

## Journal Club

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## Understanding the Hierarchical Organization of Large-Scale Networks Based on Temporal Modulations in Patterns of Neural Connectivity

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Review of Shine et al.

Marsel Mesulam's (Mesulam, 1990) vision that "complex behavior is mapped at the level of multifocal neural systems rather than specific anatomical sites" has triggered an overwhelming interest in understanding the organizational architecture of the brain. Research over the past several decades has shown that the brain is organized in hierarchical networks that shape behavior and cognition, both in healthy and pathological states (Zhou et al., 2006; Honey et al., 2007; Hagmann et al., 2008; Bullmore and Sporns, 2009). At a macroscale level, intrinsic connectivity of parcellated cortical brain regions has been used to characterize neural networks in an attempt to understand their influence on information processing. By examining fluctuations in neural activity, measured by vascular hemodynamic changes, resting-state functional magnetic resonance imaging (rs-fMRI) has reliably been used to investigate network connectivity. Networks are characterized as a collection of nodes and

edges: nodes represented by focal neural parcellations identifiable with three coordinates ( $x$ ,  $y$ , and  $z$ ) and edges as the connections between these nodes (Sporns, 2011). Networks are classified as structural (physical anatomical connections), functional (statistical dependencies among neurophysiological measurements), and effective (one region exerting a causal influence over another) (Friston et al., 2011). Investigation of effective connectivity is a relatively recent development. Corticocortical evoked potential (CCEP) research, which combines invasive (electrode-based EEG recording etc.) and noninvasive (rs-fMRI, diffusion tensor imaging, etc.) methodologies, provides a distinct advantage in such studies by allowing researchers to observe whether stimulation of one site can activate another and drive network variability (Matsumoto et al., 2012).

In a recent article, Shine et al. (2017) combined intracranial stimulation and recording with rs-fMRI to provide valuable insight into the temporal pattern of effective connectivity between three large-scale brain networks: the default-mode network (DMN), the salience network (SN), and the frontoparietal network (FPN). These three networks have received considerable attention due to their large spatial extent, which provides a common framework to understand dysfunction across multiple disorders, especially re-

lated to cognition (Menon, 2011). Additionally, their dynamic synchronization between resting and attentive states, and ease of visibility, offer a unique opportunity to study complex network dynamics at rest.

Shine et al. (2017) recruited 7 subjects that had depth electrodes implanted for a clinical epilepsy study. Each intracranial electrode was individually assigned membership to a predefined network using rs-fMRI. A pair of electrodes within the nodes of predefined networks were stimulated while evoked responses were recorded from each other pair of electrodes, for a duration of 200 ms. The authors found marked heterogeneity in the network-related temporal patterns across subjects; therefore, they used a data-driven approach to identify three spatially and temporally distinct clusters of evoked activity patterns across the 7 subjects. The unsupervised nature of the latter approach adds strength to the study because it does not depend on any a priori hypothesis for expected results, yet the identified clusters aligned with predefined network topology.

Shine et al. (2017) provide evidence that signal propagation among predefined networks occurs along distinct temporal scales. Understanding the temporal modulation of a network can provide unique perspectives on how information is

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transmitted and influenced. The most significant network pattern within the first cluster (which occurred between 11 and 69 ms after stimulation) was evoked within the SN when electrodes within the FPN and SN were stimulated. The second cluster (82–125 ms) was generated in the FPN as a result of stimulating electrodes within the SN and DMN. The third cluster (132–200 ms) was generated by stimulating electrodes in the DMN, which lead to a more general activation across all three networks: the FPN, SN, and DMN. The authors' *post hoc* analysis revealed that stimulation within the FPN and SN led to maximal activations in the early window after stimulus, whereas the DMN generated maximal activation in the delayed window after stimulus. The authors also replicated a finding from Keller et al. (2014) demonstrating that intranetwork stimulation generated higher amplitude activation (within the DMN and FPN but not SN) compared with internetwork stimulation. Temporal network connectivity at the individual subject level showed similar activation patterns (i.e., early FPN/SN and late DMN responses).

Based on the Shine et al. (2017) study, three temporal characteristics may be interpreted to the SN and FPN interaction. First, the fast-evoked activation of the SN (Shine et al., 2017, their Fig. 3) is consistent with this network's key roles (i.e., monitoring and detecting salient events and subsequent access to attention and memory resources) (Menon, 2011). This efficient SN modulation may be supported by a unique regional advantage. The von Economo neurons in the fronto-insular and cingulate cortex (Dijkstra et al., 2018) are rapidly conducting projection neurons. It is suggested that these neurons support sophisticated social-emotional-autonomic functions (Seeley, 2008). Hence, one may conclude that the fast SN activation is in part supported by its distinct cytoarchitecture, which has evolved to facilitate an efficient, "knee-jerk" assessment of changing (external and internal) environmental cues (Menon, 2011). Second, the SN appears to be tightly coupled with other networks: the DMN, but more so the FPN. This is evidenced by its prevalence across all three temporal clusters. This adaptive coupling reinforces the proposed active role of SN in network switching (Menon, 2011) and mediating internetwork connectivity (Sridharan et al., 2008). Third, FPN stimulation leads to SN activation in the early temporal cluster, suggesting that FPN depends on the SN for information acquisition. FPN has

been associated with initiating and adjusting cognitive control based on salience information (Dosenbach et al., 2008).

Shine et al. (2017) report delayed activation of the DMN (i.e., evoked responses in latter clusters). The authors suggest that this indicates a late influence of the DMN on the SN and FPN (i.e., after these networks have completed early/local processing). They also suggest that the delay in DMN influence may expedite the integration of information processing, albeit over time. While the authors attribute this delayed DMN activation to multisynaptic processing, and imply its role as an "influencer" acting after early-level processing (SN and FPN), the interpretation of delayed DMN activation appears to be more challenging. Shine et al. (2017) report that most of their DMN electrodes were located in the medial temporal lobe regions for clinical reasons. Here the polysynaptic hypothesis, as suggested by Shine et al. (2017), is well supported, as the medial temporal lobe is a convergent zone of cortical processing and receives input from all sensory modalities creating multiple subsystems (Simons and Spiers, 2003; Squire et al., 2004). This may create a bottleneck effect contributing to longer latencies. Nevertheless, there are other explanations that warrant consideration. For example, there is evidence of reduced DMN recruitment and alterations in hippocampal networks associated with epilepsy, which may also have contributed to the delayed DMN activation (James et al., 2013). Another study-specific limitation, clearly unavoidable, is the somewhat rostral placement of stimulation electrodes, which may have biased recordings toward delayed activation. Furthermore, it has been suggested that the DMN plays a role in functional integration of information across segregated brain regions (Hagmann et al., 2008). This proposed role could explain the delayed, and more importantly, sustained activation. Can the delayed yet sustained processing within the DMN be the brain's intrinsic effort to reorganize and bind information in a relational pattern to facilitate later retrieval? The absence of DMN activation in the early clusters, its large spatial span across the cortex, and its estimated 20% consumption of neural resources may hint toward a global, more robust role in information processing. However, this remains to be explored.

A few considerations suggest that the findings of Shine et al. (2017) should be interpreted with caution. One of the most evident limitations of interpreting CCEP

data from epileptic participants is the lack of generalizability to control populations (Kunieda et al., 2015). Although the stimulations do not trigger a recordable epileptic discharge, there may be unexplored alterations in brain activity with little to no consequences for behavior. Also, while it has been hypothesized that intracranial electrode stimulation elicits activation of the deep (layer IV–VI) cortical layers (Keller et al., 2014), the lack of such validation reduces interpretability and may require methodological clarification. Despite these limitations, CCEP studies offer an unprecedented advantage to advance the understanding of neural organization patterns, albeit in a pathological state.

To reiterate, the results from Shine et al. (2017) support the notion that, despite obvious heterogeneity within the connectome, the macroscale interactions within and between neural networks follow a temporal hierarchy. Their evidence lends further support to the conceptual framework that the region's architectural thresholds may help optimize signal propagation and foster an environment of efficient connectivity. Understanding this more global temporal hierarchy along with regional cytoarchitectural characteristics may provide insights about the connectome (van den Heuvel et al., 2015). CCEP is a powerful tool that can identify the time-varying "chronnectome" (Calhoun et al., 2014) and may possibly help predict behavior by increasing our understanding of the hierarchical organization underlying efficient information processing, and alterations that may manifest in pathological disease states.

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