Dynamic endophenotypes and longitudinal trajectories: capturing changing aspects of development in early psychosis

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The search for replicable markers — biological, psychological, social or their combinations — for psychiatric illnesses carries on. In psychosis, attention has been directed to at-risk and first-episode populations, given the possibility that following markers in young people will avoid the confounds of chronicity and exposure to pharmacological treatments. A variety of promising biomarkers have been evaluated, but few have passed through the rigorous screens of broader replication and demonstrated clinical utility.1,2

We argue in this editorial that such difficulties are unsurprising, given the construction of current studies and the fact that such disorders typically develop in adolescence and young adulthood. This “critical period”3 for risk and onset of major youth mental illnesses, such as psychosis, coincides with deeply interlaced neurobiological, clinical, affective and cognitive development, not to mention profound changes in social interactions and exposures. We explore the implications of this backdrop, including how it should inform both the search for biomarkers and the design of future studies.

Critical periods, neurobiology and mental health

Adolescence to early adulthood, the age range when the early course of psychotic illness is most likely to emerge, is a window in which much is dynamic. At a neurobiological level, this includes neurons and synapses, neurotransmission, trophic factors and long-term potentiation, among other phenomena.4 However, it is also a period of sexual maturation, exposure to novel social and physical environments, different forms of stress, new models and contexts for learning, and developing cognitive abilities, reasoning and affective states.5 Whether adaptive or maladaptive, changes in brain structure and function are believed to occur in response to these internal and external stimuli and demands. For example, the frequency and type of stressful life events change as children enter adolescence;6 with these changes come further alterations in neurobiology that may denote increased sensitization to such stimuli.7 If more persistently dysregulated during a critical window, the long-term set-points of neurobiological processes or pathways may become distorted. Neurodevelopment that results from interactions between endogenous neurobiological elements and exogenous environmental factors could thus profoundly alter brain processes, leading to differential trajectories and clinical outcomes.

Neurobiology, biomarkers and endophenotypes

The acknowledgement that biological, psychological and social markers are not static but may themselves be subject to critical periods of influence or change, especially in youth mental health, suggests a path forward. Rather than relying on “snapshots” of data at a single time point to differentiate groups or pathology, it draws attention to the dynamic trajectories of individual markers and how they change. As individuals pass through periods of heightened risk, sequential evaluation of 1 or more markers could bring to light differential courses or inflection points that relate to specific clinical or functional outcomes.

The suggestion that neurodevelopmental alterations may cut across traditional psychiatric diagnostic schemas (such as DSM categories)8 dovetails with major contemporary neuroscience initiatives. In response to the perceived failure of symptom-based classification, the National Institute of Mental Health Research Domain Criteria (RDoC) were pioneered to “reimagine” classification systems agnostic of traditional diagnostic schema.9 To advance the study and treatment of mental illness, RDoC advocates argue that we must identify and focus on valence systems, cognition, social processing, and arousal and regulatory systems across multiple levels of analysis — from genes and molecules to circuitry and (ultimately) behaviour. Similarly, the endophenotype concept has been articulated as a means of deconstructing unseen but measurable components “along the pathway” between genotype and syndromic illness.10 Endophenotypes are heritable traits that cosegregate with illness, but similar to RDoC categories can be

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Structural neuroimaging

The increasing use of longitudinal and repeated neuroimaging measures has allowed for the documentation of progressive brain changes in individuals with schizophrenia, reflecting the dynamic central nervous system. In a more homogeneous sample of patients with first-episode schizophrenia, Andreasen and colleagues found progressive grey and white matter loss in multiple regions over time related to the severity of psychotic symptoms, relapse duration and antispsychotic treatment intensity. The presence and persistence of negative symptoms in patients with schizophrenia and related psychoses is also associated with abnormalities in cortical thickness and white matter, cognitive deficits, and poor functional and vocational outcomes. Studies examining how putative endophenotypes interact and lead to differential effects on brain structure and functioning are therefore much needed. For example, in individuals at clinical high risk for psychosis, both negative symptoms and progressive cortical changes are frequently seen. This approach could derive clinical utility from imaging findings by enabling the early identification and targeted clinical follow-up of those likely to experience less favourable outcomes.

Advanced imaging techniques

Continuing innovations in neuroimaging have generated image processing techniques that capture salient alterations in brain structure at the group level and statistical techniques that are robust to uneven sampling among study participants. Longitudinal designs can thus improve statistical power for detecting group differences in neuroanatomical trajectories, allowing for the investigation of novel endophenotypes. For example, shape metrics are heritable and may have alterations specific to schizophrenia. Such methodologies provide insight into subcortical structures (such as the striatum) that are especially subject to confounding effects, or where standard volumetric comparisons fail to elucidate group differences. These metrics are also more sensitive to the progressive changes in neuroanatomy that characterize adolescent maturation. Advanced statistical, graph theory and community detection techniques are beginning to be used to infer underlying connectivity between brain regions, with adaptations of these methods to understand brain maturation. Future work will need to connect these different levels of observation into an integrated model. Advanced imaging techniques could provide novel insight into a variety of microscopic-level alterations in neuroanatomy — valuable indices for the study of progressive brain changes underlying the onset of psychosis.

Sleep alterations

Among the most profound changes occurring during adolescence is a diminution in the amount of deep sleep: Δ (1–4 Hz) electroencephalography power in nonrapid eye movement sleep declines massively during adolescence. It was on this basis that Feinberg originally advanced the neurodevelopmental hypothesis of schizophrenia, postulating that such alterations in sleep architecture correlate with synaptic pruning and maturation. Important changes in sleep patterns also take place during adolescence, notably a physiologic phase delay in the circadian rhythm. A dynamic defect in these maturational processes could be tracked over time and may offer another window to explore markers of risk for the emergence of psychotic symptoms. This is particularly relevant given emerging literature that sleep disorders are present in at-risk and first-episode phases of psychosis and are often early warning signs that precede the onset of illness or indicate relapse.

Stress reactivity

The hypothalamic–pituitary–adrenal axis harbours key markers of stress response and undergoes significant change during and after puberty. In psychosis, stress reactivity has been posited as a putative endophenotype owing to its cosegregation with illness and potential trait status in unaffected relatives. Well-designed laboratory-based paradigms have demonstrated short-term blunting of the dynamic cortisol response to stress, which may reflect exhaustion of the potential for a (healthy) robust phasic response to stress alongside the more tonic elevation found in at-risk populations. However, key questions remain about whether blunted stress response is a trait consistently found in unaffected family members and how the observed association between low tonic and robust phasic activity in healthy individuals changes along with longitudinal outcomes.

Challenges and conclusions

The examples we have discussed illustrate how putative endophenotypes could change dynamically over time — a phenomenon that would benefit from longitudinal studies. Such an approach presents 2 kinds of challenges. The first of these is concerned with theory and science: markers may diverge or change at different time points and rates; given
this background variation, how and at what thresholds should risk for a particular trajectory be flagged or treated? Will change be expressed as linear, step-wise, cyclical, or in other forms? How can baseline states be indexed and/or maladaptive changes be determined in the context of already changing neurobiology — by age, phase of sexual matura­tion, clinical stage or otherwise? A necessary next step here is for within-subject changes in biomarkers to be correlated with longitudinal clinical/functional observations, making it plausible to weed out spurious correlations or to derive the presumably heterogeneous pathways to illness or resolution.

A second set of challenges relates to the clinical applicability of this approach. For individuals in early stages of mental distress or subclinical illness, it is likely that single biomarkers/ endophenotype trajectories will present as abnormal while an overall panel will remain ambiguous. Such a scenario should trigger closer monitoring and clinical correlation over time to either worsening or resolution of biomarkers and outcomes. In later (clinical high-risk or first-episode) stages, biomarker panels may become increasingly abnormal, prompting more intensive interventions with different risk/benefit profiles. It will therefore be important to identify and follow overlapping cohorts with varying degrees of enrichment for risk in order to understand whether putative endophenotypes better align with a priori risk groupings or evolving clinical/functional outcomes.

Finally, are youth mental health services capable of handling an integrative and coordinated approach given current resource constraints? At the research stage, intensive assaying at frequent time points is necessary. Ultimately, however, we envision the use of biomarkers not as a standard tool to be applied uniformly at great cost, but as a measured and targeted approach that can be selected to guide risk assessments or treatment decisions at key branch-points. And the economic arguments to fund “pre-emptive” clinical infrastructure are beginning to be made. Recent developments aimed at transforming youth mental health systems, including in Canada, will almost certainly involve at-risk populations beyond DSM-specific diagnoses. They now promise to inform services ready to take up knowledge about dynamic endophenotypes can be situated.

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