

Engaging Research Domain Criteria (RDoC): Neurocircuitry in Search of Meaning

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The Research Domain Criteria (RDoC) initiative was implemented to reorient the approach to mental health research from one focused on Diagnostic and Statistical Manual of Mental Disorders (DSM) nosology to one oriented to psychological constructs constrained by neurocircuitry and molecular entities. The initiative has generated significant discussion and valuable reflection on the moorings of psychiatric research. The purpose of this article is to illustrate how a basic or clinical investigator can engage RDoC to explore the neurobiological underpinnings of psychopathology and how a research question can be formulated in RDoC's framework. We utilize a brain region with significant growing interest, the habenula, as an example for probing RDoC's utility. Opportunities to enhance neurocircuitry-psychological construct associations and problems associated with neuronal populations that enable bidirectional circuitry influence are discussed. The exercise reveals areas for further development that could move RDoC from a promising research idea to a successfully engaged foundation for catalyzing clinically relevant discoveries.

Key words: habenula/RMTg/psychiatric diagnosis/translational/DSM/endophenotype

RDoC: A Catalyst for Stalled Progress in Psychiatric Research

Recent efforts by National Institute of Mental Health (NIMH) to break through stalled advancements in the treatment of major psychiatric disorders have led to a reconceptualized research strategy that focuses on constructs of psychology and psychopathology delineated by specific neurocircuitry.^{1–3} The strategy promotes the investigation and characterization of these constructs at various levels of analysis from genes to physiology to behavior. *Constructs* are grouped into 5 larger *domains* (negative valence, positive valence, cognition, social processes, and arousal/modulation) that serve to organize closely linked psychological functions (see [figure 1](#)). For

example, the negative valence domain comprises the constructs fear, anxiety, threat, loss, and frustrative non-reward. Each construct is envisioned to function along a continuum from normal to pathological. Thus, while healthy fear serves a protective function under appropriate conditions, nonspecific, inappropriate, or extreme fear could contribute to psychopathology. Importantly, the domains and constructs are uncoupled from traditional diagnostic categories.

The development of RDoC has not been without its critics^{4–6} and calls for caution—“... it would indeed be unfortunate if the march to freedom from the DSM's ‘epistemic prison’⁷ led merely to a padded cell in the matrix penitentiary.⁸ However, the shift in experimental approach is not intended to replace the current diagnostic classification systems (see specific discussions^{6,8–11}) but rather to provide a road map for research and through this framework to “elaborate a set of psychological constructs linked to behavioral dimensions for which strong evidence exists for circuits that implement these functions, and relate the extremes of functioning along these dimensions to specified symptoms (i.e., impairment).”³

RDoC is explicitly translational and designed as a platform to engage frontline basic neuroscience, clinical neuroscience, and psychiatry. Basic science research is well positioned to investigate the molecular and neuroanatomical substrates of clinically relevant constructs, while clinical research efforts are encouraged to delineate boundaries of construct operation while remaining agnostic to psychiatric nosology. The common ground is laid out in a two-dimensional matrix (<https://www.nimh.nih.gov/research-priorities/rdoc/constructs/rdoc-matrix.shtml>). *Domains* and their underlying *constructs* are organized in rows. Within each row, columns representing *units of analysis* elaborate the genetic, molecular, cellular, circuitry, physiology, and behavioral components of each construct. Importantly, the foundation and justification of each construct is based in large part upon evidence for a neural circuit or system that contributes to an attendant

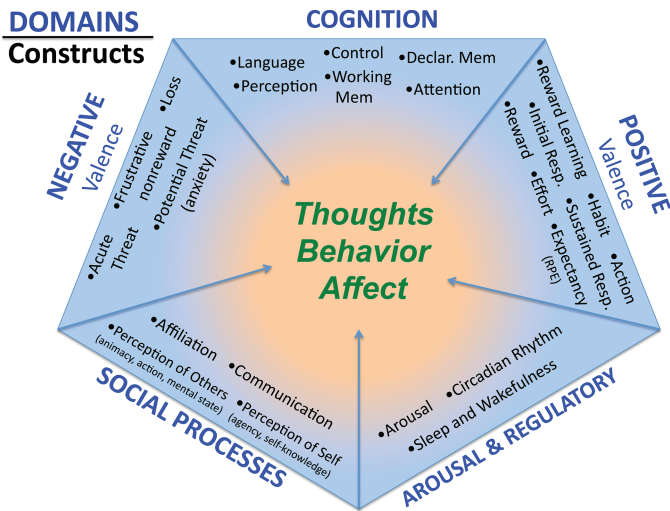


Fig. 1. RDoC envisions sets of psychological *constructs* organized into larger associated *domains*. Individual constructs are thought to function along a continuum from normal to abnormal. The balance of construct functionality contributes to the overall mental constitution of the individual. RDoC, Research Domain Criteria.

psychological function with demonstrated clinical relevance and validity.¹²

The purpose of this article is to illustrate how an investigator or clinician can engage RDoC to glimpse the current state of understanding regarding the neurobiological underpinnings of disease psychopathology and to demonstrate how a research question can be framed in the context of domains, constructs, and circuitry. Our intent is to highlight existing strengths within RDoC and to identify areas for further development that would enhance its practical utility for the scientific community.

Engaging RDoC

There are several practical objectives for which researchers and clinicians may want to engage RDoC. One may be to extract information from the matrix, for example, to answer a specific neurobiological question or to understand a clinically relevant psychopathological construct. For this purpose, simple exploration of RDoC's matrix provides a bird's-eye view of how aspects of human behavior are encompassed as the composite of 5 dynamically maintained domains, each animated by a cluster of associated psychological constructs (<http://www.nimh.nih.gov/research-priorities/rdoc/constructs/rdocmatrix.shtml>). Further exploration reveals the psychological rails that loosely define and constrain the constructs and mechanisms that drive the constructs.

Another objective may be to use it as a standardized platform to develop and test new hypotheses and to integrate new research findings into its current domains, constructs, and units of analysis or perhaps

to develop new ones. Innovative developments in methods to discover regional connections^{13–18} and functional relevance^{19–22} are rapidly advancing our understanding of circuitry dynamics and the mechanisms that drive them. Given the pace of discoveries, ensuring a viable platform to achieve this objective is of paramount importance to ensure that RDoC remains relevant as a research resource and framework for hypothesis generation and testing.

As a means of illustrating a practical engagement with RDoC, we will focus on the habenula (Hb), a region with rapidly growing implications for a broad range of psychopathology. We begin by querying RDoC using the *unit of analysis: circuit* as an entry point (<http://www.nimh.nih.gov/research-priorities/rdoc/units/circuits/index.shtml>). Selecting Hb from the list of brain regions reveals the current domain and construct associations. Selecting a specific construct reveals the information currently available related to each unit of analysis. As of this writing, the only annotation for the Hb is the positive valence domain in the construct/subconstruct “approach motivation/ expectancy_reward prediction error.” Its placement here reflects a body of evidence implicating the Hb in “states that are triggered by internal or external stimuli, experiences or contexts that predict the possibility of reward” and that “the reward expectation can alter the experience of an outcome and can influence the use of resources (eg, cognitive resources)” (<http://www.nimh.nih.gov/research-priorities/rdoc/constructs/expectancy-reward-prediction-error.shtml>). Other brain regions involved in this construct are the amygdala, basal ganglia, dorsal anterior cingulate cortex, orbital frontal cortex, rostral medial tegmentum, substantia nigra (SN)/ventral tegmental area (VTA), and ventral striatum. Notably, a specific circuit is not defined.

Elaborating and Refining RDoC constructs

In the future, especially given the importance of neurocircuitry in delineating constructs, RDoC would ideally enable investigators to curate a list of known afferent and efferent connections derived from various levels of neuroanatomical analysis. In this manner, for example, RDoC could serve as a bioinformatics platform similar to other public and commercial genomics and connectomic platforms.^{23,24} Specific queries could lead to a table of known connections with annotations noting the anatomical method of determination or to searchable diagrams with coded domain and construct associations and embedded annotations containing information from all levels of analysis (see [figure 2](#)). The capacity to query upstream and downstream connections with annotated constructs and domains would be informative at several levels. For example, understanding relevant circuits could inform a query

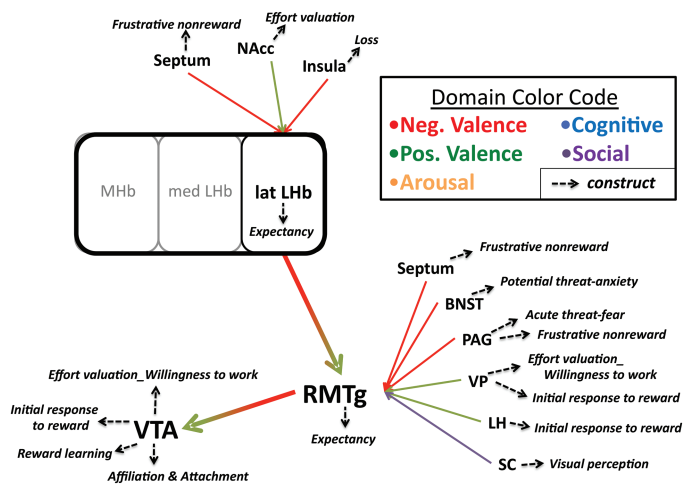


Fig. 2. Example informatics diagram based upon anatomical connections with the Hb. Clicking the region of interest (LHb) would expand afferent and efferent portions of the circuit along with associated construct annotations. Hb, habenula; BNST, bed nucleus solitarius tract; LH, lateral hypothalamus; NAcc, nucleus accumbens core; PAG, periaqueductal gray; RMTg, rostral medial tegmentum; SC, superior colliculus; VP, ventral pallidum; VTA, ventral tegmental area.

relating to psychopathology resulting from brain injury. In anatomical regions with multiple inputs and outputs, querying connected regions could inform clinical expectations and interpretations of comorbidity. Conversely, the bioinformatics approach could drive novel research questions such as, “When circuit A-X-J is activated (or inhibited) does construct Y emerge?” or “When construct Y emerges, is circuit A-X-J activated?” Nuanced questions could be considered when regions with multiple afferent and efferent connections are known. For example, “is the activation of A-X-J similar to A-X-K or B-X-J?”

Refining and elaborating RDoC constructs will require an iterative process involving multiple levels of scientific input. The Hb is an excellent example of a structure in need of elaboration and refinement. Early studies associated the region with a number of functions including sleep, feeding, affect, stress, learning motor activity, and sex.^{25,26} The recent significant rise in interest was fueled by the finding that Hb activation transiently inhibited midbrain dopamine cell firing at a population level^{27,28} and that the unexpected omission of reward increased Hb firing rates.^{29–33} While these findings undoubtedly led to its inclusion within the positive valence domain, Hb activation gives rise to more than a prediction error signal. More recent studies have suggested that the Hb is involved in governing the attribution of incentive salience and in evaluating the cost and benefits associated with different actions.^{34–37} The Hb has also been implicated in avoidance, aversion, and anhedonia with the notion that overactivation gives rise to depressive-like behaviors.^{38–41} Theoretically, high levels of tonic activity within the Hb could drive sustained thoughts of “worse-than-expected”

outcomes or contribute to avoidance and anhedonia, suggesting the region could play a prominent role in one or more constructs contained within the negative valence domain. For example, rumination and negative bias, both behavioral components of the construct loss, could be driven by this activity. Depressive-like somatic symptoms such as sleep disturbances and changes in psychomotor activity and appetite are likewise implicated in Hb function suggesting a potential role in constructs organized within the arousal and regulatory domain.^{42–44} The specific discoveries cited above have implications for the Hb’s role in specific constructs but also in the definition and implementation of circuitry as a unit of analysis within the RDoC framework.

Kozak and Cuthbert³ note that as RDoC evolves, a more satisfactory concept of circuit will be needed and that it could be considered as a “functional unit” that would account for behavioral observations. The Hb provides an excellent example of both the need and difficulty in conceptualizing a circuit in light of its unique interconnections. Recent anatomical findings place a newly characterized region, the rostromedial tegmentum (RMTg), as a critical relay in Hb communication to reward-relevant dopaminergic circuitry.⁴⁵ The RMTg receives major glutamatergic input from the lateral Hb that, in turn, provides dense GABAergic innervation of the SN and VTA.^{46–50} The functional outcome of Hb activation is excitation of RMTg inhibitory neurons that in turn inhibit midbrain dopamine neurons. While the RMTg is strongly activated by stimuli that also activate Hb neurons, the RMTg introduces its own unique hodology into the Hb-VTA/SN pathway via regions that process fear (eg, periaqueductal gray), threat (eg, bed nucleus solitarius tract), avoidance, and anxiety.^{41,45,51–53} Folding the RMTg into existing Hb circuitry adjusts our perspective and prompts different questions related to domains and constructs influenced by this structure and its connections with other brain regions. **Figure 2** shows an example informatics snapshot that would enable an investigator to visualize anatomical connections and construct associations. Exploring upstream and downstream brain regions can point to previously unexplored efferents or parallel circuits that have a similar function but are not directly connected to the region of interest. This information could be the catalyst for hypothesis generation.

Central to the success of this approach is the need to consider what level of detail is required to describe a brain region to fully capture its functional role within a particular construct without also adding an overwhelming level of detail.³ For example, the distinct anatomical connections and behavioral phenotypes attributed to the medial and lateral Hb strongly argue that these regions be considered separately within the RDoC framework.⁵⁴ Recent research suggests the same is true for the VTA, a structure that until recently has been considered relatively homogeneous.⁵⁵

For example, while a projection from the lateral Hb to the VTA via the RMTg has been shown to mediate avoidance behavior,^{51,53} Hb efferents directly targeting the VTA drive approach behavior.⁵⁶ Thus, the same structure, through differences in connectivity, can mediate diametrically opposed behavioral outcomes. Curating these important distinctions within a given brain region is an essential goal and major challenge confronting the RDoC platform.

That a single structure can support diametrically opposed behavioral phenotypes can in some instances be attributed to the firing properties of the constituent neurons. The autogenous electrical properties of Hb and VTA neurons allow them to exhibit spontaneous firing in the absence of synaptic input. Accordingly, their spontaneous “tonic” activity can be modulated bidirectionally. The lateral Hb is particularly well suited for bidirectional modulation, given its unique synaptic architecture that enables a single neuron to exhibit an excitatory or inhibitory response to afferent neurons that carry both glutamate (excitatory) and GABA (inhibitory) neurotransmitters.⁵⁷ Thus, depending upon the dynamics of the input, opposing constructs could be engaged to produce opposite effects on behavior. As RDoC moves forward, some guiding clarification will be needed as similar regions and circuits are likely to be discovered.

What Next

RDoC is an innovative concept that could function as a catalyst to organize and synergize multidisciplinary research efforts and serve as a resource for clinician scientists and as a didactic tool for graduate and residency training.⁵⁸ While RDoC seeks a paradigmatic shift in the way clinical research is conducted, it is likely to alter the trajectory of preclinical research as well. In this regard, less focus will be placed on creating and benchmarking animal models using DSM diagnostic criteria. RDoC’s reliance on phenotypes that are objective, quantifiable, and associated with specific brain circuits make it more compatible with the strengths of contemporary animal research with the added benefit of a higher order, clinically based organization.

In order to achieve its potential, RDoC will have to aggressively confront some of its shortfalls and be resourced to advance its evolution as an informatics platform. It will be important for RDoC to maintain a guided, iterative process to shape and refine constructs and units of analysis and create algorithms for adding newly discovered findings. Specific progress could be made in several areas:

- RDoC would benefit from a commitment to building a research and information platform that would include a larger breadth of annotation at all levels. It has been stated that RDoC was intended as a strategic proposal rather than a content proposal, yet investigators are encouraged to refine and expand the matrix. In our opinion, RDoC cannot live in between these realities.

- While leeway has been granted during its startup to encourage novel approaches, annotations for each unit of analysis should be consistent across domains and constructs—this is especially true for circuits. (For example, the substantia nigra and VTA are collectively listed as “SN/VTA,” “Substantia nigra/VTA,” “Ventral tegmental area/Substantia Nigra,” and “VTA/SN” in different constructs. In addition, some constructs used generic reference (eg, “reward circuit”) vs regions.)
- A new level of circuitry should be added. Currently, most constructs simply list brain regions in the circuit unit of analysis—this is informative at a general level, but a region does not make a circuit, and as discussed above, the growing awareness of subregion connectivity with diverse functional outcome makes this an imperative. At minimum, a 2-component pathway and ideally a 3-piece circuit would significantly enhance information and specificity.
- RDoC circuits need to accommodate bivalent systems operating from a tonic baseline.
- More emphasis should be given to the development of basic science research within the RDoC initiative. NIMH should provide guidance for successful application and encourage basic science research that would dovetail with the RDoC initiative.
- There needs to be a method for curating content that balances the need to incorporate new findings while acknowledging the need for reproducibility. Cloud-sourcing information at various levels (basic vs clinical science and novel vs replicated) may be a means to maintain current state of the art and engender community involvement.

Whether or not RDoC succeeds will ultimately depend on the role assigned to the strategy—research guiding philosophy or engaged dynamic experimental framework—and, of course, the veracity of the assumptions.

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