

Fractionation of reward processing into independent neural representations by novel decoding principle

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Title: Fractionation of reward processing into independent neural representations by novel decoding principle

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Abstract:

How to retrieve latent neurobehavioural processes from complex human neurobiological signals is an important yet unresolved challenge. Processing value and salience information are fundamental yet mutually confounded pathways of reward reinforcement. Here, we develop a novel analytical approach, orthogonal-Decoding multi-Cognitive Processes (DeCoP), to decode brain-wide responses into spatially overlapping, yet functionally independent, evaluation and readiness networks during the anticipation of reward or punishment. Our findings indicate that this functional independence is modulated differentially by those neural systems innervated by the cortico-mesolimbic and nigro-striatal dopamine projections. The segregation of evaluation and readiness neural responses from a unified input signal was also achievable using a theoretical neuronal population-coding model, hence further advancing our understanding of the neural basis involved in motivational processing. Furthermore, our novel theoretical approach could potentially be applied more generally to decode multiple latent neurobehavioral processes and thus advance both the design and hypothesis testing in cognitive task paradigms.

1 **Main**

2 The brain frequently engages parallel processing involving different latent
3 behavioural processes mediated by functionally distinct, though spatially overlapping,
4 neural networks ¹. Previously, human functional neuroimaging studies have had
5 difficulty in unravelling these processes from basal compound physiological signals ²⁻
6 ⁴, which has made it difficult to build process-specific and mechanistic models of the
7 brain ⁵.

8 Reward/punishment processing is perhaps the most adaptive function of the
9 behavioural control system, optimising outcomes through both positive and negative
10 reinforcement ⁶. Recent overarching frameworks propose two different cognitive
11 processes engaged in parallel during rewarded or punished behaviour, namely
12 evaluation (i.e. scaling signal values from reward to punishment) and response
13 readiness (subsuming arousal and attentional salience, contributing to response
14 preparatory processes) ^{6, 7}. The evaluation process is essential for guiding upcoming
15 action selections based on their value, for which the brain has evolved dedicated
16 regions/circuits ⁸⁻¹³. Complementary to evaluation, both reward and punishment, as
17 highly salient events, attract greater attention than neutral stimuli, also engaging greater
18 levels of motor preparation and emotional arousal ¹⁴⁻¹⁶, hence contributing to response
19 (including motor) readiness. Therefore, evaluation and readiness signals are inevitably
20 confounded with each other during reward/punishment processing. Unfortunately,
21 decomposing this compound signal, for example in human fMRI studies, has proven
22 challenging because these two components cannot be identified by using only reward
23 (or only punishment) stimuli in many experimental paradigms. Previous attempts have
24 been made to overcome this problem through identifying spatially dissociated brain
25 regions specific for either evaluation or readiness ^{2, 3, 17}. However, these approaches
26 have failed to disentangle signals in brain regions known to encode both evaluation and
27 readiness signals, for example, in the striatum and the ventromedial prefrontal cortex
28 (vmPFC) ^{7, 16, 18}. Further, the existing studies have not provided convincing evidence to
29 clarify the neural basis for the assumption of functional independence of the evaluation
30 and readiness processes.

1 In the present study, through an innovative approach, we have achieved a brain-
2 wide voxel-wise decomposition of process-specific neurobiological representations of
3 complex neurobehavioural processes for the first time. By implementing this method,
4 we identified two functionally independent yet spatially overlapping neural networks
5 that process evaluation and readiness signals, implicating distinct dopaminergic
6 pathways from VTA and SNc, respectively. We further demonstrated that most brain
7 regions only encoded the relative rank of the signal stimuli, except for the dACC and
8 insula, which coded precise input signals. Finally, based on these results, we established
9 an integrated neural network model that could encode the value-related stimuli (i.e.
10 monotonically from punishment through neutral to reward) into two separate outputs
11 of value and salience signals, hence further advancing our understanding of the neural
12 mechanisms involved in reward/punishment processing.

13 **Experiment design and behaviour outcomes**

14 A monetary incentive delay (MID) task (Fig. 1a), one of the classical and widely
15 used fMRI paradigms for reward processing, was conducted in 1939 children aged 9-
16 10 from the ABCD study¹⁹ (Extended Data Table. 1) to assess reward/punishment
17 processing with gain/loss conditions for small or large amounts of money and a neutral
18 condition (i.e., -5.0 \$, -0.2 \$, 0, 0.2 \$ or 5.0 \$). Briefly, cues are shown to indicate the
19 trial condition as well as the amount of money involved at the anticipation phase.
20 Participants are then shown a target stimulus and asked to respond to gain reward or
21 avoid loss after a jittered anticipatory delay.

22 Overall, participants demonstrated better task performance in responding to both
23 reward and punishment stimuli (Accuracy: Reward vs Neutral, $t_{1,927} = 38.96$, $p < 1E-$
24 245 , Punishment vs Neutral, $t_{1,927} = 29.41$, $p < 1E-156$; Reaction Time: Reward vs
25 Neutral, $t_{1,927} = -10.98$, $p < 1E-26$, Punishment vs Neutral, $t_{1,927} = -9.69$, $p < 1E-20$; Fig.
26 1b). Further, using a monotonic contrast [-2, -1, 0, 1, 2] to represent the step-wise
27 increment of signal values and a corresponding V-shaped contrast [2, 1, 0, 1, 2] to
28 represent the salience effect of input at each condition (i.e. the absolute value of signals)
29 (Fig. 1c), we estimated the contributions of both contrasts to task performance through

1 a general linear model at the individual level, and found both higher accuracy and faster
2 reaction times were strongly associated with a higher salience effect (Accuracy: $t_{1,927} =$
3 $42.74, p < 1E-281$; Reaction Time: $t_{1,927} = -14.23, p < 1E-43$) but not with the signal
4 value itself, i.e. no difference between reward and punishment trials (Accuracy: $t_{1,927} =$
5 $1.87, p = 0.06$; Reaction Time: $t_{1,927} = -1.63, p = 0.11$). Such a hugely differentiated result
6 hence implied that our brain may have different dedicated neural circuits underlying
7 the processes of value and salience signals.

8 **Decomposed neural representations**

9 We next introduced a novel approach, the orthogonal-Decoding of multi-Cognitive
10 Processes (DeCoP), to identify the neural representation of evaluation and readiness
11 encoding in the anticipation phase. The central idea is to note that there are five
12 experimental conditions that can evoke condition-specific neuronal responses. This
13 means that there exists four orthogonal contrasts or patterns of responses over the five
14 conditions (i.e., number of conditions minus one for the constant term). Crucially, these
15 orthogonal contrasts should have a clear interpretation, by design, in terms of latent
16 neurobehavioural processes. For instance, in the second-level analyses of BOLD
17 signals, we again implemented the two plausible contrasts evaluation (i.e., [-2, -1, 0, 1,
18 2]) and readiness (i.e., [2, 1, 0, 1, 2]) that respectively reflected putative independent
19 hypothetical processes of value and salience information across five task conditions. It
20 should be noted that not only are both evaluation and readiness contrasts orthogonal to
21 each other (i.e. their covariance equals 0), but their complementary orthogonal contrasts
22 (i.e., the N-shape model: [-1, 2, 0, -2, 1] and the W-shape model: [1, -2, 2, -2, 1]) were
23 also available and could explain information not accounted for by the hypothetical
24 contrasts for evaluation and readiness (Fig. 2a). We were thus able to retrieve the
25 decomposed signal components of those underlying confounded processes and assess
26 the significance of their respective (and collective) expression over the entire brain.
27 (Extended Data Fig. 1a, see Methods for more details). In our initial report of the results,
28 we will focus on evaluation and readiness components and then place them in the larger
29 context of a four-dimensional space of reward processing in the human brain.

1 Notably, the vmPFC (Brodmann area [BA] 10-11; Peak MNI: [-9, 49, -9], $t_{1,927} =$
2 15.69, Cohen's D = 0.36; Cluster: 615 voxels, $p_{\text{FWE-corr}} = 1.87\text{E-}08$) and ventral striatum
3 (VS, Peak MNI: [-7, 25, -3], $t_{1,927} = 14.21$, Cohen's D = 0.32; Cluster: 634 voxels, $p_{\text{FWE-}}$
4 $\text{corr} = 1.36\text{E-}08$) were the most prominent regions identified in the evaluation model (Fig.
5 2b upper left & 2c), thus being highly sensitive for tracking the entire dimension from
6 punishment to reward. These areas coincide with the terminal regions of the dopamine
7 neuron projections from the ventral tegmental area (VTA), i.e. the meso-corticolimbic
8 dopamine system²⁰⁻²². For the readiness model, however, the signals were more widely
9 dispersed across cortical and subcortical areas, including motor-somatosensory,
10 salience and attention networks, and regions such as the dorsal striatum (DS, Peak MNI:
11 [8, 10, 4], $t_{1,927} = 32.80$, Cohen's D = 0.75) and thalamus (THA, Peak MNI: [13, -6, 16],
12 $t_{1,927} = 24.66$, Cohen's D = 0.56, Cluster: 2267 voxels, $p_{\text{FWE-corr}} = 1.11\text{E-}16$) (Fig. 2b
13 upper right & 2c), consistent with their engagement in processing both reward and
14 punishment²³.

15 **Respective neural circuits**

16 We then investigated whether the neural representations of evaluation and
17 readiness processes were underpinned by putative neural circuits, in particular those
18 modulated by the midbrain dopaminergic projections originating from either the
19 substantia nigra pars compacta (SNc) or the VTA, which plays a central role in reward
20 prediction and approach^{20, 21}. We found regions of the evaluation model with higher
21 functional connectivity (FC) to VTA than to SNc (*paired t*-test: $t_{183} = 14.84$, Cohen's
22 D = 1.10, $p < 10\text{E-}32$, Fig. 3a), and regions of the readiness model with higher FC to
23 the SNc than to the VTA (*paired t*-test: $t_{183} = 3.63$, Cohen's D = 0.27, $p = 0.0004$, Fig.
24 3a) based on 7T high-resolution resting-state fMRI data from the Human Connectome
25 Project²⁴. Further, we extracted the *t*-maps of the difference between the seed-based
26 FC from VTA and SNc (Extended Data Fig. S2a-c), which was exhibited high
27 similarities, although in opposite directions, with the *t*-maps of both evaluation ($r =$
28 0.22, $p_{\text{adj}} < 10\text{E-}20$) and readiness ($r = -0.12$, $p_{\text{adj}} < 10\text{E-}12$, Fig. 3b) models. Thus, the
29 separate VTA and SNc dopamine projections could be the underlying source of

1 evaluation and readiness processes, respectively.

2 **Independence of evaluation and readiness**

3 We further demonstrated that the above spatially overlapping cognitive processes
4 modulated by distinct neural pathways were indeed functionally independent, which
5 could be directly inferred from uncorrelated signal components at the co-activated
6 regions (see Methods for the detailed proof). Based on our simulation results, only if
7 the compound signals were indeed a combination of independent signals, the
8 decomposed signals could be uncorrelated ($|r_{\text{mean}}| < 0.001$, the ‘Independent’ model).
9 Otherwise, the decomposed signals were highly correlated and hence inseparable as
10 modulations of latent signals (the ‘One Signal’ model: $r_{\text{mean}} = 0.54$; the ‘Push and Pull’
11 model: $r_{\text{mean}} = -0.45$; Extended Data Fig. 1B & Extended Data Table 2).

12 In this study, we found that the signals attributed to evaluation and N-shape models
13 (dependent signals: $r_{\text{mean}} = -0.093$, 95% CIs = $[-0.114 \text{ to } -0.073]$, $p_{\text{bootstrap}} < 0.0001$
14 based on 10000 bootstrap) together described the sensitivity of evaluation from
15 punishment to reward. Further, signals of evaluation and N-shape models were
16 independent (evaluation vs readiness: $r_{\text{mean}} = 0.006$, 95% CIs = $[-0.009, 0.021]$, $p_{\text{bootstrap}}$
17 $= 0.4142$, 99.5% voxels with $r \in [-0.1, 0.1]$, i.e. less than a meaningful small effect size;
18 evaluation vs W-shape: $r_{\text{mean}} = -0.008$, 95% CIs = $[-0.025, 0.009]$, $p_{\text{bootstrap}} = 0.3358$,
19 99.3% voxels with $r \in [-0.1, 0.1]$; N-shape vs readiness: $r_{\text{mean}} = -0.001$, 95% CIs = $[-$
20 $0.015, 0.012]$, $p_{\text{bootstrap}} = 0.8432$, 99.3% voxels with $r \in [-0.1, 0.1]$) of those attributed
21 to the readiness and W-shape models (dependent signals: $r_{\text{mean}} = -0.159$, $p_{\text{bootstrap}} <$
22 0.0001), which together described the differentiated engagement of readiness from the
23 neutral condition to reward/punishment conditions (Fig. 4a). Hence, the observed
24 unbalanced sensitivity towards reward and punishment, such as VS and vmPFC, could
25 be parsed into two spatially overlapped though functionally independent signal
26 components, i.e. a point-symmetric evaluation process and a line-symmetric readiness
27 process (Fig. 4c).

28 **Complementary components from DeCoP**

29 Additionally, the complementary N-shape and W-shape models account for the

1 deviation from the latent evaluation and readiness signals of the proposed evaluation
2 and readiness models respectively. Complementary to the evaluation model, the N-
3 shape model was only observed with significant signals in the primary visual cortex
4 (BA 17; Peak MNI: [4, -81, 1], $t_{1,927} = 7.65$, Cohen's D = 0.17, Cluster: 233 voxels,
5 $p_{\text{FWE-corr}} = 2.91\text{E-}05$, Fig. 2b lower left & 2c). For the W-shape model, as being
6 complementary to the readiness model, the most prominent regions were bilateral
7 anterior insula (aINS, BA 38, Peak MNI: [49, 25, -9], $t_{1,927} = 14.83$, Cohen's D = 0.34;
8 Cluster: 1177 voxels, $p_{\text{FWE-corr}} = 4.41\text{E-}05$) and anterior cingulate cortex (ACC, BA 32,
9 Peak MNI: [7, 49, 22], $t_{1,927} = 11.51$, Cohen's D = 0.26; Cluster: 881 voxels, $p_{\text{FWE-corr}} =$
10 $2.87\text{E-}10$, Fig. 2b lower right & 2c). Therefore, the signal strength of the additional
11 complementary orthogonal contrasts could provide a useful measurement of the
12 distance between the latent independent signals and the proposed models (see Methods
13 for more details). Converging evidence indicated that most brain regions distinguish
14 reward from punishment signals with their relative rank. In other words, the brain
15 simplifies the scales of the different values when processing reward/punishment
16 information, hence processing highly abstract information only. However, the bilateral
17 anterior insula (aINS) and dorsal anterior cingulate cortex (dACC), commonly referred
18 to as the salience network, were most likely tracking the parametric nature of the
19 experimental design (i.e. [-5, -0.2, 0, 0.2, 5]; Extended Data Fig. 3-4 & Extended Data
20 Table 3; also see Supplementary Information for more details). These results were
21 consistent with the role of aINS and dACC in updating and maintaining subjective value
22 information^{25, 26}.

23 **Differential impacts on task performance**

24 We further implemented a weighted voxel co-activation network analysis
25 (WVCNA) to capture the most informative brain-wide signal clusters²⁷ and identified
26 55 and 194 clusters for the evaluation and readiness processes, respectively. These
27 clusters could be further divided into four and five subnetworks by hierarchical
28 clustering, respectively (see Supplementary Information for more details, Extended
29 Data Fig. 5 & Extended Data Table 4). Again, we observed brain-wide low correlations

1 between evaluation and readiness clusters ($r_{\text{mean}} = 0.01$, $p_{\text{bootstrap}} = 0.1621$; Fig. 3B),
2 hence further supporting their neural functional independence.

3 We then investigated the effects of decomposed signals on task performance using
4 the canonical correlation analysis (CCA) ^{28, 29}, and found associations between
5 variations in the neural signal and task performance across three condition categories
6 (i.e. reward, neutral and punishment) for both evaluation and readiness: reaction time
7 (evaluation: adjusted η^2 ($adj\text{-}\eta^2$) = 0.020, $p_{\text{perm}} < 0.001$; readiness: $adj\text{-}\eta^2 = 0.075$, p_{perm}
8 < 0.001) and accuracy (evaluation: $adj\text{-}\eta^2 = 0.025$, $p_{\text{perm}} < 0.001$; readiness: $adj\text{-}\eta^2 =$
9 0.079 , $p_{\text{perm}} < 0.001$; Extended Data Table 5-6).

10 Next, based on identified behavioural components from the CCA results, we
11 investigated how task performance received differential contributions from
12 decomposed latent evaluation and readiness signals, especially in spatially overlapping
13 regions, for instance the vmPFC, dACC, and bilateral striatum. For the evaluation
14 signals, both faster reaction times and higher accuracy demonstrated similar first
15 behavioural components, i.e. loading heavily on both reward and punishment
16 conditions but not on the neutral condition. Both components were associated with
17 reduced sensitivity of evaluation processing in the left VS, vmPFC and dACC (Fig. 5a-
18 b & Extended Data Table 5-6), thus highlighting that the evaluation process is engaged
19 similarly for either the reward or punishment signal, which also helps to explain the
20 lack of difference in task performance between reward and punishment trials. On the
21 contrary, suppressed readiness processing in left VS and dACC was uniformly
22 associated with faster reaction times across all three experiment conditions. We also
23 observed a competing readiness process featuring with a greater engagement of a task-
24 preparatory and motor-response network (i.e. the supplementary motor area SMA and
25 somatosensory-motor cortices) that could led to faster reaction times across all three
26 experiment conditions (represented as the first behavioural component, Fig. 5d right &
27 Extended Data Table 5). In terms of accuracy, however, readiness signals in the right
28 aINS and dACC (i.e. the salience network) produced opposite effects in the presence
29 or absence of valence signals: in reward/punishment trials (i.e. the first behavioural
30 component), higher accuracy was associated with reduced sensitivity (Fig. 5d left),

1 whereas in neutral trials (i.e. the second behavioural component), the relationship was
2 reversed, with higher accuracy associated with more readiness (Fig. 5d right &
3 Extended Data Table 6).

4 In conclusion, we demonstrated that neural processing for evaluation and
5 readiness contribute differentially to task performance. The lack of cognitive
6 associations between task performance and signal values (see Fig 1c) was because
7 neural processing for both evaluation and readiness contributed equivalently to task
8 performance in reward and punishment conditions.

9 **Theoretical neuronal encoding models**

10 Above, we have demonstrated the existence of two spatially overlapping, though
11 functionally independent neural representations in the brain encoding value and
12 salience information, thus indicating two distinct patterns of neuronal response at the
13 cellular level. However, it remains unclear how neuronal populations can encode two
14 distinct aspects of information from the same input. Here, we established an integrated
15 three-layer spike neural network that could translate the same input into value and
16 salience outputs, respectively (Fig. 6).

17 Remarkably, while the models for value and salience outputs were trained
18 individually, both models demonstrate a surprisingly high convergence on the input and
19 hidden layers. Specifically, in both models, neurons in the input layer all had a linear
20 response to the value of signals (i.e. from lower to higher activations according to the
21 signal strength from punishment to reward), and neurons in the hidden layer were
22 formed in two groups: one was activated by punishment inputs (“punishment neurons”),
23 the other being activated by reward inputs (“reward neurons”) (Fig. 6). The co-
24 existence of reward and punishment neurons has been supported by findings across
25 multiple species³⁰⁻³⁴. Further, the connections between the input and hidden layers in
26 both models were also highly similar (value model: $\mu_{\text{weight}} = 0.0187$,
27 excitatory/inhibitory ratio (E/I ratio) = 1.8825, $\mu_{\text{bias}} = 3.7294$; salience model: $\mu_{\text{weight}} =$
28 0.0184, E/I ratio = 1.8409, $\mu_{\text{bias}} = 3.7489$; Extended Data Fig. 6), thus indicating that
29 the input and hidden layers of both neural networks were highly robust and

1 homogeneous, and could be represented by the same group of neurons.

2 Both neural networks mainly differ in the output layers. For the value model,
3 neurons in the output layer were inhibited if connecting to the ‘punishment neurons’ in
4 the hidden layer and activated if connecting to the ‘reward neurons’, hence
5 demonstrating linear output from punishment to reward (Fig. 6 & Extended Data Fig.
6 6). For the readiness model, however, there are two groups of neurons in the output
7 layer: one group is activated when connecting to the “punishment neurons” but
8 inhibited if connecting to the “reward neurons” from the hidden layer. In contrast, the
9 other group was reversely inhibited when connecting to the “reward neurons” but
10 activated if connecting to the “punishment neurons”. Hence, the net outputs for reward
11 and punishment stimuli were equivalent (Fig. 6 & Extended Data Fig. 6).

12 Altogether, this integrated three-layer neural network could encode the same input
13 into distinct value and salience outcomes, thus providing a plausible population
14 neuronal model underlying the spatially overlapping, though functionally independent
15 neural representations of evaluation and readiness, which may provide further insight
16 into the real-world neuronal encoding system.

17 **Discussion**

18 In the present study, we have successfully disentangled two functionally
19 independent (i.e. evaluation and readiness) processes from complex neurobehavioural
20 signals during reward/punishment processing by the novel approach. Our findings
21 provide insight into commonly ambiguous observations in fMRI tasks which involve
22 multiple inferential latent behavioural or cognitive processes. For example, vmPFC,
23 as a key node in the neural circuitry underlying reward processing and value-based
24 decision making³⁵⁻³⁷, was paradoxically ‘inactive’ during the reward vs neutral contrast.
25 This unexpected ‘inactivation’ can now be understood as a product of a trade-off
26 between two independent processes: activation by reward stimuli (i.e. of the evaluation
27 process) and deactivation as part of the default mode network (i.e. of the readiness
28 process). Furthermore, our findings also demonstrated that the independence of
29 evaluation and readiness processing in the brain, putatively is mediated by differential

1 neural circuits including the VTA and SNc respectively. This finding provides novel
2 evidence that the reward processing is linked to the midbrain dopaminergic system and
3 that evaluation and readiness processes involve distinct underlying neural mechanisms
4 ^{7, 22, 38}. Motivated by these findings, we formulated an integrated theoretical neuronal
5 population-coding model to illustrate a plausible neuronal mechanism underlying the
6 segregation of value and salience signals from the same input. The two discrete
7 neuronal populations revealed in the hidden layer, which respectively responds to
8 reward and punishment signals, were also supported by previous studies of dopamine
9 neurons in rodent and primates, especially in the VTA and the VS ³⁰⁻³⁴. We thus
10 proposed a unified dopaminergic signal processing system that could bridge the gap
11 between the neuronal signal encoding (at the cellular level) and the neural
12 representation of encoded signals (at the metabolic or functional level), which may
13 significantly advance our understanding of the neural basis of reward/punishment
14 processing.

15 Further, our novel approach also enables direct statistical inference concerning the
16 functional independence among decomposed latent neurobehavioural processes based
17 on predefined orthogonal latent contrasts. This demonstration may have radical
18 implications for analysing experimental designs of theoretical neurobehavioural models.
19 While existing research frameworks have hitherto attempted to identify spatially
20 separated brain regions that activate specifically under a particular neurobehavioural
21 model ^{5, 18}, such a ‘specific’ activation is highly subjective to the selected threshold for
22 significance and hence may not be truly ‘specific’. More problematically, even if an
23 activation is indeed specific to a particular model and null in the other, their underlying
24 signals might still be related. For instance, while the N-shape model in the present study
25 was barely activated brain-wide, it nevertheless showed strong negative correlations
26 with the evaluation model in voxels that were ‘specifically’ activated for the evaluation
27 models, and hence both signals were not independent of one another. Therefore, the
28 statistical framework provided by our novel approach is vital for any meaningful
29 inference of decomposed independent signals. Moreover, such a statistical framework
30 enables a computational decomposition for any potentially independent

1 neurobehavioural processes, as long as the experimental design allows a meaningful
2 orthogonal decomposition. We therefore expect this new approach to promote new
3 study designs for cognitive processes that are theoretically distinct but almost
4 inseparable through common experimental designs.

5 Finally, our analytical framework could comprehensively describe any possible
6 outcomes across all experimental conditions. Complementary to the proposed
7 theoretical neurobehavioural models, the other orthogonal components specified from
8 the same theoretical framework could also provide valuable information. For instance,
9 in the present study, the complementary N-shape and W-shape models, were highly
10 correlated with the evaluation and readiness components, respectively. Therefore, both
11 components modified the response to small monetary stimuli of the corresponding
12 primary models during reward/punishment anticipation. This led to the finding that
13 dACC encoded the exact monetary magnitude of the experimental design in both
14 evaluation and readiness models, whereas most other regions encode only highly
15 abstracted information. This observation is consistent with the role of dACC in updating
16 and maintaining subjective value information ^{25, 26}. Besides dACC, an unexpected
17 increase in the neutral condition was observed for bilateral aINS, rendering the best
18 model in fitness as the W-shape model, instead of the V-shape readiness model. In fact,
19 the W-shape activation has also been observed in several previous studies on reward-
20 related decision making ^{3, 17}, though without further discussion. The aINS was
21 seemingly involved in two situational processes in regulating task performance, in that
22 higher engagement of aINS could lead to better performance in the absence of reward
23 or punishment signals, but it is the suppressed aINS that was associated with better
24 performance in the reward/punishment trials. This observation may help to understand
25 the W-shape activation of aINS as a combination of two situational processes (for
26 instance, positive and negative V-shape activations). Hence, DeCoP could further
27 strengthen our understanding of latent neurobehavioural processes through additionally
28 retrieving complementary, though not necessarily independent, components.

29 In summary, we have developed and evaluated a novel signal decomposition
30 strategy, 'DeCoP', to dissociate behavioural processes that confound the observation of

1 functional neuroimaging signals. Through DeCoP we demonstrated the functional
2 independence of evaluation and readiness processing in the brain, putatively modulated
3 differentially by neural circuits targeting VTA and SNc. We also proposed an integrated
4 theoretical neuronal population-coding model to illustrate a plausible neuronal
5 mechanism underlying the segregation of evaluation and readiness by distinct midbrain
6 dopaminergic projection systems. We also found that most brain regions encoded
7 signals based on abstract information instead of the observed exact monetary magnitude
8 scale, except for the salience network, i.e. dACC and aINS. We expect that DeCoP can
9 be applied usefully to elucidate many other comparably ambiguous data-sets and
10 enhance experimental designs for complex cognitive processing.

1 **Methods**

2 **Participants**

3 The dataset used for this study was selected from Annual Curated Data Release
4 2.01 (<https://data-archive.nimh.nih.gov/abcd>) of the Adolescent Brain Cognitive
5 Development (ABCD) cohort, which recruited 11,875 children between 9–11 years of
6 age from 21 sites across the United States ¹⁹. The study conforms to each site's
7 Institutional Review Board's rules and procedures, and all participants provide
8 informed consent (parents) or informed assent (children). More details of the subjects
9 and the data collection are provided at the ABCD website
10 (<https://abcdstudy.org/scientists/protocols/>) and are also described previously ¹⁹.

11 Magnetic resonance imaging (MRI) data in the ABCD study were collected
12 from different 3T scanner platforms (i.e. Siemens Prisma, General Electric MR750 and
13 Philips Achieva dStream). To minimise biases introduced by multiple platforms, we
14 only included MRI data from the most frequent manufacturer Siemens Prisma, i.e. 5968
15 participants from 13 sites. By examining the similarity of brain activations across these
16 13 sites, we selected 2326 participants from 4 sites with consistent activation patterns.
17 Furthermore, the data with poor registration (by visual check) and high head motion
18 (mean framework displacement (FD) > 0.5 mm) were excluded. Hence, 1939 quality-
19 controlled participants were included in the following analysis and the demographic
20 characteristics of these participants are summarised in Extended Data Table 1. In the
21 further analyses, the age, gender, handedness, race, site informance and mean FD of
22 participants were included as the covariates.

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

24 A modified version of the MID task was used to examine brain activation during
25 monetary reward anticipation and receipt ³⁹, which consists of five levels of incentive:
26 large loss, small loss, neutral, small win and large win (i.e., -5.0 \$, -0.2 \$, 0 \$, 0.2 \$ and
27 5.0 \$ respectively). In each trial, participants were first presented with one of three cue
28 shapes (circle, square or triangle) that indicated the trial condition (of win, loss, or
29 neutral, respectively), as well as the amount of money involved. This cue presentation

1 (2,000 ms) was followed by a jittered anticipatory delay (1,500-4,000 ms) of fixation
2 on a black crosshair. Subsequently, a blank target cue (with the same shape as the
3 previously presented cue) emerged and required the participants to press a response
4 button before the target disappeared to win or avoid loss. With a tracking algorithm, the
5 time of the target on the screen was dynamically manipulated (i.e., 150 ms-500 ms) to
6 maintain a 60% success rate for each participant. After a short delay, feedback of the
7 current trial (i.e., the amount of monetary gain or loss) and the accumulated reward so
8 far were presented for 1,500-1,850 ms (Fig. 1a). During the anticipation phase,
9 participants underwent 50 trials in total (i.e., 10 trials per incentive degree). Participants
10 had first completed a practice session outside the scanner before completing two
11 sessions of the MID task with fMRI recording (approximately 5.5 minutes each).

12 We first compared the task behaviour outcomes (i.e., the reaction time (RT) and
13 accuracy) between the salient (i.e., reward or punishment) and neutral conditions across
14 the participants by paired *t-test*. We next set a general linear model to estimate the
15 effects of the putative contrasts (i.e., value model: [-2 -1 0 1 2] and salience model: [2
16 1 0 1 2]) on the task behaviour outcomes at the individual level. Then the group-level
17 effects were obtained by one sample *t-test*.

18 **Image acquisition and preprocessing**

19 Imaging protocols were harmonised across sites and scanners. 3-dimensional
20 T1-weighted images (1.0 mm isotropic, TR = 2500 ms, TE = 2.88 ms) were acquired
21 with a gradient-echo sequence for anatomical localisation and high spatial (2.4 mm
22 isotropic) and temporal (TR = 800 ms) resolution MID-task-based fMRI was acquired
23 with echo-planar imaging (EPI) sequence. The detailed MRI acquisition protocol is
24 described elsewhere ¹⁹. All functional images were preprocessed with the same
25 preprocessing procedure by suggested protocols from FMRIB's Software Library (FSL
26 v5.0.9), Advanced Normalization Tools (ANTs v1.9.2) and Analysis of Functional
27 NeuroImages (AFNI v18.3.03). Concretely, the whole preprocessing procedure
28 included the following steps: (i) brain extraction (ANTs antsBrainExtraction), nonlinear
29 registration to MNI space (ANTs antsRegistrationSyN) for structural images; (ii) rigid

1 realignment to adjust for motion (FSL MCFLIRT) and field map correction (FSL
2 TOPUP) for functional images; (iii) co-registration to a high-resolution T1 image and
3 normalisation to 3 mm isotropic MNI standard space (ANTs antsRegistrationSyN) and
4 (iv) spatial smoothing with a 6mm full-width at half-maximum (FWHM) Gaussian
5 kernel (AFNI 3dBlurToFWHM) and detrending (AFNI 3dDetrend).

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

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

18 **Orthogonally Decoding multi-Cognitive Processes**

19 In the present study, we propose a novel approach to decompose each participant's
20 brain activations at varied conditions (denoted as y) with a set of orthogonal basis
21 $\mathbf{x} = (x_1, \dots, x_k)$, where each vector could represent a predefined signal model, e.g.
22 evaluation or readiness. Specifically, 'orthogonal' here means that any pairwise
23 covariances of vectors all equal zero, i.e. $Cov(x_i, x_j) = E(x_i, x_j) - E(x_i)E(x_j) = 0$.
24 In this way, the regression coefficients $\boldsymbol{\beta} = (\beta_1, \dots, \beta_k)$ (i.e. the strength of signals for
25 each individual) estimated from a multiple linear model with all vectors were the same
26 as those estimated univariately (of simple linear models), i.e.

1 $T(\beta_i | y, \mathbf{x}, \boldsymbol{\beta}_{-i}) = T(\beta_i | y, x_i)$, where $T(\cdot)$ stands for the best linear unbiased
 2 estimator.

3 We propose that the above individual level orthogonal decomposition eliminates
 4 spurious correlations of signal components (i.e. β) introduced by related vectors (i.e. x_i
 5 are correlated), thus allowing us to make meaningful inferences regarding signal
 6 independence at the population level. Here, we only describe a simple proof for the
 7 purpose of illustration, where only two orthogonal vectors were involved. Let y_i denote
 8 the vector of activations across all conditions of the i th individual, and x_{1i} and x_{2i} denote
 9 the two predefined independent orthogonal models of input signals, where
 10 $Cov(x_{1i}, x_{2i}) = 0$, and β_{1i} and β_{2i} denote the corresponding regression coefficients, i.e.
 11 the signal strengths. We could then have the following linear model:

$$y_i = \beta_{1i}x_{1i} + \beta_{2i}x_{2i} + \varepsilon$$

12
 13 We first show that orthogonal settings of x_{1i} and x_{2i} are necessary for a meaningful
 14 independent decomposition of signals. The covariance and variance conditioned on x_{2i}
 15 could be easily derived as:

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

17 and

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

18
 19 Similar results can also be acquired for the covariance/variance conditioned on x_{2i} .
 20 Therefore, we establish the following relationship between the least square estimations
 21 (LSEs) of β_{1i} and β_{2i} :

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

1

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

5 Meanwhile, by setting $\text{Cov}(x_{1i}, x_{2i}) = 0$, and realising that the LSE of β_{1i} in the
6 absence of x_{2i} from the regression model (1) (i.e. reduced to a simple linear model) is
7 $\beta'_{1i} = \text{Cov}(x_{1i}, y_i) / \text{Var}(x_{1i})$, we could immediately have $\beta'_{1i} = \beta_{1i}$. Thus with
8 orthogonal x_{1i} and x_{2i} , estimations from a multiple linear model will be the same as
9 those univariately estimated from simple linear models.

10 We then show that the population-level correlation analyses of β_{1i} and β_{2i} derived
11 above provide meaningful statistical inferences of signal independence. For simplicity,
12 assume that we could rewrite β_{2i} , the signal strength of model x_{2i} , into a sum of two
13 independent components, i.e. $\beta_{2i} = \beta'_{2i} + \gamma_i \beta_{1i}$ where β'_{2i} is independent of β_{1i} , i.e.
14 $\text{Cov}(\beta'_{2i}, \beta_{1i}) = 0$, and the parameter γ_i denotes the proportion of overlapped signals

1 with model x_{1i} , and is independent of β_{1i} , i.e. $Cov(\gamma, \beta_1) = 0$. We could then calculate
 2 the population-level correlation of β_{1i} and β_{2i} as:

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

3
 4 which only equals 0 if either $E(\gamma) = 0$, i.e. the population mean signals of both models
 5 are not a function of each other, or $Var(\beta_1) = 0$, i.e. one of the signals is invariant
 6 across the population, and hence again is not dependent on the other signal. Either way,
 7 the signals of both models are indeed independent, and hence we could have
 8 $Cov(\beta_1, \beta_2) = 0 \Leftrightarrow \beta_1 \perp \beta_2$. In conclusion, this two-stage approach can be used for
 9 statistical inference concerning the dependence of decomposed signals.

10 To illustrate the above theoretical derivations, we conducted simulation analyses to
 11 evaluate the performance of DeCoP on three different complex signal models, namely
 12 the "independent-signal model", where each observed signal consists of two
 13 independent inputs ($\beta_{1i} \perp \beta_{2i}$, and hence $Cov(\beta_1, \beta_2) = 0$), and two dependent-signal
 14 models, i.e. the 'single signal model' with $\beta_{2i} = \gamma_i \beta_{1i}$ (assume $E(\gamma) > 0$, and hence
 15 $Cov(\beta_1, \beta_2) > 0$), and the 'push-and-pull model' with $\beta_{2i} + \gamma_i \beta_{1i} = c_i$ (assume
 16 $E(\gamma) > 0$, and hence $Cov(\beta_1, \beta_2) < 0$). For simplicity, the signal overlapping
 17 parameter γ_i was set as a constant (i.e. 0 for the independent-signal model and 1 in
 18 both dependent-signal models); the observed signals (y_i in equation 1) were simulated
 19 from $[-2, -1, 0, 1, 2]$ to $[2, 1, 0, 1, 2]$ with an increment of $[0.2, 0.1, 0, 0, 0]$ at each step,
 20 i.e. $[-2, -1, 0, 1, 2], [-1.8, -0.9, 0, 1, 2], \dots, [1.8, 0.9, 0, 1, 2], [2, 1, 0, 1, 2]$; and the
 21 orthogonal vectors were fixed as $x_{1i} = [2, 1, 0, 1, 2]$ and $x_{2i} = [-2, -1, 0, 1, 2]$.
 22 Therefore, for the 'single signal model', y_i was directly set as the observed signals
 23 plus a random noise $N(0,1)$. For the 'push-and-pull model', at the i th simulation step,

1 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
 2 could then simulate $y_i = \beta_{1i}x_{1i} + \beta_{2i}x_{2i}$. Clearly, at the population-level, we have
 3 $Cov(\beta_1, \beta_2) = E_i^2(c) - E_i(c^2) = -Var(c) \neq 0$, and hence β_1 and β_2 are dependent. For
 4 the 'independent-signal model', at the i th simulation, by setting $c = 0.05 \times (i-1)$,
 5 $\beta_{1i} = c + N(0,1)$ and $\beta_{2i} = 1 - c + N(0,1)$, we could also simulate $y_i = \beta_{1i}x_{1i} + \beta_{2i}x_{2i}$,
 6 where, however, one would expect $Cov(\beta_1, \beta_2) = 0$. It is notable that while the
 7 simulation models of 'push-and-pull' and 'independent-signal' are rather similar, they
 8 are fundamentally different. The reason lies in the fact that the constant c in the
 9 'independent-signal model' is invariant across individuals and solely determined by the
 10 predefined form of observed signals, and hence the signal strengths β_{1i} and β_{2i} do
 11 not depend on each other. However, in the 'push-and-pull model', the form of
 12 observed signals only determined the expectation $E_i(c)$, and hence both β_{1i} and β_{2i}
 13 are nevertheless the functions of c_i that varies across individuals, and thus are not
 14 independent. For each model, we simulated with 1000 independent individuals for 1000
 15 times, and the detailed results of simulations are shown in Extended Data Fig. 1b and
 16 Extended Data Table 2.

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

18 A total of 184 participants' preprocessed high-resolution (7T) resting-state fMRI
 19 data were collected from the Human Connectome Project (HCP) dataset²⁴. Total FC
 20 (Z-score of Pearson Correlation) from evaluation and readiness activation activations
 21 regions to the voxels in the brainstem (Fig. 3a left), which were masked with AAL3
 22 atlas⁴⁰ and from VTA and SNc to the whole brain were examined respectively. A paired
 23 *t-test* was used to estimate which brain region had a stronger FC with the activation
 24 region. The similarity between patterns of activation and FC was represented by
 25 Pearson's correlation coefficient r . Due to the high correlations among voxels, the
 26 degree of freedom was adjusted to the number of components that accumulatively

1 explain over 95% variance of brain activations based on the principal component
2 analysis.

3 **Measurement of modifying effects**

4 As N-shape and W-shape models can be considered modifiers of evaluation and
5 readiness models, respectively, their strength (i.e. the activation level) can be used to
6 measure the deviance from the proposed model settings, i.e. the exact scale or the
7 relative scale. Thus we computed the standardised mean
8 $((t_{relative} + t_{exact})/\max(\text{abs}(t_{relative}), \text{abs}(t_{exact})))$ to evaluate the prevailing
9 settings, where $t_{relative}$ represents the t-statistics of the activation for N-shape or W-
10 shape models under the relative-scale setting, and t_{exact} represents the t-statistics of
11 the activation for N-shape or W-shape models under the relative-scale setting. When
12 the observed signal falls somewhere between the relative-scale and exact-scale settings,
13 it is readily observable that $t_{relative}$ would be positive and t_{exact} would be negative.
14 Therefore, $\text{abs}(t_{relative}) > \text{abs}(t_{exact})$ would prefer the exact-scale setting, and the
15 standardised mean ranges from 0 to 1, whereas $\text{abs}(t_{relative}) < \text{abs}(t_{exact})$ would
16 prefer the relative-scale setting, and the standardised mean ranges from -1 to 0. It is
17 notable that meaningful settings other than this may also be possible. For instance, the
18 response of the small stimuli could be identical to the large stimuli in the observed
19 signal, thus $t_{exact} < t_{relative} < 0$, and the standardised mean could be well smaller
20 than -1. We further performed *F-test* on the strength of the decomposed components in
21 the brain regions which favour the different scale settings and compared the coincidence
22 degree for the task performance.

23 **Weighted voxel coactivation network analysis (WVCNA)**

24 The R package WGCNA⁴¹ was implemented to conduct the WVCNA²⁷, which
25 identifies activation clusters of both evaluation and readiness models across the brain.
26 The final dataset used for WVCNA included 1,912 participants with 3365 voxels for
27 the evaluation model and 1900 participants with 10932 voxels for the readiness model
28 after removing null data and outliers. We transferred most parameters as default settings
29 from previous studies⁴², except for the soft-threshold parameters, which were set to

1 seven based on the scale-free topology criteria, incidentally identical to those estimated
2 for the MID task from a different cohort ⁴². The stabilities of the generated modules
3 were assessed through bootstrapping. The hierarchical clustering was then applied on
4 the distance matrices among the identified activation modules and, together with the
5 cut-tree function, these modules were further clustered into subnetworks with a static
6 cut at the half of the maximum cluster height (i.e. 1.5 for the evaluation and 2.3 for the
7 readiness).

8 **Canonical correlation analysis (CCA)**

9 CCA has been widely used to investigate the overall correlation between two sets
10 of standardised variables ²⁸. Due to high intra-correlations in both brain networks and
11 task performances, multicollinearity was a potential risk factor jeopardising the validity
12 of the subsequent statistical inference. Therefore, we adopted the ridge-restricted CCA
13 proposed in a previous reference, where the regularised parameter was set as 0.1 ⁴². For
14 each correlation, the P-value or significance level was determined using the
15 permutation test, where the individual IDs of task performances were randomly
16 shuffled at each iteration to generate the null distribution of the corresponding test
17 statistics.

18 Further, we used the adjusted eta square (η^2) to represent the proportion of mutually
19 explained variance between the two sets of variables, as well as controlling for the
20 inflation in η^2 caused by overfitting:

$$21 \quad \eta_{adj}^2 = 1 - \frac{1 - \eta^2}{1 - \eta_0^2}$$

22 where η_0^2 represents the expected η^2 under the null hypothesis that there is no
23 relationship between the two sets of variables (that is, it acts as a measure of inflation
24 in η^2), and can be directly estimated through the permutation test.

25 **Simulation of neuronal population-coding models by artificial neural networks** 26 **(ANN)**

27 We trained a three-layer ANN with full connected fashion which contains 24
28 neurons in each layer and we used Relu function as activate function. For evaluation

1 model networks, we expected the model can map the input $[-20, 20]$ into $[0, 4]$. For
2 readiness model networks, we split the output response into two parts, each part
3 response to positive and negative input respectively. To obtain the identical mean output
4 as evaluation model, the readiness model mapped $[-20, 0]$, $[0, 20]$ into $[2, 0]$, $[0, 2]$. We
5 designed a training task sequence from easy to hard (curriculum learning) in training.
6 For example, we first set 5 points the model need to learn, then 10 points, 100 points
7 and etc. To constrain the distribution of presynaptic weight of each output neuron, we
8 set a regularization term base on the variance of weight matrix of the last layer for
9 training. When visualization, we sorted the position of neurons in hidden layer and
10 output layer base on the sum of their presynaptic weight.

11 **Data availability**

12 ABCD data are available from a dedicated database: <https://abcdstudy.org>. Human
13 Connectome Projects data are available from: <https://www.humanconnectome.org>.

14 **Code availability**

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

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11 Conception or Design of the Study: C.X., T.J., T.W.R. and J.F.. Manuscript Writing
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13 all authors critically reviewed the manuscript. Imaging Data Preprocessing: S.X., C.X.
14 and W.C.. Visualization: S.X., C.X. and T.J.. Data Analysis: S.X., C.X., Z.Z., J.K. and
15 G.S. conducted all the statistical analyses, under the instruction of T.J. and J.F.. Results
16 Interpretation: T.J., T.W.R. and J.F.. Supervision of the Study: T.J. and J.F.. Funding
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18 **Competing Interests:**

19 The authors declare no competing interests.

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Figures:

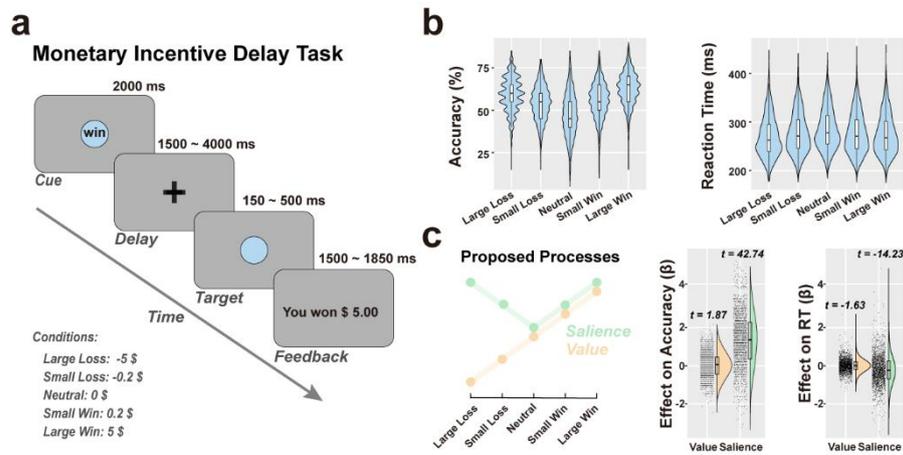


Figure 1. Experimental design and task performances. **a.** The procedure of the monetary incentive delay task (MID); **b.** Distributions of task performance in different conditions; **c.** Proposed value and salience processes and their impacts on task performances.

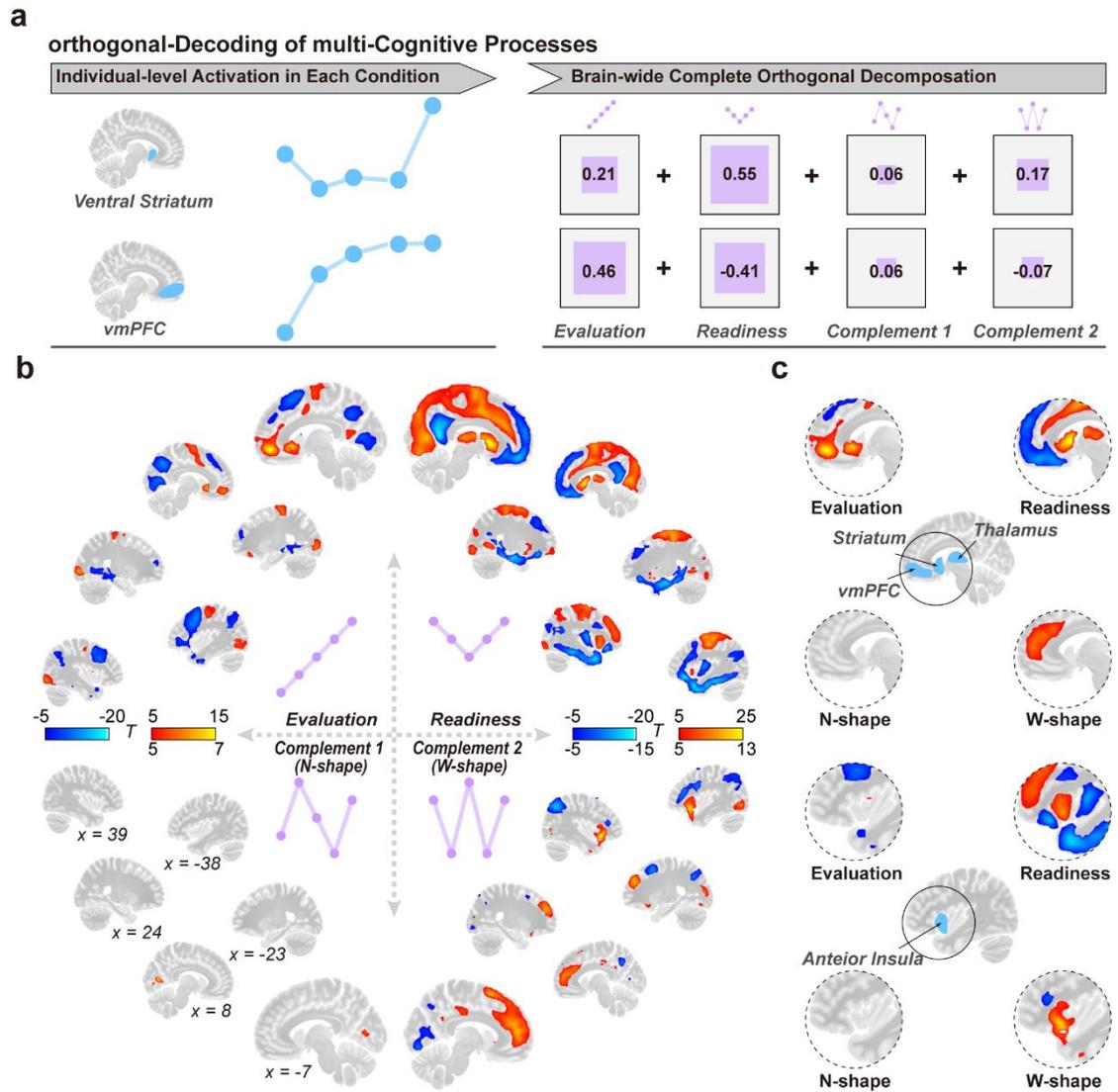


Figure 2. Neural representations of the orthogonal decomposition. **a.** An illustration of the Orthogonally Decoding of multi-Cognitive Process (DeCoP); **b.** Brain-wide T-maps of decomposed signals for orthogonal contrasts. Brain-wide significance was set as $|T| > 5$. The MNI coordinates of brain slices were shown at the lower left; **c.** Decomposed signals in key brain regions.

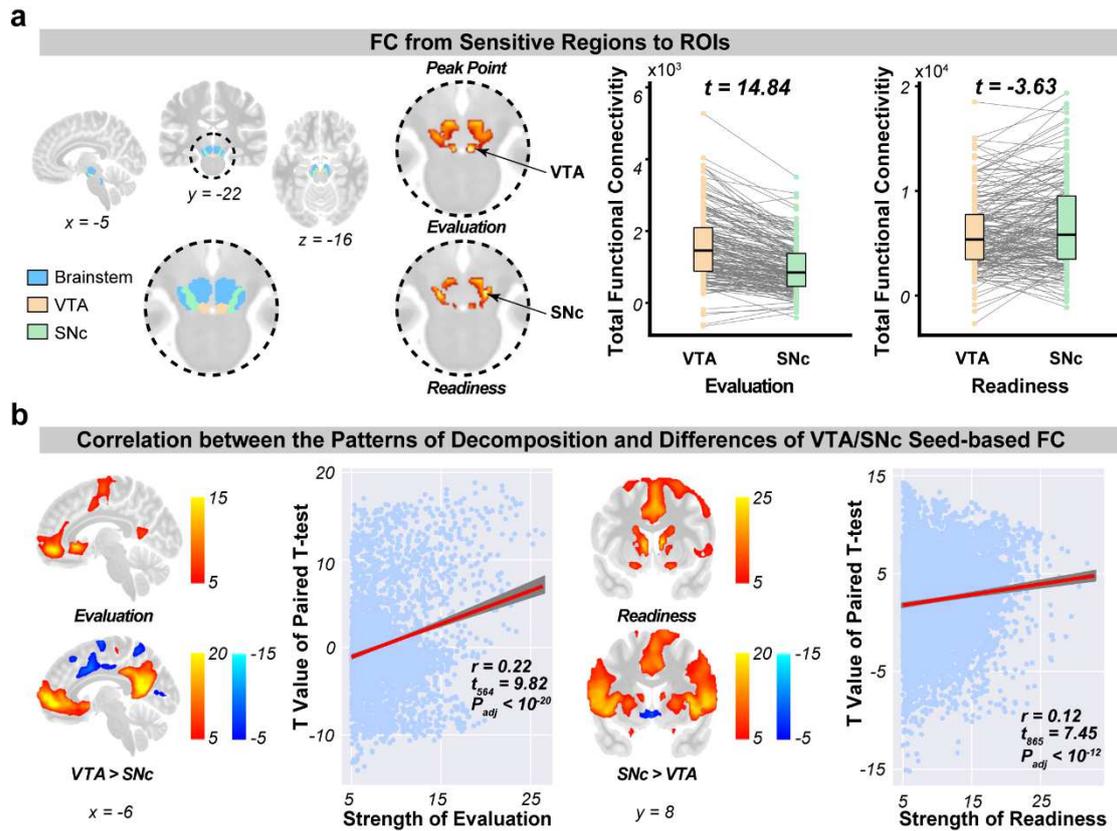


Figure 3. Decomposed evaluation and readiness processes target VTA and SNc neural circuits respectively. **a.** Left: the masks of Brainstem, VTA and SNc from AAL3 atlas; Middle: the strength of functional connectivity (FC) to VTA and SNc from the evaluation and readiness regions identified in Fig. 2B; Right: Paired t-tests between FCs to VTA and SNc from evaluation and readiness; **b.** Brain-wide pattern correlations between the strength of decomposition signals (left: evaluation; right: readiness) and the differentiation of seed-based FCs from VTA and SNc. Brain-wide significance was set as $|T| > 5$.

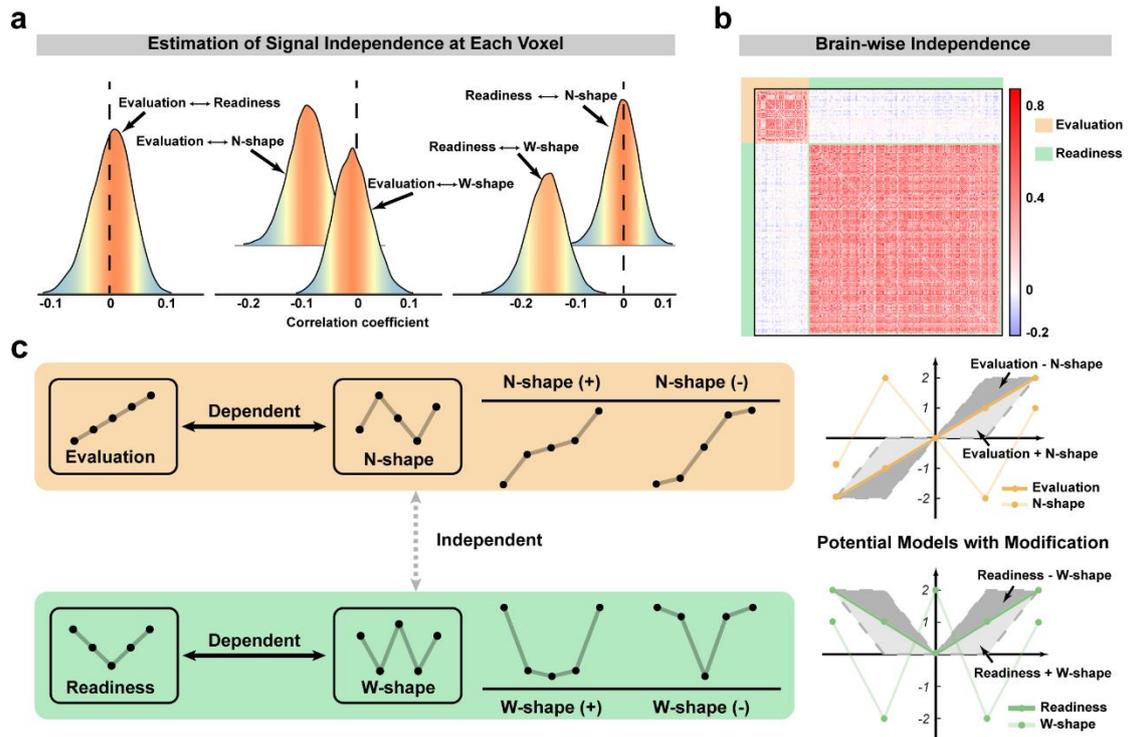


Figure 4. Inference for independence among orthogonally decomposed signals during reward anticipation. **a.** Distributions of pair-wise correlations between signals of orthogonal contrasts at each voxel, where mean correlations deviated from 0 indicate related signals; **b.** The correlation matrix of signals from the evaluation and readiness clusters identified by WVCNA; **c.** An illustration of how related signals could describe the evaluation-related and readiness-related processing.

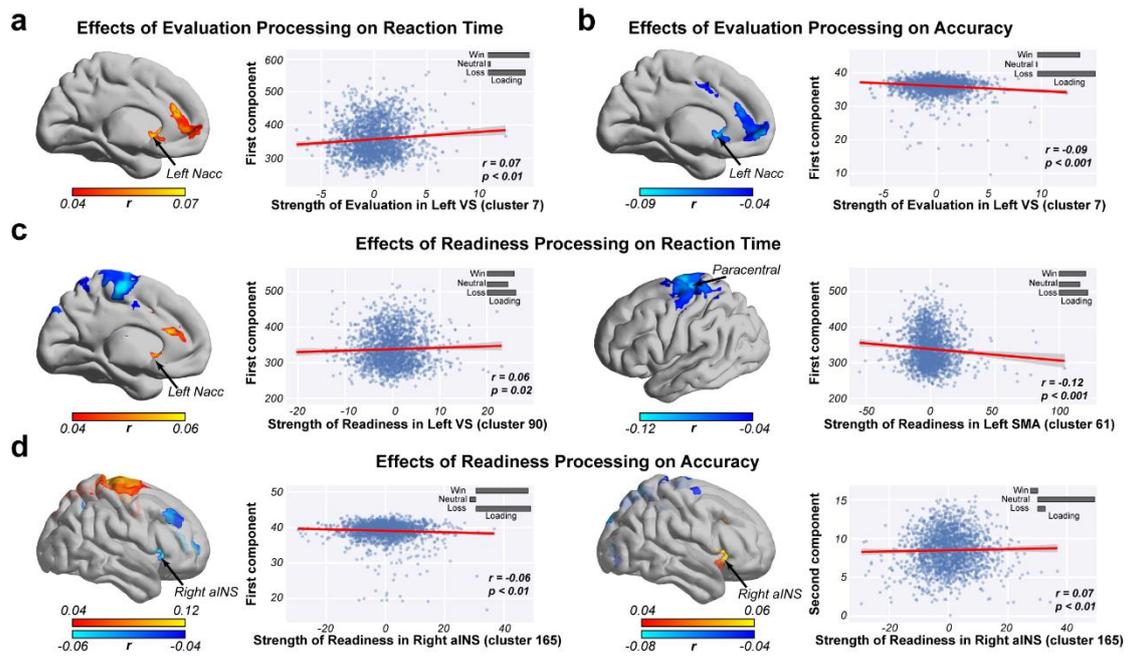


Figure 5. Impacts of evaluation and readiness processing on task performance. The canonical correlation analysis (CCA) was implemented to further segregate signal components that demonstrated differential associations with task performance under different experimental conditions (a-d). The loadings of behavioural components (i.e. of reaction time or accuracy) are shown in the upper right corner of the subgraphs. The black arrow points to the most significant cluster. Also see Extended Data Table 5-6.

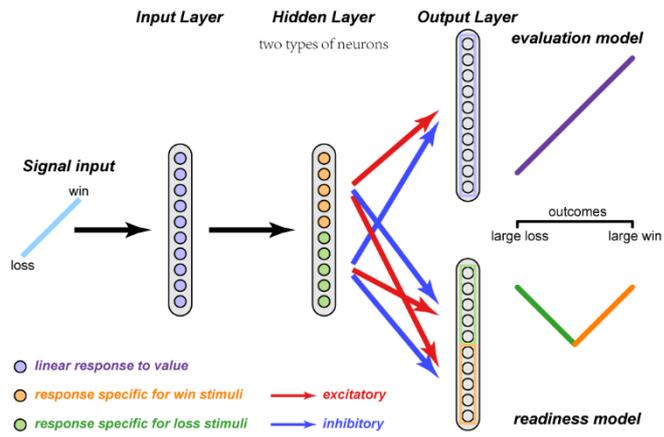
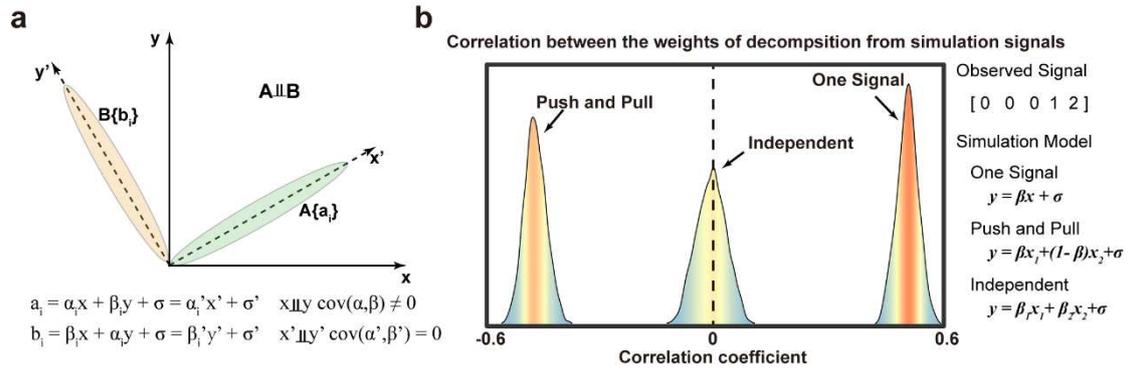


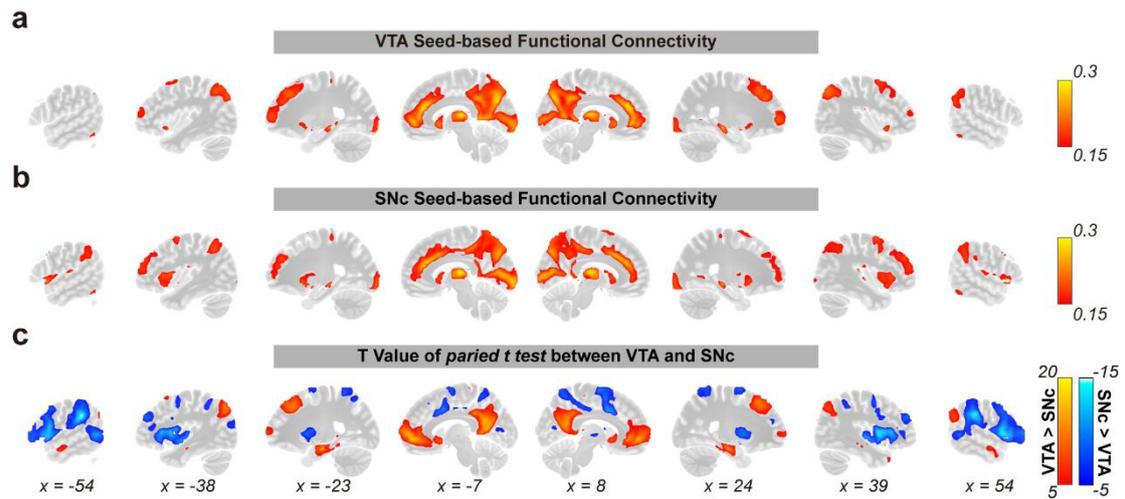
Figure 6. An illustration of the theoretical neuronal encoding model. Although trained individually, both neural networks (i.e. for evaluation and readiness outputs respectively) only differ at the output layer. They hence form an integrated neural network that could decompose the same input (i.e. a value input) into two distinct outputs (i.e. evaluation vs readiness models). Dots represent neurons; red lines represent positive connections, and blue lines represent negative ones. Also see Extended Data Fig. S6.

Extended Data Figures

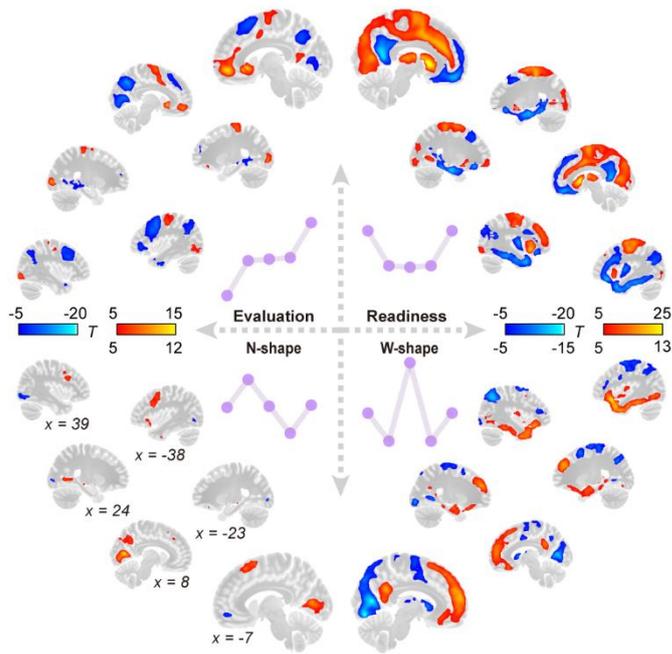


Extended Data Fig. 1. a. Only the “correct” orthogonal vectors could retrieve latent independent signals.

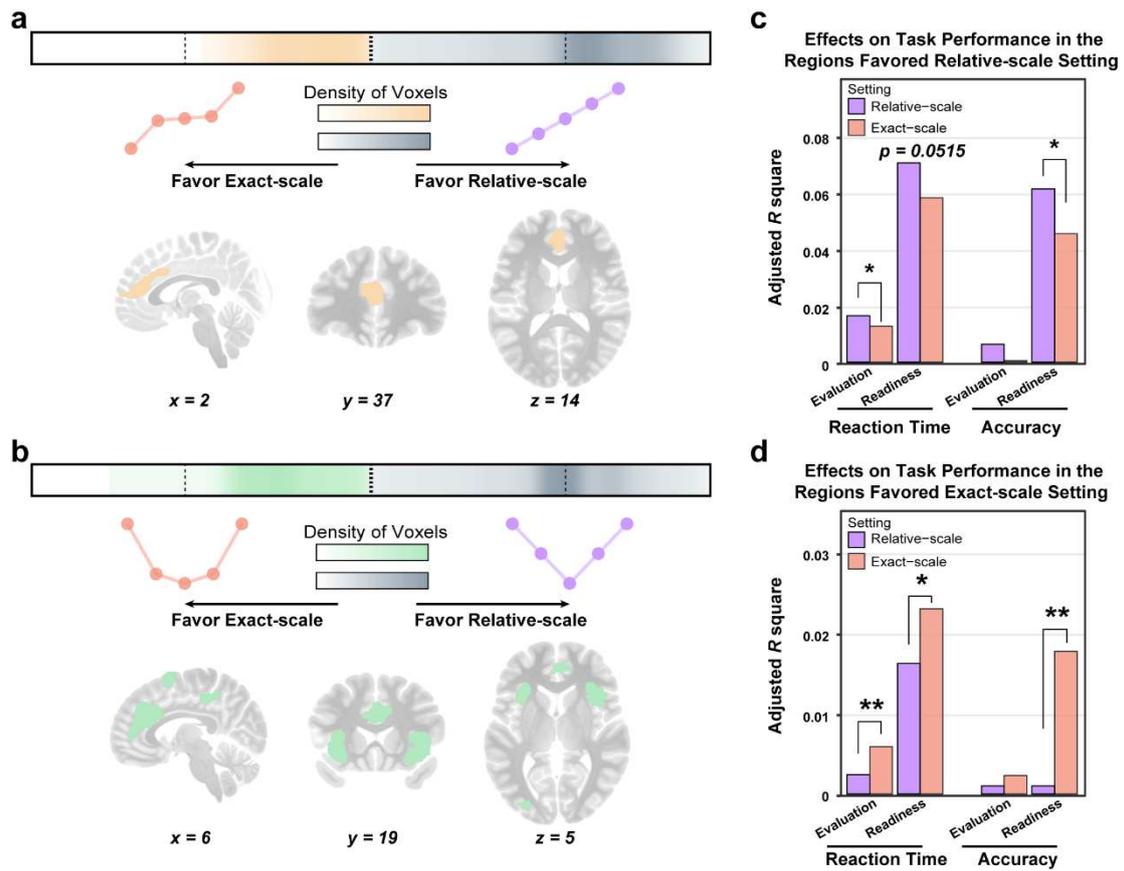
b. Correlations between decomposed signals based on different simulation models. Also see Extended Data Table 2.



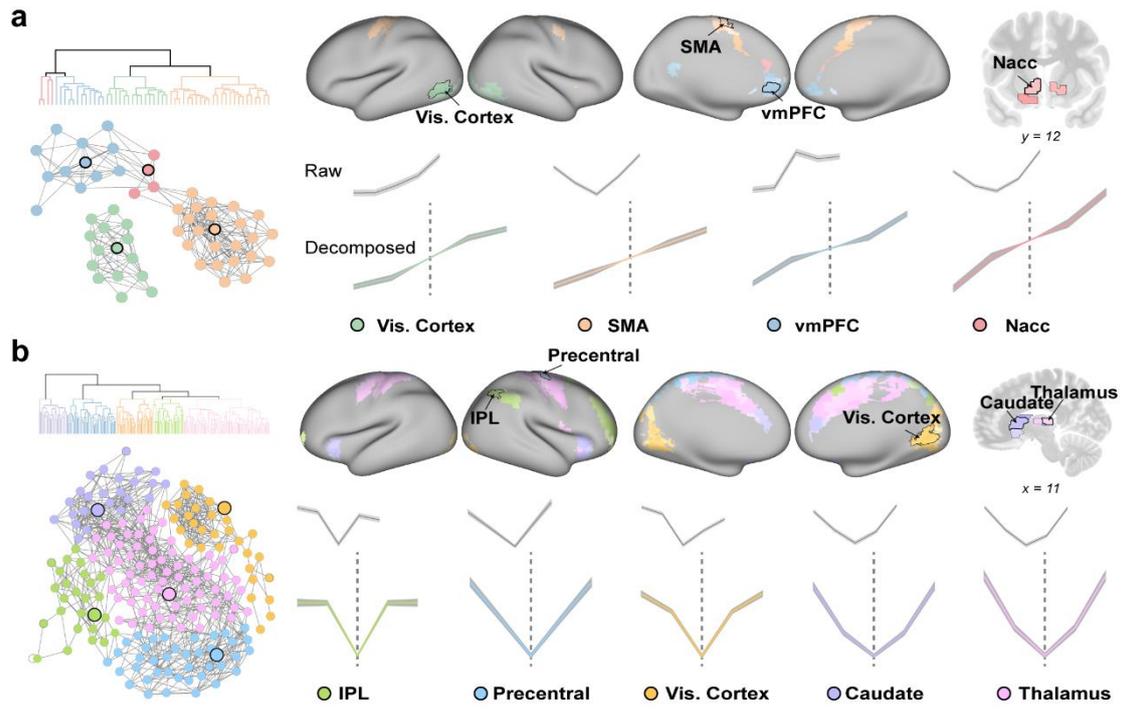
Extended Data Fig. 2. The brain-wide patterns of seed-based functional connectivity (FC) strength from regions of interest (ROIs). a. VTA seed-based FC strength; **b.** SNc seed-based FC strength; **C.** The T-map of differentiated seed-based FC from VTA and SNc.



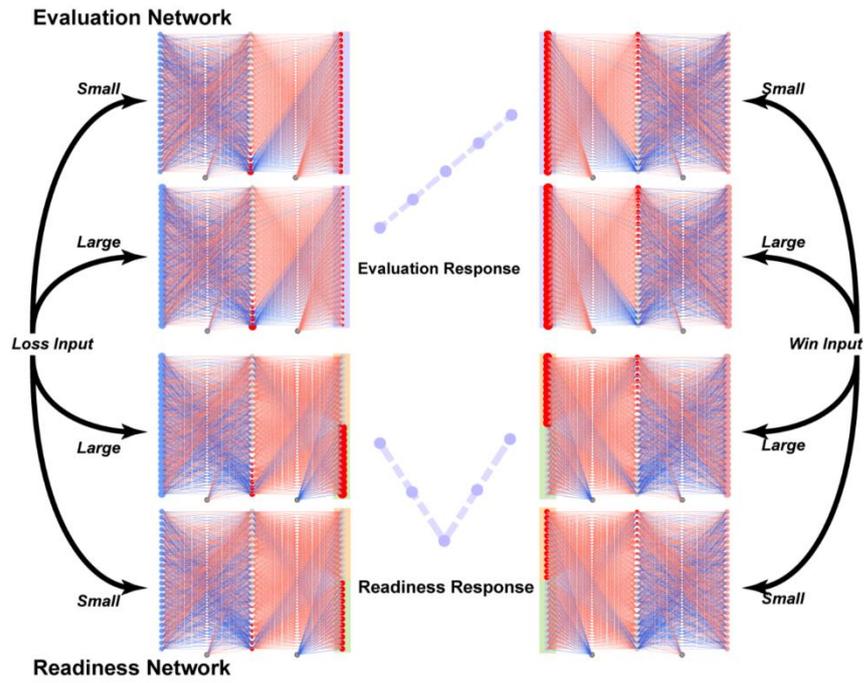
Extended Data Fig. 3. Brain-wide neural representations of signal decomposition with the exact-scale setting (e.g. the evaluation contrast takes the form $[-5, -0.2, 0, 0.2, 5]$ from large-loss to large-win, consistent with the parametric nature of the exact monetary magnitude in the experimental design).



Extended Data Fig. 4. The optimal models for latent neurobehavioural signals. 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 favoured 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.



Extended Data Fig. 5. The brain clusters and subnetworks of evaluation **(a)** and readiness **(b)** processing. Also see Extended Data Table 4. 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.



Extended Data Fig. 6. Graphical visualisation of the artificial neural networks. Dots represent neurons; red lines represent positive connections and blue lines represent negative ones. The red dots in the hidden and output layers represent activated neurons. The larger the dot is, the stronger the neuron activates.

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [ExtendedDataTable16.xlsx](#)
- [SupplementaryInformation.pdf](#)