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Glial cells in schizophrenia: A unified hypothesis

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Abstract

The cellular neurobiology of schizophrenia remains poorly understood. We discuss neuroimaging studies, pathological findings, and experimental work supporting the idea that glial cells might contribute to the development of schizophrenia. Experimental studies suggest that abnormalities in the differentiation competence of glial progenitor cells lead to failure in the morphological and functional maturation of oligodendrocytes and astrocytes. We propose that immune activation of microglial cells during development, superimposed upon genetic risk factors, could contribute to defective differentiation competence of glial progenitor cells. The resulting hypomyelination and disrupted white matter integrity might contribute to transmission desynchronisation and dysconnectivity, whereas the failure of astrocytic differentiation results in abnormal glial coverage and support of synapses. The delayed and deficient maturation of astrocytes might, in parallel, lead to disruption of glutamatergic, potassium, and neuromodulatory homeostasis, resulting in dysregulated synaptic transmission. By highlighting a role for glial cells in schizophrenia, these studies potentially point to new mechanisms for disease modification.

Introduction

Schizophrenia is a chronic and debilitating psychiatric disorder, with accelerated mortality and profound morbidity¹. As a syndrome, schizophrenia progresses through at least three phases: the prodromal (prepsychotic) phase, the initial onset of psychosis, and chronic illness². The core symptomology of schizophrenia includes social, emotional, perceptual, and cognitive domains, and its clinical phenotype can be subdivided into positive and negative symptoms, and those of cognitive impairment². In addition, most patients have abnormal sleep patterns, and sleep disturbances tend to precede clinical onset³. Cognitive impairments are apparent early in the prodromal phase, and can be used to follow disease progression and therapeutic outcomes in patients⁴. An established core feature of cognitive impairment in schizophrenia is diminished working memory, which can occur in patients who are drug naive or medicated, and in both early and chronic phases of the disease⁵. The

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dorsolateral prefrontal cortex is a primary site involved in working memory, and its functional impairment parallels the cognitive decline in schizophrenia⁶. Patients with schizophrenia have altered neural oscillatory patterns in the dorsolateral prefrontal cortex when performing working memory tasks⁷, and abnormal neural oscillations occur in other cortical areas throughout all clinical phases of schizophrenia, independent of disease phenotype⁷. This observation has led to the hypothesis that oscillatory abnormalities are a core endophenotype of schizophrenia, which suggests that broad cortical dysconnectivity underpins the symptomology of this disorder⁷.

Schizophrenia has a substantial genetic component, but its penetrance, phenotype, and severity are environmentally modulated⁸; its clinical manifestations comprise a final common pathway of multiple genetic and epigenetic influences⁸. Twin studies have established a strong genetic basis for schizophrenia, with concordance rates estimated at 33% for monozygotic twin pairs, compared with 7% in dizygotic twin pairs⁹. In adoption studies, children with a high genetic risk of developing schizophrenia are more sensitive to adverse rearing than children at lower genetic risk¹⁰. The genetic susceptibility for schizophrenia is therefore sensitive to environmental and social risk factors, such that a high prevalence of adverse developmental events increases the risk of schizophrenia later in life⁸. Developmental risk factors include physical or emotional maternal stress during the first trimester of pregnancy, maternal infections during the second and third trimesters, obstetric complications, and exposure to various infectious agents and their attendant inflammatory responses¹¹. To what extent any single factor increases susceptibility to schizophrenia remains unclear^{8,11}.

Disease phenotype and the associated genetic, histopathological, and imaging biomarkers vary greatly across patients, as well as by brain region and even in between cortical layers¹²⁻¹⁷. Complicating matters further, schizophrenia research is highly diverse, and the heterogeneity in study designs has led to a difficulty in controlling and standardising variables, such as the brain regions studied, the rate of disease progression, drug treatment history, lifestyle factors, and cause of death^{12-15,18,19}.

White matter pathology in schizophrenia

At the onset of psychosis, the anatomic pathology of schizophrenia includes structural changes in both grey and white matter across multiple brain regions, with the most severe changes being in forebrain tract connectivity^{18,20}. White matter loss can precede obvious clinical symptoms in patients who are drug naive²⁰, and progresses throughout the course of clinical manifestation²⁰. The structural environment of axons can be assessed using diffusion tensor imaging, which is a technique based on MRI that assesses changes in white matter fractional anisotropy—a measure of molecular water diffusibility²¹. In patients with schizophrenia, abnormal fractional anisotropy appears in multiple subcortical tracts, suggesting reduced white matter tract integrity. Much of the forebrain subcortical white matter is affected, including the cingulum, as is the white matter underlying the parietal regions, occipital regions, and prefrontal cortex²¹. Fractional anisotropy reductions are especially evident in the white matter below the cingulum and prefrontal cortex, and across the entire superior longitudinal fasciculus, which is the principal frontoparietal white matter

connection²¹, as well as in the arcuate and uncinate fasciculi, which connect limbic regions²¹.

The frontoparietal circuitry is particularly associated with working memory⁶. Using a combination of diffusion tensor imaging and functional MRI during a memory task, a direct correlation was noted between reduced fractional anisotropy in the right medial temporal and right frontal lobes and relative hypoactivation in the prefrontal, superior parietal, and occipital regions of patients with schizophrenia (with hypoactivation detected as a diminished blood-oxygen-level dependent signal)²². Thus, deficits in frontoparietal network coordination might underlie the working memory deficits often observed in patients with schizophrenia²³.

Abnormalities in fractional anisotropy are associated with other typical symptoms of schizophrenia, including auditory hallucinations²³, the passivity of abulia, and impaired executive functions²⁴. Furthermore, the extent of white matter abnormalities is associated with the severity of symptoms in schizophrenia; in patients with first-episode schizophrenia who are drug naive, those patients with more widespread white matter abnormalities show more severe negative symptoms than those with less widely distributed abnormalities²⁵. This finding is consistent with observations that the internal capsule is smaller in patients with schizophrenia who have a poor outcome than in those who have a good outcome²⁶. Chronic administration of the dopamine, serotonin, norepinephrine antagonist clozapine leads to increased fractional anisotropy values in several brain regions, suggesting that clozapine treatment can restore white matter microstructural integrity²⁷, with attendant symptomatic improvement.

Symptoms of psychosis are observed in other diseases affecting white matter integrity, including the leukodystrophies, demyelinating disorders, infiltrating glial neoplasms, and callosal anomalies, among others²⁸. These diverse diseases share a disruption of normal myelin development or maintenance, and psychosis is more severe in diseases with nonfunctional myelin formation than in those with disrupted myelinated structures²⁸. In addition, the concentration of N-acetylaspartate is raised in the white matter of patients with schizophrenia, suggesting both myelin loss and axonal abnormalities²⁹. Together, these data demonstrate an association between white matter changes, myelin loss, and altered circuit activation in distal regions²², thereby linking intracortical dysconnectivity with the core features of schizophrenia. An open question is whether defective oligodendrocytic function is primary or secondary to dysfunctional neural circuits in schizophrenia. We will argue that white matter pathology is a common feature of schizophrenia, and that it results from a primary, cell-intrinsic failure in the differentiation of glial progenitor cells.

Glial dysfunction in schizophrenia

The hypothesis that higher brain function solely comprises the integrated product of neuronal activity has been challenged by evidence showing the importance of glial cells to both the development and structure of local neural networks. Both macroglia—including oligodendrocytes³⁰, glial progenitor cells³¹, and astrocytes³²—and microglial cells³³ contribute to the modulation and synchronization of neuronal activity. As such, macroglia

and microglial cells influence information processing and the structural and functional plasticity of neural networks. Together, these functional roles, the evolution of astrocyte complexity with phylogeny, and the coincident appearance of neuropsychiatric disorders with hominid evolution all combine to suggest the strong contribution of glial pathology to psychiatric diseases^{34,35}.

Although glial involvement in schizophrenia has long been considered^{12,14,15,18,19,34-36}, the nature of its contribution to schizophrenia pathology has remained unclear. All four glial cell types have been reported as showing dysregulated patterns of gene expression in postmortem tissue from patients with schizophrenia, in both cortical and subcortical regions³⁵. The involvement of glial cells in schizophrenia is supported by RNA sequencing data^{37,38} and genome-wide association studies^{39,40}. These data point to the enrichment of genes related to development in schizophrenia risk loci^{37,40}, and in particular suggest the dysregulation of neuroinflammatory pathways and upregulation of genes related to astrocytes in schizophrenia^{38,39}. RNA expression data suggested a predominant neuronal pathogenesis of schizophrenia⁴¹, but that particular study was limited to postmortem analysis of adult brain tissue. Limiting data to postmortem analysis of adult brain tissue is insensitive to the transcriptional dysregulation of glial development, which might have profound consequences for adult neural network structure⁴². Indeed, substantial glial developmental pathology and consequent neural network dysfunction might leave only a weak RNA expression signature in adult glial cells. As such, defining the extent to which glial pathology in schizophrenia is causal or merely secondary to neuronal pathology has been difficult to establish.

Fundamentally, an absence of adequate experimental tools³² and the evolutionary gap between rodent and human astrocytes⁴³ have underlain the difficulties in assessing the relative causal contribution of glial pathology to schizophrenia. To address this question, Windrem and colleagues⁴² developed a humanised chimeric mouse model by engrafting glial progenitor cells—prepared from induced pluripotent stem cells derived from patients with juvenile-onset schizophrenia or their age-matched controls—into neonatal, congenitally hypomyelinated mice. The human glial progenitor cells outcompeted their host counterparts to colonise the recipient mouse brains, where they differentiated into myelinating oligodendrocytes and astrocytes; the resultant mice developed patient-specific, largely humanised forebrain white matter⁴². The schizophrenia-derived human glial chimeras manifested delayed and deficient glial differentiation, which was associated with diminished oligodendrocyte maturation and impaired central myelination⁴². The structural maturation of astrocytes was likewise impaired, and showed fewer primary processes, less proximal branching, and less coherent domain structure than did littermate controls⁴². These schizophrenia-derived glial chimeric mice displayed higher anxiety levels, diminished auditory prepulse inhibition, anhedonia, social deficits, impairment in executive memory, hyperactivity, and disrupted sleep compared with littermate controls⁴². These signs are all consistent with the hypothesis that a deficit in glial functions contributes to the behavioural phenotype of these mice. Furthermore, RNA sequencing of glial progenitor cells derived from patients with schizophrenia—the parental source of both oligodendrocytes and astrocytes—revealed substantially downregulated expression of a host of genes associated with glial differentiation relative to normal glia⁴². The defective differentiation of glial

progenitor cells was linked to the impairment of potassium uptake, which was rescued by the shRNAi-mediated knock-down of the REST repressor, binding sites that were noted in the regulatory regions of multiple glial potassium transporters. These data indicated that targeting REST-dependent transcriptional repression might rescue astroglial differentiation and potassium transport in schizophrenia⁴⁴. This group of experimental observations suggests that glial progenitor cells contribute to schizophrenia, or at least to juvenile-onset disease in an in vivo, humanised glial chimeric mouse devoid of pre-existing neuronal pathology. In the following sections, we discuss each of the four types of glial cells in terms of their function and their connection to schizophrenia.

Oligodendrocytes

Myelination is a dynamic process that adapts to external stimuli, conforming to the activity-dependent requirements of the maturing neural network long after early ontogeny. Acquiring new skills is associated with an increase in myelination⁴⁵, whereas sensory deprivation⁴⁶ and maternal rearing of mice results in hypomyelination⁴⁷, including reduced myelin thickness and gene expression in the prefrontal cortex⁴⁶, and irreversible myelination alterations in the medial prefrontal cortex⁴⁷. The medial prefrontal cortex alterations are accompanied by clinically significant social and cognitive deficits, indicating that proper myelination of the prefrontal cortex is central to cognition, and that myelination dependent on social experience can regulate cognitive functions⁴⁷. Furthermore, myelinated and unmyelinated segments occur on the same axon in the neocortex in mice³⁰, and linear myelin ensheathment varies between cortical layers of the same region⁴⁸. The internodal distances of single myelin segments can also vary substantially, with major implications for conduction properties and neuronal synchrony within distributed neural networks⁴⁹. Such complexity in myelination predisposes it to the types of desynchronising events that are characteristics of schizophrenia.

Oligodendrocytes are the principal myelinating cell type of the CNS⁵⁰. Myelination might depend in part upon neuronal activity, through activation of oligodendrocytic NMDA receptors⁵⁰. The presence of activity-dependent growth factors (eg, neuregulin) accelerates myelination, and neuronal activity is therefore a key determinant of myelinogenesis. Accordingly, myelination is reduced by neuronal NMDA receptor inhibition and by suppression of action potentials⁵⁰. In mice reared alone, mRNA levels of neuregulin 1 are lower than in mice reared together in groups, suggesting a link between social isolation and central hypomyelination⁴⁷. However, knocking out oligodendrocyte-specific *ErbB3*, a growth factor receptor, produces a phenotype only when done during early postnatal development⁴⁷; similarly, developmental overexpression of neuregulin 1 leads to hypermyelination⁵¹. These observations suggest that a crucial time window exists during ontogeny, during which oligodendrocytes are particularly sensitive to changes in the neural environment and developmental myelination is most at risk of disruption.

In patients with schizophrenia, postmortem stereological analyses have shown a substantial decrease in the numbers and density of oligodendrocytes in cortical layers III⁵² and V16 of the dorsolateral and medial prefrontal cortices, and in the superior frontal gyrus¹⁶. Other postmortem findings in the brains of patients with schizophrenia include a substantial

decrease in both volume and mitochondrial number of oligodendrocytes within the caudate nucleus and prefrontal areas⁵³. Similarly, in patients with chronic schizophrenia treated with antipsychotics, both oligodendrocyte numbers and myelin volume were significantly diminished in the anterior thalamic nucleus⁵⁴. Other postmortem studies have documented differentially diminished mRNA expression of several proteins associated with oligodendrocytes and myelin (eg, QKI⁵⁵ and CNP⁵⁶) in patients with schizophrenia as compared with healthy controls. This decline in mRNA expression of oligodendrocyte markers might reflect a developmental failure in oligodendrocyte maturation, as has been shown in oligodendrocyte progenitor cells derived from patients^{42,44}. Taken together, these studies indicate that schizophrenia is associated with impaired oligodendrocyte differentiation and consequent dysmyelination.

Glial cell progenitors

In the adult CNS, myelinating oligodendrocytes can be replenished from a pool of glial progenitor cells⁵⁷. Glial progenitor cells represent the majority of proliferating cells in the adult CNS; most persist at the G1 phase of the cell cycle⁵⁸. The transition of glial progenitor cells into the S phase is positively regulated by cyclins D and E, and negatively regulated by members of the Kip family of cell cycle inhibitors⁵⁹. Both adult and fetal glial progenitor cells retain the capacity to differentiate into oligodendrocytes and astrocytes (figure),³¹ following a tightly regulated differentiation pathway³¹. Oligodendrocyte lineage transcription factor 2 (*Olig2*) is required for glial progenitor cells to initiate the oligodendrocytic differentiation programme⁶⁰, while simultaneously inhibiting astrocytic differentiation, as signalled by the transcription of astrocytic glial fibrillary acidic protein³¹. Overexpression of *Olig2* in adult glial progenitor cells in mice enhances differentiation and remyelination, while overexpression in embryonic glial progenitor cells boosts the generation of oligodendrocytes, which is accompanied by an increase in glial progenitor cell migration velocity⁶⁰. In contrast, deletion of *Olig2* in glial progenitor cells of adult mice potentiates their astrocytic differentiation, significantly reducing the number of oligodendrocytes and thereby suppressing myelination³¹. Other elements crucial in determining the fate of glial progenitor cells include growth factors, neurotransmitters, redox state³¹, and neural activity^{31,61}. The developmental fate of glial progenitor cells is therefore highly sensitive to the surrounding neural environment^{31,61}.

In postmortem brain samples from patients with chronic and severe forms of schizophrenia, the expression of the genes encoding cyclins D1 and D2 is increased, whereas expression of those genes encoding the Kip cycle cell inhibitors is decreased³⁶. In patients with less severe symptoms, expression of genes associated with maintenance of the cell cycle in glial progenitor cells is increased, whereas expression of genes associated with cell cycle arrest is decreased¹⁴. Thus, glial progenitor cells tend to remain proliferative, and are less likely to exit the cell cycle in patients with schizophrenia. In genetically unrelated patients with schizophrenia, genotyping and polymorphism detection have revealed that an interaction between *OLIG2* and *ERBB4*, and an interaction between *OLIG2* and *CNP* both affect disease risk⁶². A linkage analysis for trans-effects on the expression of these two pairs of genes suggests that each locus regulates the expression of the other⁶². Furthermore, diffusion tensor imaging studies have shown an association of a single nucleotide polymorphism in

OLIG2 with decreased white matter integrity⁶³, implicating impaired glial progenitor cell differentiation in white matter pathology. These findings collectively suggest that the cell cycle of glial progenitor cells is dysregulated in patients with schizophrenia, resulting in arrested differentiation and delayed maturation of oligodendrocytes and astrocytes.

Astrocytes

Astrocytes play several key roles in the CNS during development and in adulthood. During development, they are essential for synapse formation⁶⁴. Astrocytes also produce and deliver glutamate to neurons⁶⁵, buffer extra-cellular potassium⁶⁶, and respond to neuromodulator activity³². Inhibition of the $\alpha 1$ noradrenaline receptor eliminates more than 90% of spontaneous astrocytic calcium signalling in awake behaving mice³². In turn, astrocytic calcium signalling modulates potassium buffering, glycogen mobilisation, and functional hyperaemia³². Astrocytes are therefore crucial to the regulation of multiple homeostatic functions that determine the precision of synaptic activity. They not only provide metabolic support for synapses, but are also necessary for synapse formation and maintenance during development and adulthood⁶⁴.

In rodent models, changes in astrocytic numbers and morphological phenotype have been shown to trigger cognitive dysfunction. Selective elimination of astrocytes using a toxin specific to astrocytes⁶⁷, or in a transgenic mouse line,⁶⁸ reduces the density of astrocytes in the prefrontal cortex, resulting in mice with deficits in attentional set-shifting, working memory, reversal learning⁶⁷, and recognition memory, and abnormal cortical gamma oscillations⁶⁸. Furthermore, mice overexpressing S100-b (a calcium-binding protein, which is important for the migration, maintenance, and morphology of mature astrocytes) show altered patterns of spatial and temporal exploration, suggesting an impairment in short term memory⁶⁹. In contrast, S100-b deletion enhances longterm potentiation in the CA1 hippocampal region, accompanied by improved spatial memory and fear-associated memory⁷⁰. Similarly, diminished expression of the astrocytic glutamate transporter reduces prepulse inhibition of the acoustic startle response⁷¹, a well established feature of schizophrenia.

Postmortem data regarding disease-dependent changes in protoplasmic and fibrous astrocytes vary greatly. In one study of the anterior cingulate gyrus of patients with schizophrenia, significant abnormalities in various astrocytic markers (DIO2, AQP-4, S100-B, glutaminase, thrombospondin, and excitatory amino acid transporter 2) were observed in the deep cortical layers, as compared with those of healthy individuals¹³. No corresponding differences were found in the superficial layers or in the underlying white matter of the anterior cingulate gyrus¹³, suggesting that astrocytic pathology in those cases was region specific, with a regional selectivity that predicted symptomatic heterogeneity. In another study, fewer fibrous astrocytes were observed in the subgenual cingulate cortex of patients with schizophrenia than in healthy controls⁷². This finding was of particular interest since other white matter diseases associated with psychosis, such as some leukodystrophies, have been associated with abnormalities in fibrous astrocytes⁷³, suggesting a possible connection between thought disorders, white matter alterations, and astrocyte pathology.

We propose that failure of glial differentiation is an initial event in schizophrenia, whereas astrogliosis and microgliosis might occur as secondary events later in disease development, possibly triggered by disturbances in network connectivity, homeostatic failure of glutamate and potassium regulation, changes in neuromodulator release, neuroinflammation, sleep deficits, and drug-related effects (figure). The effects associated with drugs might explain the large variation in postmortem data from patients with schizophrenia, as reactive gliosis is not a prerequisite for developing schizophrenia. Thus, the focus on reactive glia, rather than on abnormal glial developmental trajectories, might constitute a major obstacle in determining the role of glial cells in schizophrenia.

The contribution of glutamate to the pathology of schizophrenia has been studied extensively. Postmortem analyses suggest layer-specific alterations in both glutamatergic neurons and receptors, as well as in the glutamate metabolic pathway⁶⁵. These include reductions that are associated with schizophrenia in the expression of the astrocytic glutamate transporter⁶⁵, glutamine synthetase⁶⁵, glutamate dehydrogenase⁷⁴, glutaminase⁶⁵, and D-serine⁷⁵. However, the extent to which these changes in glutamate homeostasis occur autonomously in neurons, rather than being downstream of astrocytic failure in glutamate reuptake and processing, remains unclear. S100-B expression is increased in astrogliosis, in neurodegenerative and neuroinflammatory diseases⁷⁶, and in the glial cells of patients with schizophrenia⁷⁷. When compared with healthy individuals, patients with schizophrenia who have frequent hallucinations have more abundant glial cells containing S100-B in the grey matter of the dorsolateral prefrontal cortex, whereas other studies found significantly fewer astrocytes in patients with predominantly negative symptoms⁷⁷. In sum, these data suggest an association between astrocyte pathology, glutamatergic signalling, and schizophrenia.

Astrocytes might regulate oligodendrocyte function, whether by directly regulating oligodendrocyte differentiation, or by indirectly affecting myelination competence, or both⁷⁸. During development, the ATP-sensitive inward rectifier potassium channel (Kir4.1) is regulated in both astrocytes and oligodendrocytes, resulting in distinct potassium influxes at embryonic and postnatal stages in the oligodendrocyte cell lineage⁷⁹. Kir4.1 is an important regulator of oligodendrocyte differentiation⁸⁰ and myelinating function⁷⁹. Kir4.1 regulation of potassium influxes influences the thickness and extent of the myelin sheath according to the maturation stage of oligodendrocytes⁸¹. Oligodendrocytes cultured from Kir4.1 knockout mice display an immature morphology and a depolarised membrane potential⁷⁹, and these mice show pronounced pathology in the entire CNS, and die before reaching adulthood⁷⁹. Immunohistochemical studies of rat white matter have shown Kir4.1 immunoreactivity on the fine processes of astrocytes, beginning at postnatal day 10 (P10) and peaking at P15⁸⁰. These changes in potassium channel expression correspond to developmental shifts in resting membrane potential, whereby the resting membrane potential declines as glial progenitor cells differentiate into mature oligodendrocytes⁸¹. Kir4.1 can also modulate the shape of action potentials⁸², and two papers have reported the transcriptional downregulation of *KCNJ10* (the gene encoding Kir4.1) in glial cells derived from patients with childhood-onset schizophrenia^{42,44}.

Astrocytes can affect oligodendrocyte differentiation by secreting growth factors, including platelet-derived growth factor⁸³. Decreased platelet-derived growth factor secretion results in sustained glial progenitor cell maintenance at the expense of oligodendrocyte maturation⁸³. Astrocytes also secrete the signal effectors neuregulin and brain-derived neurotrophic factor, both of which mediate neural activity-dependent myelination⁵⁰. Polymorphisms of both *NRG1* and *ERBB4* have been observed in patients with schizophrenia⁸⁴, and mice with impaired ErbB signalling show alterations in oligodendrocyte number, morphology, and myelin thickness, with correspondingly reduced conduction velocity⁸⁵, as well as schizophreniform behavioural abnormalities and cognitive impairments⁸⁶. These mice also show perturbation of mesolimbic dopaminergic function, a common feature of clinical schizophrenia⁸⁵. Together, this literature suggests that downregulation of Kir4.1 in oligodendrocytes and astrocytes leads to hypomyelination. Furthermore, astrocytes secrete growth factors that promote oligodendrocytic differentiation, a shortage of which might exacerbate oligodendrocyte pathology in schizophrenia, at least in part by altering myelination-associated plasticity.

Neuromodulator-related abnormalities are an established feature of schizophrenia pathology⁸⁷, and are likely to contribute to the sleep deficits³ and cognitive impairments^{2,4,5} of patients with schizophrenia. Evidence from two-photon imaging experiments on awake behaving mice have shown that neuromodulators—in particular noradrenaline—regulate astrocytic calcium signalling, which in turn regulates extracellular potassium concentrations during the sleep–wake cycle⁶⁶. As such, it is possible that the defects in neuromodulator signalling in schizophrenia lead at least partially to downstream abnormalities in extracellular potassium concentrations, with unpredictable consequences for synaptic transmission.

Finally, astrocytic complexity has expanded markedly with phylogenetic evolution, with both increased cell autonomous complexity and pleomorphism⁴³. Human astrocytes are distinguishable both structurally and functionally from those of other primates^{43,88}, suggesting that astrocytes are crucial to the cognitive capabilities of the human species⁴³. In humanised chimeric mice, engrafted human glial progenitor cells differentiated into astrocytes that maintained the large size and complexity of human astrocytes, and humanised chimeric mice outperform mice transplanted with mouse glial progenitor cells on all cognitive tests⁸⁹. This observation suggests that evolutionary changes in astrocytes are central to higher brain function, possibly implicating astrocytes in the pathology of schizophrenia—a disease unique to humans⁴³. The increased structural and functional complexity of human astrocytes is expected to render these highly differentiated cells more vulnerable to environmental and social risk factors, such as maternal stress and inflammation during pregnancy, which are known to increase the risk of schizophrenia^{8,11}.

Microglia

Microglia are the resident immune cells of the CNS⁹⁰, and are responsible for synaptic pruning during development³³. The role of microglial activation and neuroinflammation in schizophrenia has been a recent focus of investigation. Although some meta-analyses suggest an increase of both cellular and molecular concentrations of inflammatory markers

in the brains of patients with schizophrenia¹⁵, other analyses have found no consistent association with microglial activation^{12,17}. Genes associated with positive loci for schizophrenia in genome-wide association studies have been shown to be enriched among genes dynamically regulated in the prefrontal cortex during early brain development; this enrichment was notable given the many genes in the dorsolateral prefrontal cortex that undergo fetal-to-postnatal isoform switching during second trimester development³⁷. Thus, genetic alterations during early brain development could result in altered developmental trajectories¹⁹.

Activated microglia have been shown to attenuate the proliferation of glial progenitor cells⁹¹, and maternal infection can cause glial progenitor cell death in two waves⁹². Inflammation has been shown to increase expression of *Olig2* in mice, and inhibit the expression of other factors important in oligodendrocyte maturation, including the cell cycle factor P27Kip1, resulting in disrupted oligodendrocyte maturation⁹³. Microglial activation during embryogenesis predicts perturbed white matter integrity⁹⁴, which is particularly vulnerable to inflammatory responses early in the third trimester of gestation, during oligodendrocytic production and maturation⁹⁴. Neonatal neuroinflammation induced by intracerebral injection of lipopolysaccharide provokes white matter necrosis at P14⁹⁵. Likewise, neonatal mice treated with IL-1 beta display long lasting hypomyelination at P35, characterised by increased abundance of non-myelinated axons, reduced myelin thickness, reduced white matter fractional anisotropy, higher densities of glial progenitor cells, and cognitive deficits⁹³. In sum, these studies suggest that activated microglial cells contribute to the deficits in macroglial differentiation that are characteristic of schizophrenia.

An integrated model for glial cells in schizophrenia

On the basis of this collection of insights into the four types of glial cells, aberrant glial function might be a central element of schizophrenia pathology. We hypothesise that microglial activation in crucial periods during embryogenesis disrupts glial progenitor cell proliferation and differentiation competence, resulting in delayed oligodendrocyte and astrocyte differentiation, and aberrant maturation of those cells (figure)⁴². In this scenario, abnormalities in oligodendrocyte numbers and maturation competence lead to structural hypomyelination, which results in cortical and subcortical abnormalities in white matter integrity. Impaired astrocyte differentiation leads to glutamatergic dysfunction, abnormal potassium regulation, and dysregulation of growth factors and neuromodulators, providing a general explanation for the homeostatic disruptions reported in schizophrenia. The dysfunction of various glial cells in parallel would result in oscillatory abnormalities and dysconnectivity across the brain, and global dyssynchrony, which together account for the hallmark characteristics of schizophrenia: positive and negative symptoms, and symptoms of cognitive impairment.

Since the completion of gliogenesis occurs relatively late in development⁹⁶, infections occurring during late fetal development would be likely to have selectively greater effects on glial cell development than on resident neurons. The period of final maturation of oligodendrocytes, during which they are vulnerable to microglial activation⁹⁴, corresponds to that period in which maternal infection and stress have been most strongly associated with

schizophrenia⁹⁷. As discussed in this Review, several of the secondary events in schizophrenia—such as abnormal glutamate and potassium homeostasis, abnormalities in sleep patterns, or even drug treatment— can lead to later reactive changes in glial cells, which might be quite independent of the initial defects in glial differentiation. These secondary effects might explain the inconsistencies in postmortem analyses of glial cells in the brains of patients with schizophrenia. Importantly, schizophrenia remains a highly heterogeneous disorder in its causation, such that the proportion of schizophrenia cases in which glial cell dysfunction is causally involved is yet to be established. Nonetheless, the studies discussed here suggest that targeting glial pathology offers a new approach towards better understanding and ultimately modifying the complex pathogenesis of schizophrenia.

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Key messages

- Emerging studies show that microglial cells, oligodendrocytes, and astrocytes are all active participants in higher brain function
- Maternal infections can activate microglia cells, which in turn negatively affect the differentiation of glial progenitor cells
- Studies of humanised chimeric mice—generated using glial progenitor cells that are derived from the stem cells of patients with schizophrenia—show that glial progenitor cells fail to differentiate into oligodendrocytes and astrocytes, despite normal neuronal architecture
- We propose a model in which the failure of oligodendrocytes and astrocytes to differentiate contributes to several of the key characteristics of schizophrenia, including hypomyelination and abnormalities in glutamate and potassium homeostasis

Search strategy and selection criteria

We identified references for this review through searches of PubMed for articles and reviews published from Feb 1, 1989, to June 30, 2019, using the terms “schizophrenia”, “oligodendrocytes”, “astrocytes”, “NG2 cells”, “glial progenitor cells”, “microglial”, and “macroglial.” Relevant articles published between Feb 2, 1989, and Jun 25, 2019, were identified through searches in Google Scholar with the same terms. We reviewed the articles resulting from these searches and the relevant references cited in those articles, and included articles published in English only.

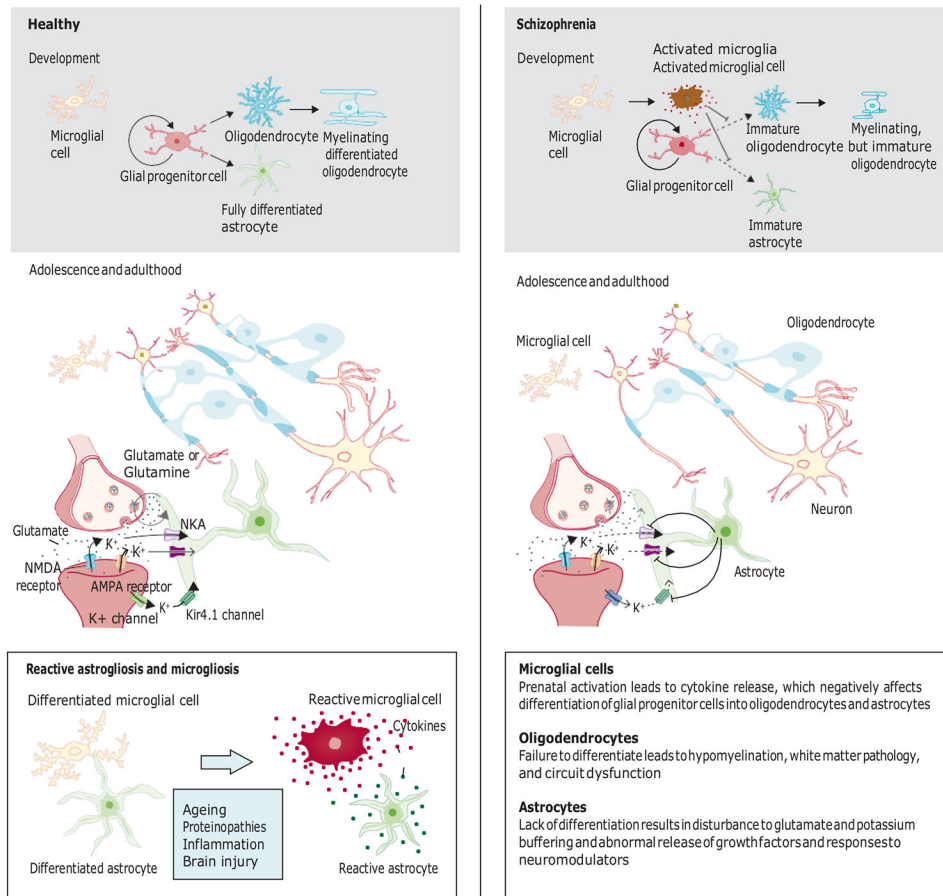


Figure: A unified gliocentric model of schizophrenia

In development, glial progenitor cells give rise to oligodendrocytes and astrocytes, following a tightly regulated differentiation pathway. We hypothesise that untimely immune activation of microglia during fetal development suppresses the differentiation of glial progenitor cells, resulting in immature morphology and abnormal functionality and numbers of oligodendrocytes and astrocytes. Deficits in oligodendrocyte maturation competence lead to structural hypomyelination and loss of white matter integrity. Failure of astrocytic differentiation results in reduced synaptic coverage and abnormal buffering of glutamate and potassium. The proposed model links disrupted glial cell differentiation to the initial event in schizophrenia, which differs from other conditions involving glial cells, in which fully differentiated glial cells undergo reactive changes (lower left panel). Reactive changes in glial cells might, however, occur in schizophrenia as secondary responses to, for example, synaptic dysregulation, lack of sleep, or drug exposure. NKA=sodium–potassium pump. K +=potassium ions.