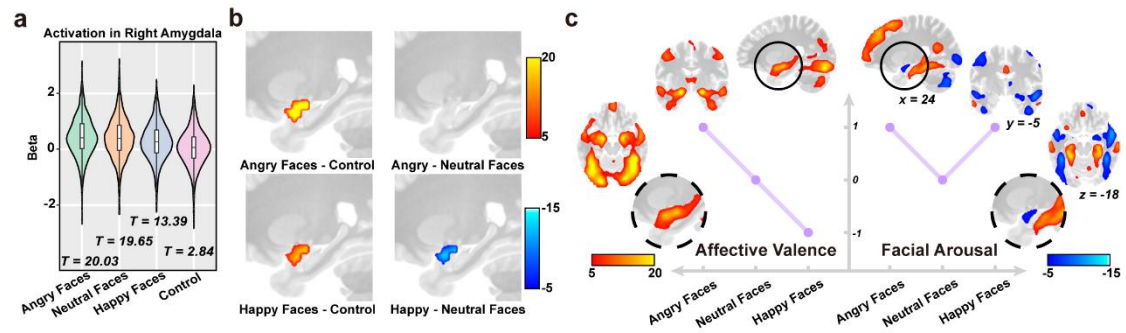
359 **the reward anticipation. a.** The distributions of pair-wise correlations between signals

360 **of orthogonal contrasts at each voxel. Mean correlations deviated from 0 would indicate**

361 **a pair of related signals; b.** The correlation matrix of signals from the evaluation and

362 **readiness clusters identified by WVCNA. c.** An illustration of how related signals could

363 **describe the evaluation-related and readiness-related processing.**



364

365

Fig. 4 Decoding affective valence and facial arousal of emotional face processing

366

in amygdala by DeCoP. a. BOLD signals of the right amygdala under different task

367

conditions. **b.** Activations of the right amygdala under different contrasts. **c.** The neural

368

representations of affective valence and facial arousal based on orthogonal

369

decompositions. Brain-wide significance was set as $|T| > 5$.

370

371 **Methods**

372 **Participants**

373 The dataset used for this study was selected from Annual Curated Data Release
374 2.01 (<https://data-archive.nimh.nih.gov/abcd>) of the Adolescent Brain Cognitive
375 Development (ABCD) cohort, which recruited 11,875 children between 9–11 years of
376 age from 21 sites across the United States¹⁷. The study conforms to each site's
377 Institutional Review Board's rules and procedures, and all participants provide
378 informed consent (parents) or informed assent (children). More details of the subjects
379 and the data collection are provided at the ABCD website
380 (<https://abcdstudy.org/scientists/protocols>) and are also described previously¹⁷.

381 Magnetic resonance imaging (MRI) data in the ABCD study were collected from
382 different 3T scanner platforms (i.e. Siemens Prisma, General Electric (GE) MR750 and
383 Philips Achieva dStream). To minimize biases introduced by multiple platforms, we
384 only included MRI data from the most frequent manufacturer Siemens Prisma, i.e. 5968
385 participants from 13 sites. By examining the similarity of brain activations across these
386 13 sites, we selected 2326 participants from 4 sites with consistent activation patterns.
387 Furthermore, the data with poor registration (by visual check) and high head motion
388 (mean framework displacement (FD) > 0.5 mm) were excluded. Hence, 1939 quality-
389 controlled participants were included in the following analysis and the demographic
390 characteristics of these participants are summarized in Extended Data Table 1.

391 **Monetary incentive delay (MID) task design**

392 A modified version of the MID task was used to examine brain activation during
393 monetary reward anticipation and receipt²⁹, which consists of five levels of incentive:
394 large loss, small loss, neutral, small win and large win (i.e., -5.0 \$, -0.2 \$, 0\$, 0.2 \$ and
395 5.0\$ respectively). In each trial, participants were first presented with one of three cue
396 shapes (circle, square or triangle) that indicated the trial condition (of win, loss, or
397 neutral, respectively), as well as the amount of money involved. This cue presentation
398 (2,000 ms) was followed by a jittered anticipatory delay (1,500-4,000 ms) of fixation
399 on a black crosshair. Subsequently, a blank target cue (with the same shape as the

400 previously presented cue) emerged and required the participants to press a response
401 button before the target disappeared to win or avoid loss. With a tracking algorithm, the
402 time of the target on the screen was dynamically manipulated (i.e., 150 ms-500 ms) to
403 maintain a 60% success rate for each participant. After a short delay, feedback of the
404 current trial (i.e., the amount of monetary gain or loss) and the accumulated reward so
405 far were presented for 1,500-1,850 ms (Fig. 1A). During the anticipation phase,
406 participants underwent 50 trials in total (i.e., 10 trials per incentive degree). Participants
407 had first completed a practice session outside the scanner before completing two
408 sessions of the MID task with fMRI recording (approximately 5.5 minutes each).

409 **Image acquisition and preprocessing**

410 Imaging protocols were harmonized across sites and scanners. 3-dimensional T1-
411 weighted images (1.0 mm isotropic, TR = 2500 ms, TE = 2.88 ms) were acquired with
412 a gradient-echo sequence for anatomical localization and high spatial (2.4 mm isotropic)
413 and temporal (TR = 800 ms) resolution MID-task-based fMRI was acquired with echo-
414 planar imaging (EPI) sequence in two separate runs (approximately 5.5 minutes each).
415 The detailed MRI acquisition protocol is described elsewhere¹⁷. All functional images
416 were preprocessed with the same preprocessing procedure by suggested protocols from
417 FMRIB's Software Library (FSL v5.0.9), Advanced Normalization Tools (ANTs v1.9.2)
418 and Analysis of Functional NeuroImages (AFNI v18.3.03). Concretely, the whole
419 preprocessing procedure included the following steps: (i) brain extraction (ANTs
420 antsBrainExtraction), nonlinear registration to MNI space (ANTs antsRegistrationSyN)
421 for structural images; (ii) rigid realignment to adjust for motion (FSL MCFLIRT) and
422 field map correction (FSL TOPUP) for functional images; (iii) co-registration to a high-
423 resolution T1 image and normalization to 3 mm isotropic MNI standard space (ANTs
424 antsRegistrationSyN) and (iv) spatial smoothing with a 6mm full-width at half-
425 maximum (FWHM) Gaussian kernel (AFNI 3dBlurToFWHM) and detrending (AFNI
426 3dDetrend).

427 **First-level analysis of task-based fMRI**

428 At the first-level analysis, we set up a general linear model to estimate the effects
429 of the task conditions at the individual level with SPM, which contained 20 task
430 condition regressors (i.e. Target (Hit or Miss) * Phases (Anticipation, Feedback) * Task
431 Conditions (Large-Loss, Small-Loss, Neutral, Small-Win or Large-Win)) and
432 additional covariate regressors (i.e., 24 motion-related parameters: 6 rigid-body motion
433 parameters, their first temporal derivatives and 12 corresponding squared items; and
434 mean signals of both white matter and ventricles). All regressors were convolved with
435 a double-gamma HRF, and no orthogonalization of regressors was applied. For BOLD
436 signals, drift was modeled with the DCT basis, and a cut-off of 128 s was applied (SPM
437 defaults). Finally, the autocorrelation was modeled as a global approximate AR(1) in
438 SPM.

439 **Orthogonally Decoding multi-Cognitive Processes**

440 In the present study, we propose a novel approach to decompose each participant's
441 brain activations at varied conditions (denoted as y) with a set of orthogonal basis
442 $\mathbf{x} = (x_1, \dots, x_k)$, where each vector could represent a predefined signal model, e.g.
443 evaluation or readiness. Specifically, 'orthogonal' here means that the pairwise
444 covariances of vectors were all zero, i.e. $Cov(x_i, x_j) = E(x_i, x_j) - E(x_i)E(x_j) = 0$.
445 In this way, the regression coefficients $\boldsymbol{\beta} = (\beta_1, \dots, \beta_k)$ (i.e. the strength of signals for
446 each individual) estimated from a multiple linear model with all vectors were the same
447 as those estimated univariately (of simple linear models), i.e.
448 $T(\beta_i | y, \mathbf{x}, \boldsymbol{\beta}_{-i}) = T(\beta_i | y, x_i)$, where $T(\cdot)$ stands for the best linear unbiased
449 estimator.

450 We propose that the above individual level orthogonal disconfounding eliminates
451 spurious correlations of signal components (i.e. β) introduced by related vectors (i.e. X_i
452 are correlated), thus allowing us to make meaningful inferences regarding signal
453 independence at the population level. Here, we only describe a simple proof for the
454 purpose of illustration, where only two orthogonal vectors were involved. Let y_i denote
455 the activations of the i th individual across all conditions, and x_{1i} and x_{2i} denote the two

456 predefined independent orthogonal models of input signals, where $Cov(x_{1i}, x_{2i}) = 0$,
 457 and β_{1i} and β_{2i} denotes the corresponding regression coefficients, i.e. the signal
 458 strengths. We have the following linear model:

$$459 \quad y_i = \beta_{1i}x_{1i} + \beta_{2i}x_{2i} + \varepsilon. \quad (1)$$

460 We first show that orthogonal settings of x_{1i} and x_{2i} are necessary for a meaningful
 461 signal decomposition. The covariance and variance conditional on x_{2i} are easily derived.

$$462 \quad Cov(x_{1i}, y_i | x_{2i}) = Cov(x_{1i}, y_i) - \frac{Cov(x_{1i}, x_{2i}) \cdot Cov(y_i, x_{2i})}{Var(x_{2i})}$$

463 and

$$464 \quad Var(x_{1i} | x_{2i}) = Var(x_{1i}) - \frac{Cov(x_{1i}, x_{2i})^2}{Var(x_{2i})}.$$

465 Similar results are easily acquired for the covariance/variance conditional on x_{1i} .

466 Therefore, we establish the following relationship between the least square estimations
 467 (LSEs) of β_{1i} and β_{2i} :

$$\begin{aligned} & \frac{Cov(x_{1i}, y_i)}{Var(x_{1i})} - \beta_{1i} \\ &= \frac{Cov(x_{1i}, y_i)}{Var(x_{1i})} - \frac{Cov(x_{1i}, y_i | x_{2i})}{Var(x_{1i} | x_{2i})} \\ &= \frac{Cov(x_{1i}, y_i)}{Var(x_{1i})} - \frac{Cov(x_{1i}, y_i) - \frac{Cov(x_{1i}, x_{2i}) \cdot Cov(x_{2i}, y_i)}{Var(x_{2i})}}{Var(x_{1i}) - \frac{Cov(x_{1i}, x_{2i})^2}{Var(x_{2i})}} \\ &= \frac{Cov(x_{1i}, x_{2i}) \cdot Cov(x_{2i}, y_i) \cdot Var(x_{1i}) - Cov(x_{1i}, y_i) \cdot Cov(x_{1i}, x_{2i})^2}{Var(x_{1i})^2 \cdot Var(x_{2i}) - Var(x_{1i}) \cdot Cov(x_{1i}, x_{2i})^2} \\ &= \frac{Cov(x_{1i}, x_{2i})}{Var(x_{1i})} \cdot \frac{Cov(x_{2i}, y_i) - \frac{Cov(x_{1i}, x_{2i}) \cdot Cov(x_{1i}, y_i)}{Var(x_{1i})}}{Var(x_{2i}) - \frac{Cov(x_{1i}, x_{2i})^2}{Var(x_{1i})}} \\ &= \frac{Cov(x_{1i}, x_{2i})}{Var(x_{1i})} \cdot \frac{Cov(x_{2i}, y_i | x_{1i})}{Var(x_{2i} | x_{1i})} \\ &= \frac{Cov(x_{1i}, x_{2i})}{Var(x_{1i})} \cdot \beta_{2i} \end{aligned}$$

468

469 Clearly, unless $Cov(x_{1i}, x_{2i}) = 0$, β_{1i} could always be expressed as a function of β_{2i} , and
 470 hence can never be independent of each other. The above derivations, therefore, prove
 471 the necessity of using orthogonal vectors for meaningful signal decomposition.

472 Meanwhile, by setting $Cov(x_{1i}, x_{2i}) = 0$, and realizing that the LSE of β_{1i} in the
 473 absence of x_{2i} from the regression model (1) (i.e. reduced to a simple linear model) is
 474 $\beta'_{1i} = Cov(x_{1i}, y_i) / Var(x_{1i})$, we could immediately have $\beta'_{1i} = \beta_{1i}$. Thus with
 475 orthogonal x_{1i} and x_{2i} , estimations from a multiple linear model would be the same as
 476 those univariately estimated from simple linear models.

477 We then show that the population-level correlation analyses of β_{1i} and β_{2i} derived
 478 above provide meaningful statistical inferences of signal independence. For simplicity,
 479 assume that we rewrite β_{2i} , the signal strength of model x_{2i} , into a sum of two
 480 independent components, i.e., $\beta_{2i} = \beta'_{2i} + \gamma_i \beta_{1i}$ where β'_{2i} is independent of β_{1i} , i.e.,
 481 $Cov(\beta'_{2i}, \beta_{1i}) = 0$, and the parameter γ_i denotes the proportion of overlapped signals
 482 with model x_{1i} , and is independent of β_{1i} , i.e., $Cov(\gamma, \beta_1) = 0$. We then calculate the
 483 population-level correlation of β_{1i} and β_{2i} as:

$$\begin{aligned}
 Cov(\beta_1, \beta_2) &= Cov(\beta_1, \beta'_2 + \gamma\beta_1) \\
 &= Cov(\beta_1, \beta'_2) + Cov(\beta_1, \gamma\beta_1) \\
 &= E(\gamma\beta_1^2) - E(\beta_1)E(\gamma\beta_1) \\
 484 &= E(\gamma\beta_1^2) - E(\gamma)E(\beta_1^2) + (E(\gamma)E(\beta_1^2) - E(\beta_1)E(\gamma)E(\beta_1)) \\
 &= Cov(\gamma, \beta_1^2) + E(\gamma)Var(\beta_1) \\
 &= E(\gamma)Var(\beta_1)
 \end{aligned}$$

485 which only equals 0 if either $E(\gamma) = 0$, i.e. the population mean signals of both models
 486 are not a function of each other, or $Var(\beta_1) = 0$, i.e. one of the signals is invariant
 487 across the population, and hence again is not dependent on each other. Either way, the
 488 signals of both models are indeed independent, and thus $Cov(\beta_1, \beta_2) = 0 \Leftrightarrow \beta_1 \perp \beta_2$,
 489 where orthogonal decomposition at the individual level permits the expression

490 $Cov(\beta_1, \beta_2) = 0$. In conclusion, this two-stage approach can be used for statistical
491 inference concerning the signal dependence or independence of two signals.

492 To illustrate the above theoretical derivations, we conducted simulation analyses to
493 evaluate three models, namely the “independent-signal model”, where each observed
494 signal consists of two independent inputs ($\beta_{1i} \perp \beta_{2i}$, and hence $Cov(\beta_1, \beta_2) = 0$), and
495 two dependent-signal models, i.e. the ‘single signal model’ with $\beta_{2i} = \gamma_i \beta_{1i}$ (assume
496 $E(\gamma) > 0$, and hence $Cov(\beta_1, \beta_2) > 0$), and the ‘push-and-pull model’ with $\beta_{2i} +$
497 $\gamma_i \beta_{1i} = c_i$ (assume $E(\gamma) > 0$, and hence $Cov(\beta_1, \beta_2) < 0$). For simplicity, the signal
498 overlapping parameter γ_i was set as a constant (i.e. 0 for the independent-signal model
499 and 1 in both dependent-signal models); the observed signals (y in equation 1) were
500 simulated from $[-2, -1, 0, 1, 2]$ to $[2, 1, 0, 1, 2]$ with an increment of $[0.2, 0.1, 0, 0, 0]$
501 at each step, i.e. $[-2, -1, 0, 1, 2], [-1.8, -0.9, 0, 1, 2], \dots, [1.8, 0.9, 0, 1, 2], [2, 1, 0, 1, 2]$;
502 and the orthogonal vectors were fixed as $x_{1i} = [2, 1, 0, 1, 2]$ and $x_{2i} = [-2, -1, 0, 1,$
503 $2]$. Therefore, for the ‘single signal model’, y_i was directly set as the observed signals
504 plus a random noise $N(0,1)$. For the ‘push-and-pull model’, at the i th simulation step,
505 by setting $c_i = 0.05 \times (i-1) + N(0,1)$, $\beta_{1i} = c_i + N(0,1)$ and $\beta_{2i} = 1 - c_i + N(0,1)$, we
506 could then simulate $y_i = \beta_{1i} x_{1i} + \beta_{2i} x_{2i}$. Clearly, at the population-level, we have
507 $Cov(\beta_1, \beta_2) = E_i^2(c) - E_i(c^2) = -Var(c)$, and hence β_1 and β_2 are dependent. For the
508 ‘independent-signal model’, at the i th simulation, by setting $c = 0.05 \times (i-1)$,
509 $\beta_{1i} = c + N(0,1)$ and $\beta_{2i} = 1 - c + N(0,1)$, we could also simulate $y_i = \beta_{1i} x_{1i} + \beta_{2i} x_{2i}$,
510 where, however, one would expect $Cov(\beta_1, \beta_2) = 0$. It is notable that while the
511 simulation models of ‘push-and-pull’ and ‘independent-signal’ are rather similar, they
512 are fundamentally different. The reason lies in the fact that the constant c in the
513 ‘independent-signal model’ is invariant across individuals and solely determined by the
514 predefined form of observed signals, and hence the signal strengths β_{1i} and β_{2i} do
515 not dependent on each other. However, in the ‘push-and-pull model’, the form of

516 observed signals only determined the expectation $E_i(c)$, and hence both β_{1i} and β_{2i}
517 are nevertheless the functions of c_i that varies across individuals, and thus are not
518 independent.

519 For each model, we simulated with 1000 independent individuals for 1000 times,
520 and the detailed results of the simulation are shown in Extended Data Fig. 1b and
521 Extended Data Table 2.

522 **Functional connectivity (FC) based on resting-state fMRI data**

523 A total of 184 participants' preprocessed high-resolution (7T) resting-state fMRI
524 data were collected from the Human Connectome Project (HCP) dataset ²¹. Total FC
525 (Pearson Correlation) from evaluation and readiness activation activations regions to
526 the voxels in brainstem (Fig. 4A), which were masked with AAL3 atlas ³⁰ and from
527 VTA and SNc to the whole brain were examined respectively. A paired t-test was used
528 to estimate which brain region had a stronger FC with the activation region. The
529 similarity between patterns of activation and FC was represented by Pearson's
530 correlation coefficient r . Considering the high correlation between voxels, the degree
531 of freedom was adjusted according to the number of components which had over 95%
532 interpretation of brain activation by principal component analysis.

533 **Measurement of modifying effects**

534 As N-shape and W-shape models can be considered as modifiers of evaluation and
535 readiness models, respectively, their strength (i.e. the activation level) can be used to
536 measure the deviance from the proposed model settings, i.e. the exact scale or the
537 relative scale. Thus we computed the standardized mean
538 $((t_{relative} + t_{exact})/\max(\text{abs}(t_{relative}), \text{abs}(t_{exact})))$ to evaluate the prevailing
539 settings, where $t_{relative}$ represents the t-statistics of the activation for N-shape or W-
540 shape models under the relative-scale setting, and t_{exact} represents the t-statistics of
541 the activation for N-shape or W-shape models under the relative-scale setting. When
542 the observed signal falls somewhere between the relative-scale and exact-scale settings,
543 it is readily observable that $t_{relative}$ would be positive and t_{exact} would be negative.
544 Therefore, $\text{abs}(t_{relative}) > \text{abs}(t_{exact})$ would prefer the exact-scale setting, and the

545 standardized mean ranges from 0 to 1, whereas $abs(t_{relative}) < abs(t_{exact})$ would
546 prefer the relative-scale setting, and the standardized mean ranges from -1 to 0. It is
547 notable that meaningful settings other than this may also be possible. For instance, the
548 response of the small stimuli could be identical to the large stimuli in the observed
549 signal, thus $t_{exact} < t_{relative} < 0$, and the standardized mean could be well smaller
550 than -1.

551 **Weighted voxel co-activation network analysis (WVCNA)**

552 The R package WGCNA³¹ was implemented to conduct the WVCNA³², which
553 identifies activation modules of both evaluation and readiness models across the brain.
554 The final dataset used for WVCNA included 1,912 participants with 3365 voxels for
555 the evaluation model and 1900 participants with 10932 voxels for the readiness model
556 after removing null data and outliers. We transferred most parameters as default settings
557 from previous studies³³, except for the soft-threshold parameters, which were set to
558 seven based on the scale-free topology criteria (Fig. S5), incidentally identical to those
559 estimated for the MID task from a different cohort³³. The stabilities of the generated
560 modules were assessed through bootstrapping.

561 **Canonical correlation analysis (CCA)**

562 CCA has been widely used to investigate the overall correlation between two sets
563 of standardized variables³⁴. Due to high intra-correlations in both brain networks and
564 task performances, multicollinearity was a potential risk factor jeopardizing the validity
565 of the subsequent statistical inference. Therefore, we adopted the CCA proposed in a
566 previous reference³³. Briefly, we used the eta square (η^2) to represent the proportion of
567 mutually explained variance between the two sets of variables. For each correlation, the
568 *P* value or significance level was determined using the permutation test, where the
569 individual IDs of task performances were randomly shuffled at each iteration to
570 generate the null distribution of the corresponding test statistics. We further included
571 an adjusted η^2 to correct for the inflation in η^2 caused by the increased number of
572 variables as:

$$573 \quad \eta_{adj}^2 = 1 - \frac{1 - \eta^2}{1 - \eta_0^2}$$

574 where η_0^2 represents the expected η^2 under the null hypothesis that there is no
575 relationship between the two sets of variables (that is, it acts as a measure of inflation
576 in η^2), and can be directly estimated through the permutation test. As the effect of one
577 task performance was significant, we also estimated the effects of each component of
578 the behavior with the significant levels at $P < 0.05$.

579 **Emotional faces task in the IMAGEN project**

580 Emotional face processing was investigated using the “emotional faces task” in
581 the IMAGEN project ²⁶. Participants were exposed to a sequence of stimuli which
582 consisted of short (2–5 s) black-and-white video clips showing male and female faces
583 with varying facial expressions. Stimuli showed human faces which started with the
584 expression of a neutral expression and then either turned angry/happy or displayed a
585 neutral movement without a particular emotional content (for example, twitching the
586 nose). Stimuli were arranged in 18 s blocks, each block including 4–7 video clips
587 depicting faces of the same emotion or neutral. Altogether, there were 3 blocks of
588 neutral faces and 3 blocks containing angry and happy faces respectively. In between 2
589 blocks of face clips, an 18 s non-biological control video clip was presented. The control
590 stimuli consisted of expanding and contracting black-and-white concentric circles of
591 various contrasts, roughly matching the contrast and motion characteristics of the faces
592 clips.

593 The recruitment procedures employed in the IMAGEN project and demographic
594 information have been described previously ²⁶. The standard operating procedures for
595 the IMAGEN project are available at [http://www.imagen-](http://www.imagen-europe.com/en/Publications_and_SOP.php)
596 [europe.com/en/Publications_and_SOP.php](http://www.imagen-europe.com/en/Publications_and_SOP.php), which contain details on ethics,
597 recruitment, neuropsychological tests and preprocessing protocols of MRI data. In brief,
598 task-based functional MRI data were analyzed with SPM12 (Statistical Parametric
599 Mapping, <http://www.fil.ion.ucl.ac.uk/spm>). Preprocessing included: slice time
600 correction to adjust for time differences due to multi-slice imaging acquisition,
601 realignment to the first volume in line, non-linearly warping to the MNI space (based
602 on a custom EPI template (53x63x46 voxels) created out of an average of the mean

603 images of 400 adolescents), resampling at a resolution of $3 \times 3 \times 3 \text{mm}^3$ and smoothing
604 with an isotropic Gaussian kernel of 5 mm full-width at half-maximum. And the
605 activations of each task condition, i.e. happy, angry, neutral faces and control stimuli,
606 were evaluated for each individual at the first-level analysis (The details can be found
607 at https://github.com/imagen2/imagen_processing). And these results were conducted
608 for the further DeCoP analysis.

609 **Data availability**

610 ABCD data are available from a dedicated database: <https://abcdstudy.org>. Human
611 Connectome Projects data are available from: <https://www.humanconnectome.org>. The
612 IMAGEN project are available from a dedicated database: <https://imagen2.cea.fr>.

613 **Code availability**

614 Custom code that supports the findings of this study is available from the
615 corresponding author upon request. All data needed to evaluate the conclusions in the
616 paper are present in the paper and/or the Supplementary Information. Additional data
617 related to this paper may be requested from the authors.

618

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639

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649 **Author Contributions:**

650 Conception or Design of the Study: T.J., T.W.R. and J.F.. Manuscript Writing and
651 Editing: S.X. and T.J. wrote the manuscript; T.W.R. and J.F. edited the first draft; all
652 authors critically reviewed the manuscript. Imaging Data Preprocessing: S.X., C.X. and
653 W.C.. Visualization: S.X., C.X. and T.J.. Data Analysis: S.X., C.X., Z.Z., J.K. and G.S.
654 conducted all the statistical analyses, under the instruction of T.J. and J.F.. Results
655 Interpretation: T.J., T.W.R. and J.F.. Supervision of the Study: T.J. and J.F.. Funding
656 Acquisition: T.J. and J.F..

657 **Competing Interests:**

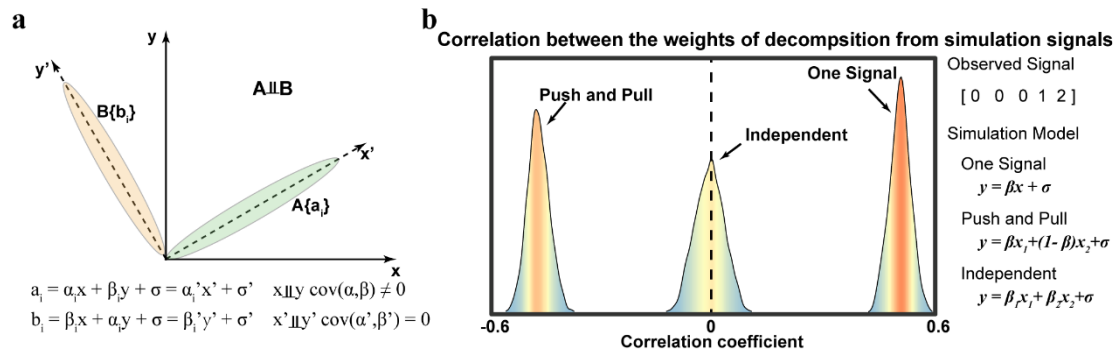
658 The authors declare no competing interests.

659 **Additional Information**

660 Supplementary Information is available for this paper.

661

662 **Extended Data figures and tables**



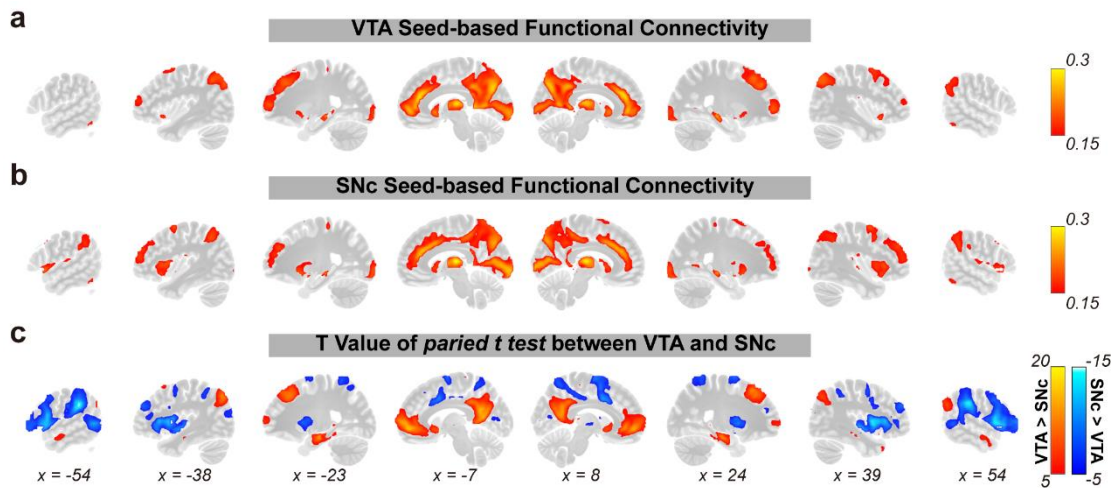
663

664 **Extended Data Fig. 1. a.** Only the “correct” orthogonal vectors could retrieve latent

665 independent signals. **d.** Correlations between decomposed signals based on different

666 simulation models. Also see Extended Data Table 2.

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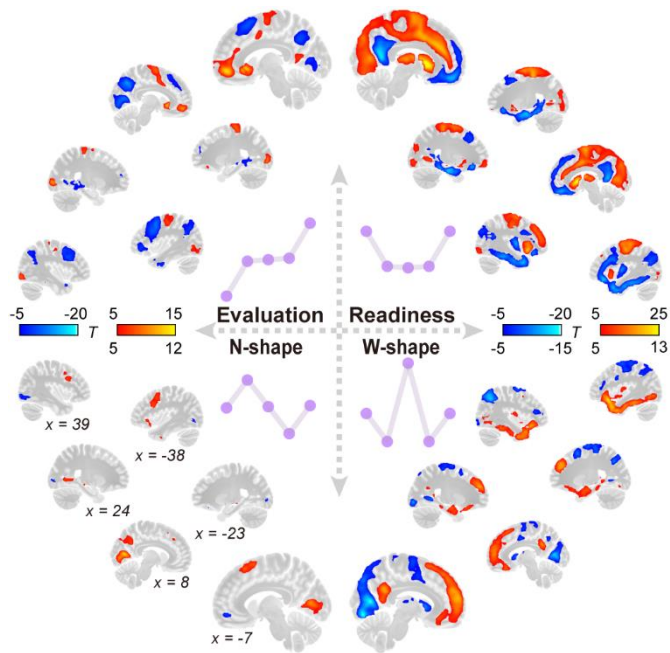
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Extended Data Fig. 2 The brain-wide patterns of seed-based functional connectivity (FC) strength from regions of interest (ROIs): **a.** VTA seed-based; **b.** SNc seed-based; **c.** The T-map of the differences between seed-based FCs patterns from VTA and SNc.



673

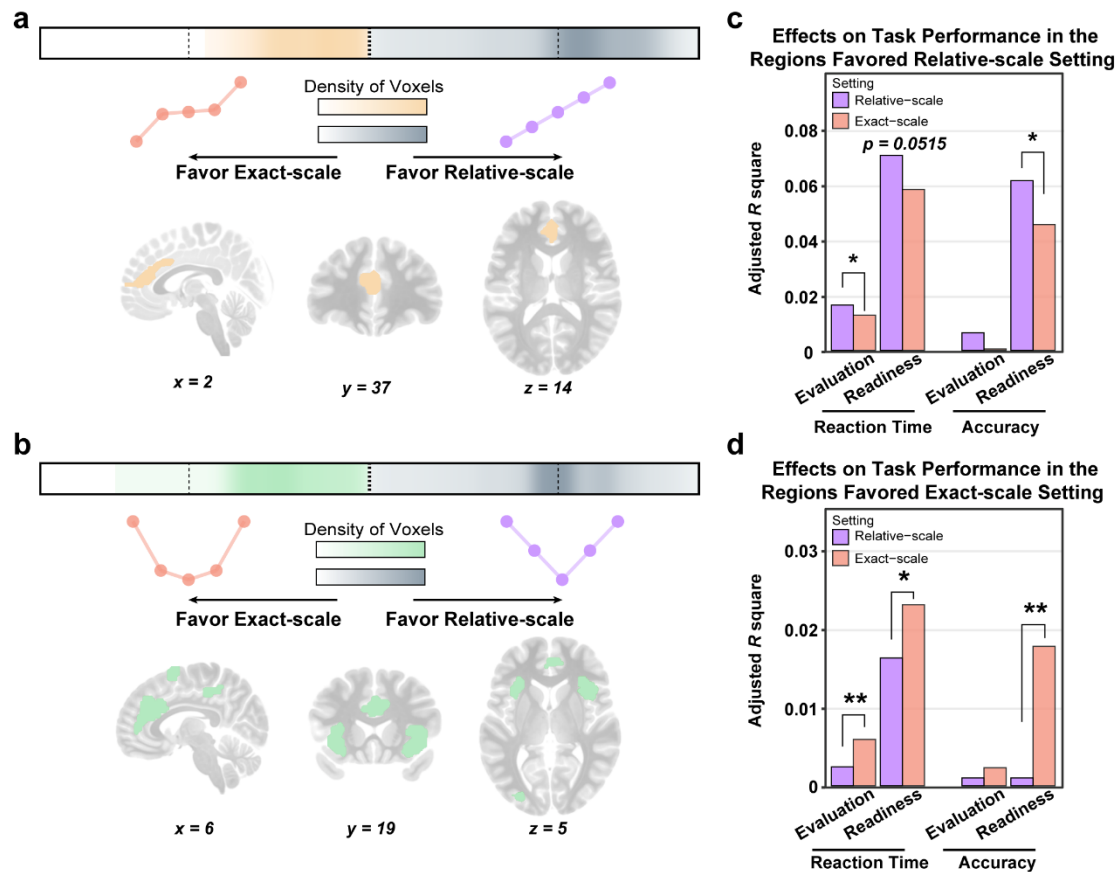
674 **Extended Data Fig. 3** Brain-wide neural representations of signal decomposition with

675 the exact-scale setting (e.g. the evaluation contrast takes the form [-5, -0.2, 0, 0.2, 5]

676 from large-loss to large-win, consistent with the parametric nature of the exact

677 monetary magnitude in the experimental design).

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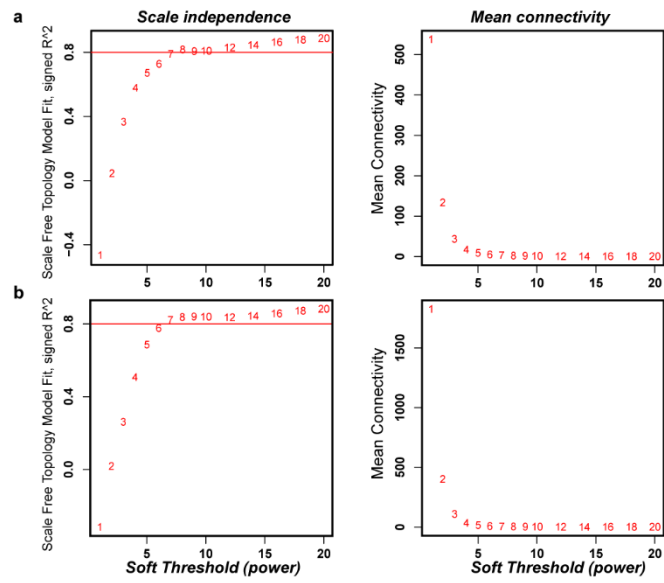
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Extended Data Fig. 4 The bimodal distribution of voxels favoring the exact-scale or relative-scale settings in evaluation (a) and readiness (b) processing. The typical regions favoring the exact-scale setting were illustrated in the corresponding lower subplots. c&d. The favored neural representations (i.e. relative-scale vs exact-scale) demonstrated better predictions for task performance. *significant at level 0.05, ** significant at level 0.01.

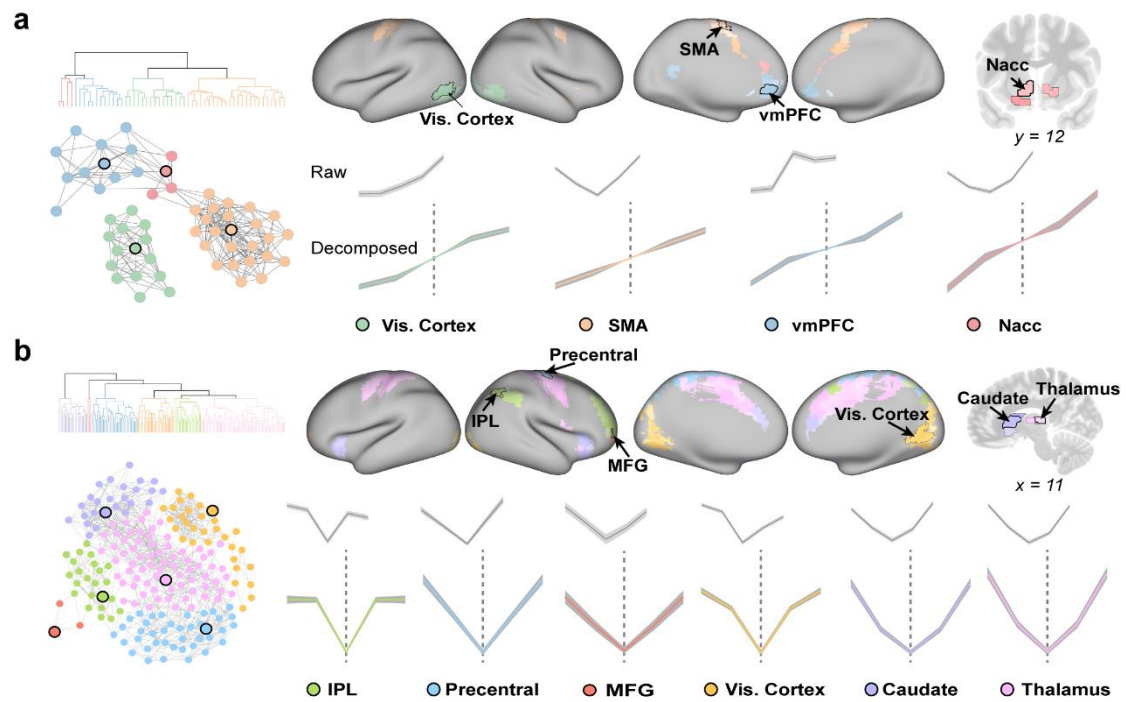


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Extended Data Fig.5 Plots of soft-threshold for WVCNA.

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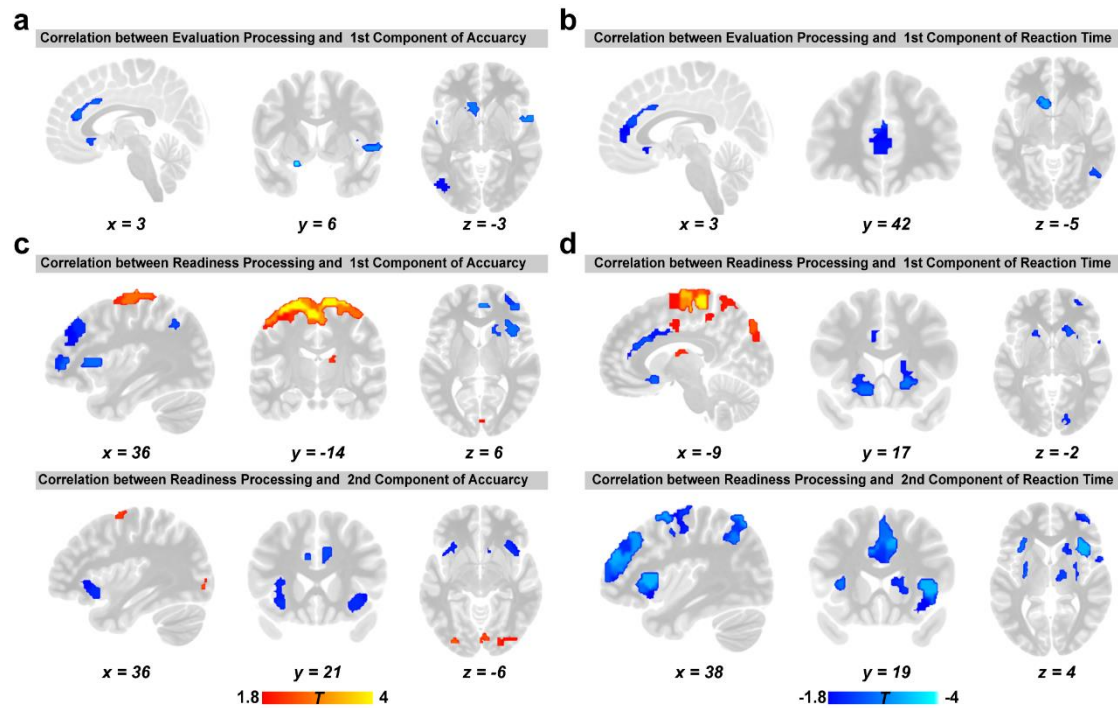
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Extended Data Fig. 6 The brain clusters and subnetworks of evaluation (a) and readiness (b) processing. Also see Extended Data Table 3. The core brain region of each subnetwork is marked by a black bound and illustrated in both the networks (left) and brain templates (right). Raw and decomposed signals of the core brain regions are presented below the brain.



697

698 **Extended Data Fig. 7** The T-maps of evaluation and readiness processing's impacts
699 on the task performances. The canonical correlation analysis (CCA) was implemented
700 to further segregate signal components that demonstrated differential associations with
701 task performances under different experimental conditions.

702

703 **Extended Data Table 1** Demographic characteristics of the samples in this study.

Characteristics	Total (n=1939)	Site1 (n= 415)	Site2 (n= 362)	Site3 (n= 396)	Site4 (n= 766)
Gender (male)	984 (50.75%)	219 (52.77%)	169 (46.69%)	191 (48.23%)	405 (52.87%)
Race/ethnicity					
White	1473 (75.97%)	282 (67.95% ^{***})	246 (67.96% ^{**})	313 (79.04%)	632 (82.51% ^{***})
Black	39 (2.01%)	3 (0.72%)	10 (2.76%)	18 (4.55% ^{**})	8 (1.04%)
Hispanic	233 (12.02%)	84 (20.24% ^{***})	52 (14.36%)	24 (6.06% ^{***})	73 (9.53%)
Asian	20 (1.03%)	2 (0.48%)	13 (3.59% ^{***})	4 (1.01%)	1 (0.13% [*])
Other/Missing	174 (8.97%)	44 (10.6%)	41 (11.33%)	37 (9.34%)	52 (6.79%)

Superscripted * means the variable differ significantly from the total population on the corresponding variable. * p < 0.05, ** p < 0.01, *** p < 0.001 by Fisher's Exact Tests.

704

705 **Extended Data Table 2** Simulation results of the correlations between the weights of
 706 decomposition.

Observerd Signals	One Signal Model			Push and Pull Model			Independent Model		
	<i>r</i>	<i>T</i> (Readiness)	<i>T</i> (Evaluation)	<i>r</i>	<i>T</i> (Readiness)	<i>T</i> (Evaluation)	<i>r</i>	<i>T</i> (Readiness)	<i>T</i> (Evaluation)
2 1 0 1 2	-0.001	30.232	0.032	-0.448	21.868	0.024	0.010	30.185	0.025
1.8 0.9 0 1 2	0.080	30.033	2.669	-0.449	20.768	1.033	0.000	28.696	1.353
1.6 0.8 0 1 2	0.155	29.875	5.245	-0.449	19.667	2.086	0.000	27.154	2.751
1.4 0.7 0 1 2	0.230	29.652	7.735	-0.449	18.620	3.113	-0.002	25.624	4.087
1.2 0.6 0 1 2	0.296	29.433	10.073	-0.449	17.437	4.138	0.000	24.137	5.484
1 0.5 0 1 2	0.355	29.158	12.230	-0.450	16.356	5.174	-0.001	22.665	6.785
0.8 0.4 0 1 2	0.410	28.779	14.196	-0.449	15.333	6.195	0.001	21.067	8.178
0.6 0.3 0 1 2	0.454	28.442	15.998	-0.450	14.230	7.275	-0.002	19.631	9.527
0.4 0.2 0 1 2	0.492	28.067	17.658	-0.449	13.133	8.253	-0.001	18.060	10.893
0.2 0.1 0 1 2	0.522	27.449	19.051	-0.449	12.062	9.255	0.000	16.603	12.216
0 0 0 1 2	0.542	26.735	20.315	-0.450	10.937	10.321	0.001	15.109	13.553
-0.2 -0.1 0 1 2	0.555	25.962	21.468	-0.449	9.817	11.328	0.000	13.565	14.977
-0.4 -0.2 0 1 2	0.555	24.847	22.457	-0.451	8.789	12.310	0.000	12.072	16.320
-0.6 -0.3 0 1 2	0.547	23.471	23.307	-0.450	7.645	13.390	-0.001	10.559	17.724
-0.8 -0.4 0 1 2	0.523	21.816	24.111	-0.449	6.563	14.409	-0.001	9.154	19.057
-1 -0.5 0 1 2	0.485	19.635	24.733	-0.450	5.440	15.470	0.001	7.570	20.393
-1.2 -0.6 0 1 2	0.429	16.942	25.369	-0.450	4.382	16.495	0.001	6.033	21.723
-1.4 -0.7 0 1 2	0.351	13.595	25.931	-0.450	3.230	17.528	-0.001	4.541	23.125
-1.6 -0.8 0 1 2	0.252	9.561	26.360	-0.448	2.225	18.543	-0.001	3.031	24.491
-1.8 -0.9 0 1 2	0.132	4.959	26.780	-0.449	1.097	19.576	0.000	1.490	25.792
-2 -1 0 1 2	0.001	0.014	27.205	-0.449	-0.028	20.638	-0.001	0.000	27.234

707

708 **Extended Data Table 3** Coincidence degree of both evaluation and readiness
 709 processing for the task performance under the different model settings.

Models		Reaction Time				Accuracy			
		Adj. R ²	P	Adj. R ² diff.	P _{adj.*}	Adj. R ²	P	Adj. R ² diff.	P _{adj.*}
<i>All Modules</i>									
Relative-scale Models	Readiness (control W-shape)	0.0849	0.0000	0.0119	0.0476	0.0617	0.0000	0.0123	0.0506
	Evaluation (control N-shape)	0.0252	0.0138	0.0087	0.0646	0.0055	0.1607	0.0044	0.0577
Exact-scale Models	Readiness (control W-shape)	0.0730	0.0000			0.0494	0.0000		
	Evaluation (control N-shape)	0.0165	0.0044			0.0011	0.5570		
<i>Modules Favored Relative-scale Models</i>									
Relative-scale Models	Readiness (control W-shape)	0.0706	0.0000	0.0124	0.0349	0.0614	0.0000	0.0159	0.0196
	Evaluation (control N-shape)	0.0164	0.0037	0.0037	0.0515	0.0063	0.1245	0.0058	0.0360
Exact-scale Models	Readiness (control W-shape)	0.0582	0.0000			0.0455	0.0000		
	Evaluation (control N-shape)	0.0127	0.0080			0.0005	0.4398		
<i>Modules Favored Exact-scale Models</i>									
Relative-scale Models	Readiness (control W-shape)	0.0163	0.0004	-0.0068	0.0463	0.0010	0.3637	-0.0168	0.0021
	Evaluation (control N-shape)	0.0024	0.0108	-0.0035	0.0073	0.0010	0.0949	-0.0013	0.1645
Exact-scale Models	Readiness (control W-shape)	0.0231	0.0000			0.0178	0.0002		
	Evaluation (control N-shape)	0.0059	0.0041			0.0023	0.0767		
* Adjusted P values based on 10000 bootstrap; Adj. R ² diff. The difference between the adjusted R square of relative-scale and exact-scale models.									

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