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Mapping the pathophysiology of schizophrenia: interactions between multiple cellular pathways

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Abstract
Schizophrenia is a complex disorder involving dysregulation of multiple pathways in its pathophysiology. Dopaminergic, glutamatergic and GABAergic neurotransmitter systems are affected in schizophrenia and interactions between these receptors contribute to the pathophysiology of the disease. Deficits in acetylcholine muscarinic receptors have been identified in a sub-group of individuals with schizophrenia. Inflammation has also been found to play a major role in the development and exacerbation of psychotic symptoms in schizophrenia. Additionally, evidence from genetic, post-mortem and animal studies over the past decade has identified a number of susceptibility factors for schizophrenia, including neuregulin 1 (Nrg1) and its receptor ErbB4, disrupted-in-schizophrenia-1 (DISK1), dysbindin-1, catechol-O-methyl tranferase (COMT), BDNF, and Akt. These factors and related pathways interact closely with dopaminergic, glutamatergic and GABAergic neurotransmitter systems. A key question is how do these interactions contribute to the pathophysiology of schizophrenia? More specifically, how do these components interact during early brain development based on the view of schizophrenia as a developmental disorder? Therefore, this special issue aims to map the pathophysiology of schizophrenia by illuminating the interactive nature of specific pathways on different levels of the brain from cellular pathways and neural circuits to functional deficits.

Keywords
between, multiple, cellular, pathways, interactions, schizophrenia, mapping, pathophysiology

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Mapping the pathophysiology of schizophrenia: interactions between multiple cellular pathways

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Schizophrenia is a complex disorder involving dysregulation of multiple pathways in its pathophysiology with strong evidence to support roles for dopaminergic, glutamatergic, GABAergic, and cholinergic neurotransmitter systems and their interactions in the pathophysiology of the disorder (Benes, 2009; Karam et al., 2010; Gibbons et al., 2013). Additionally, evidence from genetic, post-mortem and animal studies over the past decade has identified a number of susceptibility factors for schizophrenia, including neuregulin 1 (Nrg1) and its receptor ErbB4, disrupted-in-schizophrenia-1 (DISK1), catechol-O-methyl transferase (COMT), BDNF, and Akt, along with their related pathways, that interact closely with dopaminergic, glutamatergic, and GABAergic neurotransmitter systems (Karam et al., 2010). Hence a key question is how these neurotransmitter systems and their interactions contribute to the pathophysiology of schizophrenia and whether interactive changes in these pathways occur in early brain development, based on the view of schizophrenia as a developmental disorder?

This Frontier Research Topic has brought together leading experts in the field to address these questions from different angles in nine reviews, one theoretic article and two research articles. The first four articles focus on the roles and interactions of neurotransmitters in the pathophysiology of psychiatric disorders. Snyder and Gao (2013) provide an excellent review of NMDA receptor hypofunction hypothesis, suggesting NMDA receptor hypofunction as a convergence point for progression of schizophrenia, including neuregulin 1 (Nrg1) and its receptor ErbB4, disrupted-in-schizophrenia-1 (DISK1), catechol-O-methyl transferase (COMT), BDNF, and Akt, along with their related pathways, that interact closely with dopaminergic, glutamatergic, and GABAergic neurotransmitter systems (Karam et al., 2010). Hence a key question is how these neurotransmitter systems and their interactions contribute to the pathophysiology of schizophrenia and whether interactive changes in these pathways occur in early brain development, based on the view of schizophrenia as a developmental disorder?

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In the following three papers, Chana et al. have provided an excellent review of the current progress in the search for biomarkers for schizophrenia and psychosis, focusing on biomarkers for major neurotransmitter systems in post-mortem brain studies, as well as covering some recent and exciting studies in microRNA dysregulation in both the blood and brain of schizophrenia patients (Chana et al., 2013). It is followed by a very timely review of current knowledge on the features of polysialic acid and their synthesizing enzymes (specially ST8SIA2), their functions in regulations of cell adhesion, ion channels, neurotrophins (such as BDNF) and catecholamine neurotransmitters (particularly dopamine), in light of several recent lines of evidence linking polysialic acid to schizophrenia (Sato and Kitajima, 2013). Iwakura and Nawa provide a very clear overview of the ErbB1-dependent EGF/neuregulin signals, their role in regulating the development and function of the central nervous system, and the contribution of deficits in ErbB signaling to schizophrenia and neurological disease (Iwakura and Nawa, 2013).

Four articles discuss the animal model of schizophrenia, two of which address the Nrg1-cannabinoid interaction in a hypomorphic Nrg1 (Nrg1 HET) mouse model of schizophrenia. In an original study, Spencer et al. investigated Nrg1-cannabinoid interaction in the hippocampus using proteomics, in which they identified alterations of some proteins involved in vesicular release of neurotransmitters, serotonergic neurotransmission, and growth factor release in response to Nrg1 hypomorphism and...
of schizophrenia. We believe that this Frontier Research Topic will stimulate the development of future collaborative and interdisciplinary research to reveal the unknown mechanisms underlying the pathophysiology of schizophrenia.

REFERENCES


Kesby, J. P., Cui, X., Burne, T. H. J., and Eyles, D. W. (2013). Altered dopamine ontology in the developmentally vitamin D deficient rat model and its relevance to schizophrenia were reviewed (Kesby et al., 2013). This review suggests that early alterations in dopamine ontology are a core feature in the pathophysiology of schizophrenia representing a critical aspect useful to a model of this disease.

Finally, in the context of schizophrenia as a neurodevelopmental disorder, Catts et al. (2013) discuss elegantly the normal development of the prefrontal cortex on the molecular and cellular levels in line with cognitive development, as well as the timing of cognitive decline in schizophrenia. They proposed to reconsider schizophrenia as an outcome from a failure to reach the final state of cortical maturation resulting in retention of an immature cortex rather than resulting from an excess of adolescent synaptic pruning (Catts et al., 2013).

In summary, these studies illustrate clearly the interactive nature of specific pathways on different levels of the brain from molecular and cellular pathways, and neural circuits to functional deficits contributing to the pathophysiology of schizophrenia. We believe that this Frontier Research Topic will stimulate the development of future collaborative and interdisciplinary research to reveal the unknown mechanisms underlying the pathophysiology of schizophrenia.

Nrg1-cannabinoid interaction (Spencer et al., 2013). In a short review, Karl and Arnold further discussed the complex neuro-behavioral effects of Nrg1-cannabinoid interaction and its clinical implications (Karl and Arnold, 2013). In a BDNF heterozygous mouse model, Manning and van den Buuse investigated the effects of chronic methamphetamine treatment during late adolescence/early adulthood on a behavioral endophenotype related to the positive symptoms of schizophrenia, prepulse inhibition (PPI) of the acoustic startle reflex (Manning and van den Buuse, 2013). In the fourth animal model paper, altered dopamine ontology in the developmentally vitamin D deficient rat model and its relevance to schizophrenia were reviewed (Kesby et al., 2013). This review suggests that early alterations in dopamine ontology are a core feature in the pathophysiology of schizophrenia representing a critical aspect useful to a model of this disease.

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