Role of Neuro-Immunological Factors in the Pathophysiology of Mood Disorders: Implications for Novel Therapeutics for Treatment Resistant Depression

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Abstract Mood disorders are associated with persistently high rates of morbidity and mortality, despite the widespread availability of antidepressant treatments. One limitation to extant therapeutic options has been that nearly all approved antidepressant pharmacotherapies exert a similar primary action of blocking monoamine transporters, and few options exist for transitioning treatment resistant patients to alternatives with distinct mechanisms. An emerging area of science that promises novel pathways to antidepressant and mood-stabilizing therapies has followed from evidence that immunological factors play major roles in the pathophysiology of at least some mood disorder subtypes. Here we review evidence that the compounds that reduce the release or signaling of neuroactive cytokines, particularly IL-1β, IL-6, and TNF- α , can exert antidepressant effects in subgroups of depressed patients who are identified by blood-based biomarkers associated with inflammation. Within this context we discuss the role of microglia in central neuroinflammation, and the interaction between the peripheral immune system and the central synaptic microenvironment during and after neuroinflammation. Finally we review data using preclinical neuroinflammation models that produce depression-like behaviors in experimental animals to guide the discovery of novel neuro-immune drug targets.

Keywords Depression • IL-1 β • IL-6 • Microglia • Mood disorders • Neuroimmunology • Neuroinflammation • P2X7 • TNF- α

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Mood disorders constitute clinically pleomorphic syndromes consisting of behavioral and experiential changes in the emotional, cognitive, visceral, and appetitive domains that show moderate to high heritability, but remain idiopathic with respect to etiology. The main mood disorders, major depressive disorder (MDD) and bipolar disorder (BD) show relatively high lifetime prevalence rates [1] and despite the availability of many antidepressant drugs, MDD is ranked by the World Health Organization (WHO) as the highest global cause of "years of life lived with disability" for all age groups. The persistence of this global public health problem partly reflects the limited efficacy of extant therapies, as about one third of MDD patients do not achieve remission despite multiple trials using different treatments, while another third experience illness relapse and recurrence despite continued adherence to initially effective treatments [2–4]. One limitation of extant antidepressant pharmacotherapies is that they essentially all target biogenic amine based mechanisms, so that for patients who do not respond to such mechanisms, therapeutic options with distinct mechanisms have been largely unavailable.

Notably, the results of studies that compared depressed patients who respond to monoamine reuptake inhibiting agents versus those who do not consistently have shown that the non-responders manifest abnormal elevations in a variety of pro-inflammatory immunological markers [5–9]. These data converge with evidence that factors within the innate and the adaptive immune system play roles in the pathophysiology of MDD and BD, potentially thereby illuminating new targets for novel therapeutics in mood disorders [10, 11]. As reviewed below the findings that administration of pro-inflammatory cytokines such as interferon-alpha or low dose endotoxin can induce depressive symptoms in a subset of humans who have not previously been depressed [12], along with the implication of immune pathway dysregulation by genome-wide association studies (GWAS) of primary mood disorders, suggest that some individuals have a biological diathesis to manifest depressive symptoms under immune challenge [13, 14]. Such conclusions have been corroborated pre-clinically by similar phenomena, specifically by showing that immune activation produces depression-like behaviors in repeatedly stressed animals and that these behaviors can be prevented or reversed by anti-inflammatory treatments [15]. Similarly, an emerging clinical literature provides evidence that some types of anti-inflammatory treatments can produce antidepressant effects in depressed patients with peripheral blood evidence of inflammation [16].

1 Interplay of the Immune System and the Central Nervous System (CNS)

The emerging neuro-immunological literature suggests that immune cells in the periphery and/or the brain interact with neurons in the CNS to play roles in the pathophysiology of mood disorders [11, 17]. These data point to the existence of a bi-directional immune-connectivity between the peripheral and central compartments [18-20]. The interplay of the immune system and the CNS involving pro-inflammatory cytokines, chemokines, and related molecular processes that lead to microglial activation and astrogliosis is referred to as *neuroinflammation*. However, in the CNS the biological concomitants of an inflammatory state differ in many respects from conventional inflammation involving peripheral immune cells [21]. Thus the neuroimmunology field has broadened in perspective to also encompass the mechanisms by which the peripheral immune system modulates central neurophysiology. In contrast, the neuroinflammatory changes in microglia, astrocytes, and oligodendrocytes that putatively contribute to the causal mechanisms underlying multiple sclerosis, Parkinson's disease, and epilepsy are generally absent in mood disorders. For example, post mortem studies of glial cell function, structure and density do not show the astrocytosis and amoeboid microglial morphology that is manifest in multiple sclerosis, trauma, or neurodegeneration. Instead such studies have demonstrated reductions in oligodendroglia, impaired astroglial function, and intermediate morphologies of activated microglia [22]. One exception to this general set of findings in post mortem studies of mood disorders involves elderly patients characterized by a late age of depression onset; such patients show clinical and neuropathological evidence for a pathophysiological process mediated via cerebrovascular disease, including astrogliosis, inflammation, and other histopathological correlates of ischemic disease [23]. Nevertheless, debate remains whether neuroinflammatory processes play pathological or adaptive/compensatory roles in the pathophysiology underlying early onset mood disorders, which instead have been associated with a combination of genetic and environmental (e.g., early life trauma) risk factors [24-26].

In the CNS, bone marrow derived immune cells have a restricted access due to an intact blood-brain barrier (BBB) and blood-CSF barrier. During an injury or infection when this barrier is compromised, peripheral immune cells can penetrate the CNS causing neuroinflammation. Nevertheless, other conditions exist in which macrophages and monocytes from the periphery can migrate into the CNS [27, 28] and the CNS lymphatics may serve as conduits of peripheral to central cellular migration [29, 30]. In addition, and probably more pertinent to the topic of neuroinflammation, microglia constitute the critical cell types that change from a "surveillance" mode to a "response" mode during injury and disease pathology. Resting microglia manifest a distinct "ramified" morphology whose function is to sense the local environment and maintain homeostasis among the neurons, astrocytes, and oligodendrocytes that participate in synaptic function [31] and transmission. During pathology associated with neuroinflammation, microglia respond by adopting an amoeboid morphology and release gliotransmitters such as pro-inflammatory cytokines (IL-1 β , IL-6, TNF- α , IFN- γ), chemokines, glutamate, ATP, nitric oxide, and reactive oxygen and nitrogen species that alter the neuro-glia functional interactions [32]. Microglial activation also involves up-regulation of cellular markers, increased microglial proliferation and migration, and a shift in structure and function towards an "M1" pro-inflammatory phenotype. The M1 phenotype occurs in response to tissue injury, stress, and infection as part of the adaptive immune response which ultimately leads to reparative processes. The repair is mediated by microglia, predominantly of the anti-inflammatory (M2) phenotype, which are more phagocytic in nature. Nevertheless, microglia more commonly exist in a range of phenotypes that are intermediate in morphology between the M1 and M2 [33–35]; the differential roles of M1/M2 microglia and their role in CNS (patho)physiology are reviewed elsewhere [36, 37]. Notably, post mortem assessment of brain tissue from patients with MDD and BD revealed that the microglia manifest such an intermediate "activated" morphology associated with greater quinolinic acid expression (implying pro-inflammatory activation of the kynurenine pathway) in the subgenual anterior cingulate cortex area consistently implicated in the pathophysiology of mood disorders [38].

One type of phagocytic activity performed by the microglia involves synaptic pruning that regulates interneuronal connectivity and restores the optimal multipartite synaptic function from the altered states that arise during neuroinflammatory states [39–41]. Thus in mood disorders it has remained unclear whether microglial activation manifests as a reparative response or instead comprises a pathological mechanism that initiates disruption of normative neurophysiology [42]. In other neuropathological conditions, the extant data suggest that neuroinflammation can play pathological roles under some conditions and adaptive/restorative roles in others, with both roles potentially co-existing within the context of a particular CNS disease states. Nevertheless, chronic and/or dysregulated neuroinflammation eventually contributes to a pathological phenotype within the CNS. For example, the extant preclinical and emerging clinical data suggest that glial factors released from microglia and astrocytes during neuroinflammation modulate synaptic plasticity and neurogenesis and impact the neurocircuitry in a manner that can manifest behaviorally in much of the symptomology that defines mood disorders.

Peripheral immune cells also play roles in CNS function that include supporting learning and memory [43–48], protecting against pathogens (e.g., as evidenced by IFN- γ -mediated control of *Toxoplasma gondii* [49], and inducing neuropathology (e.g., in multiple sclerosis). The peripheral immune system also can play a beneficial or healing role in CNS pathology. For example, a controlled amplification of the autoimmune response was associated with improved neuronal survival in rodent models of acute CNS injury [50] and chronic neurodegeneration [51, 52]. The

complex interplay between the peripheral immune system and the CNS in mediating beneficial components of the immune response to CNS pathology is only beginning to be elucidated.

Conversely, changes in peripheral immune cell populations have also been associated with CNS pathologies that do not feature clear penetration of the blood–brain or blood–CSF barrier by circulating cells. Pro-inflammatory cytokines released during peripheral infection are associated with behavioral correlates of depressive mood—termed sickness behavior [53]. Links between sickness behavior and a tryptophan metabolizing enzyme, namely indoleamine 2,3 dioxygenase (IDO), have been demonstrated [54], establishing a potential association between cytokines and monoamine deficiency.

Moreover, post-traumatic stress disorder (PTSD), a syndrome characterized by chronic anxiety, depression, and hyperarousal arising in the aftermath of traumatic stress constitutes a condition that links CNS pathology, stress, and immune system dysregulation at the level of inflammatory cascades and gene networks. For example, in soldiers studied before and after deployment in areas of active conflict, CD14+ monocyte-associated factors were differentially regulated in PTSD sufferers [55]. Notably monocytes are primary producers of the neuroactive cytokines IL-6, IL-1 β , and TNF- α , which have been linked to mood disorders in preclinical studies and clinical populations (reviewed elsewhere in this volume). CD14+ monocytes are mobilized into circulation primarily by CCL2, a chemokine produced by glial [56] and blood–brain barrier cells [57] during neuroinflammation. These findings implicate a link between glial activation and loss of peripheral immune homeostasis leading to chronic feedback between the CNS and periphery in PTSD.

2 Abnormalities in Immunological Factors in a Subset of Patients with Mood Disorders Suggest Novel Antidepressant Targets for Such Subgroups

A rapidly expanding scientific literature suggests that alterations in immune system function and neuroinflammation play major roles in the pathophysiology of at least some subtypes of mood disorders [17, 58–61]. This evidence has encouraged targeted and rational drug discovery efforts with a view to intervene using immune modulating treatments for mood disorders [16]. Immune mediators for which the mean concentrations are increased in the blood and cerebrospinal fluid (CSF) of patients with mood disorders versus healthy controls, both when assessed at baseline and after exposure to stressors, include IL-6, IL-1 β , IFN- α , TNF- α , prostaglandin E2, and the chemokine CCL2 [62–66]. The mRNA transcripts for these cytokines and for other related innate immune system genes have also been elevated in peripheral blood cells in patients with mood disorders relative to healthy controls matched for age, BMI, smoking, and comorbid medical conditions [6, 22, 67–70]. The clinical significance of these findings is supported by evidence that the elevations of these cytokines in the plasma or CSF of patients with MDD or BD relative to controls are correlated with illness severity and/or suicidality ratings (reviewed elsewhere in this volume). Moreover, *successful* response to conventional antidepressant drugs is associated with reductions in these cytokine levels in depressed patients, although *non-response* to conventional antidepressants is predicted by higher IL1- β , IL- β , and CRP levels in the pre-treatment baseline [6, 9, 71]. The preliminary evidence reviewed below suggests that in depressed patients who manifest both resistance to monoamine reuptake inhibitor antidepressant agents and elevation in pro-inflammatory cytokines/acute phase proteins, certain classes of anti-inflammatory agents can produce antidepressant effects.

The relationship between immune challenge and the development of "sickness behavior" as well as other more clearly pathological depressive symptomatology is instructive in considering the etiology of mood disorders. In one of the clearest examples indicating that elevated cytokine signaling can cause depressive symptoms, immune challenge with interferon- α (IFN- α) during the treatment of hepatitis C or other medical conditions reliably has induced the major depressive syndrome (and less commonly manic symptoms) in 30-40% of previously non-depressed humans [64]. This neuropsychiatric sequelae of IFN- α converges with other types of evidence to suggest that elevated signaling of some neuroactive cytokines can play a causal role in inducing depressive symptoms [72]. Within the days following IFN- α administration, previously non-depressed patients show a behavioral complex that includes anorexia, fatigue, lower mood, reduced social interaction, and reduced engagement in pleasurable activities, a symptom complex referred to in the research literature as "sickness behavior" [73]. Notably, in the subset of patients who receive IFN- α who go on to develop major depressive episodes, more specific depressive symptoms such as pessimism, anxiety, and suicide ideation arise later than the initial appearance of sickness behavior, and the likelihood of developing an MDE continues to rise with longer time spent receiving IFN- α [74]. Thus the symptoms of the major depressive episode differ from those of sickness behavior by magnitude in some cases (e.g., more severely depressed mood and pervasive anhedonia in MDE) and by quality in others (e.g., pathological anxiety and suicide ideation). In addition there is also evidence that IFN- α induced depression differs from depression arising in medically healthy MDD subjects by the presence of pathological guilt in the latter, but not in the former condition [75, 76]. These data are notable within the context of evidence that primary MDD is heterogeneous, with subgroups that are distinguishable on the basis of clinical symptomatology as well as immunological markers [58], as reviewed below.

The prevailing hypothesis holds that IFN- α induced major depressive episodes constitute an example of a "pro-inflammatory state induced mood disorder" that manifests in some individuals exposed to IFN- α on the basis of a biological predisposition. A corollary to this hypothesis posits that a subset of the "primary" MDD population also manifests depressive symptoms due to the influence of elevated neuroactive cytokine signaling caused via other etiologies. An example of the existence of a biological predisposition toward IFN- α induced depression

was provided by the report of a single nucleotide polymorphism (SNP) in the IL-6 receptor gene that resulted in lower IL-6 expression, and also was associated with decreased susceptibility to the development of depressive symptoms during IFN- α treatment [77].

Notably, some patients who manifest IFN- α induced major depressive episodes improve during SSRI treatment, leading physicians to prophylactically initiate SSRIs in the weeks prior to initiating IFN- α for some patients [78]. Of the symptom domains affected by IFN- α , however, the depressed mood symptom dimension is most responsive, whereas the anxiety, cognitive and neurovegetative symptoms appear less responsive (or unresponsive) to prophylactic SSRI treatment [79]. These observations hold intriguing therapeutic implications, because in patients with primary mood disorders, higher blood levels of proinflammatory cytokines or their mRNA transcripts predict non-response to treatment with SSRIs or other conventional antidepressants [6–9]. It is conceivable that depressed patients who both manifest chronic inflammation and prove nonresponsive to conventional antidepressant drug treatment may benefit from immune modulating treatments.

The hypothesis that anti-inflammatory agents may exert antidepressant effects in depressed patients has been tested both in patients who have primary MDD and in *patients with autoimmune disorders who manifest clinically significant depressive symptoms* [16]. For example, patients with psoriasis who received the anti-TNF- α agent etanercept showed significant improvement in depressive symptoms in response to drug versus placebo as assessed using conventional depression rating scales, and this difference was evident earlier than the associated changes in pain or skin lesions [80], implying the antidepressant response occurred independently of psychological benefits related to the improvement in the skin lesions per se. In this study patients treated with etanercept also had significant improvements in fatigue. Notably, while the improvements in fatigue correlated with decreasing joint pain, the improvements in depression were less correlated with objective measures of skin clearance or joint pain.

3 Immunological Biomarker Data from Mood Disordered Samples Shows Heterogeneity That May Hold Therapeutic Implications

In studies of *primary MDD patients* treated using anti-inflammatory agents, the extant data suggest that subgroups characterized by high levels of pro-inflammatory biomarkers are most likely to benefit (see below). This observation appears intuitive when the findings that mean concentrations of cytokine levels are elevated between depressed and control samples are considered in further detail. The distribution of the immunological data from these studies suggests that the differences reported in *mean values* are attributable to a subset(s) of the depressed patients. This observation appears consistent with accumulating evidence of biological and genetic

differences between subtypes of depressed subjects with MDD, who otherwise appear phenotypically homogeneous in many aspects of symptom presentation. For example, from the Netherlands Study of Depression and Anxiety database, Lamers et al. [58] used subgroups defined initially using cluster analysis of depressive signs and symptoms, and then further differentiated these subtypes based on serum protein profiles. The identified analytes consisted largely of inflammatory (e.g., CRP) and metabolic markers (e.g., insulin), supporting the conceptualization of a subtype(s) characterized by metabolic disturbances and inflammation. These researchers [81] also showed that these subgroups appeared stable across time, with patients moving between different levels of severity, but not between subtypes, during longitudinal follow-up. In another example, data from the Mood Inflame Consortium identified three MDD subtypes: one manifest in MDD patients aged >28 years that was characterized by *increased* expression of monocyte genes and decreased expression of glucocorticoid receptor (GR) α versus β subunit ratio, a second in MDD patients <28 years of age who showed a severe course of depression (characterized by recurrent type, illness onset <15 years of age, history of childhood trauma, and prominent panic/arousal symptoms) but monocyte gene expression similar to healthy controls, and a third also manifest in MDD patients <28 years of age characterized by a milder illness course (most with first episode of depression, age at onset ≥ 15 years, and absent panic symptoms) that exhibited a strongly *reduced* inflammatory monocyte activation compared to controls [82].

Within the bipolar spectrum of mood disorders, another study from the Mood Inflame Consortium identified a biomarker signature composed of multiple immunological factors that discriminated the majority of BD patients from healthy controls. Using whole-genome expression profiling of RNA obtained from purified CD14+ monocytes, Padmos and colleagues reported elevated mRNAs of inflammatory (e.g., TNF, PDE4B, IL-1 β , IL6, TNFAIP3), trafficking, survival (e.g., BCL2A), and mitogen-activated protein kinase pathway (e.g., MAPK6, ATF3) genes in BD subjects in various illness phases, as well as in affected offspring of BD parents [69]. Notably, in peripheral blood mononuclear cells (PBMC) from the same subjects assessed via fluorescence-activated cell sorting (FACS) analysis, the percentages of anti-inflammatory CD4+CD25highFoxP3+ regulatory T cells were higher in BD patients <40 years of age, while percentages of Th1, Th2, and Th17 cells were normal. Together these results thus showed enhancement of both pro-inflammatory monocyte and anti-inflammatory T cell mediators in BD [83].

4 Novel Drug Targets at the Crossroads of Neuroimmunology and Mood Disorders

With continued and refined understanding of the role of immune cells and their mediators in the periphery and the CNS, it is anticipated that new mechanisms will be discovered that can exert antidepressant and mood-stabilizing effects in primary

mood disorders. Several comprehensive reviews have highlighted potential drug targets in neuroimmunology for mood disorders [11, 21, 84]. In this chapter, we summarize evidence that highlights TNF- α , IL-6, and IL-1 β signaling in the pathophysiology of mood disorders.

<u>TNF- α </u>: TNF- α signaling appears to play a major role in mood disorders [85]. In meta-analyses of clinical studies, plasma TNF- α correlated with depression severity and level of resistance to conventional antidepressants [62]. A causal relationship between TNF- α elevation and depressive symptoms was suggested by observations that in patients with immunological diseases such as rheumatoid arthritis and psoriasis, anti-TNF- α treatment alleviates depressed mood; as reviewed above, these antidepressant effects do not appear attributable simply to improvement in sickness symptoms, such as fatigue, or in the underlying autoimmune disorder [86]. Consistent with these observations, the TNF- α receptor 1, TNF- α receptor 2, and TNF- α knockout mouse models all show antidepressant-like phenotypes [87, 88]. Likewise, systemic administration of antibodies targeting TNF- α in chronic models of stress reversed the anhedonic behaviors, suggesting that TNF- α signaling contributes to depressogenic behaviors in rodents [89, 90].

Nevertheless, a clinical study of the efficacy of infliximab (a monoclonal antibody against TNF- α) in depressed patients generated negative results on depressive symptoms rating using a conventional depression rating scale [91]. A *post hoc* investigation of data from this study, however, revealed a significant positive correlation between clinical improvement and pre-treatment levels of the nonspecific inflammation marker, CRP, raising the possibility that antidepressant effects may be limited to individuals who manifest a pro-inflammatory diathesis. Nevertheless, because the test of the *a priori* hypothesis in this study was negative, the question has remained whether targeting TNF- α via large molecules introduced in the periphery alone can produce an antidepressant effect (since very low proportions of peripherally administered monoclonal antibodies enter the brain following acute treatment), or whether therapies that reduce TNF- α signaling must instead directly engage targets in the CNS.

<u>IL-6</u>: In studies of MDD or BD one of the more highly replicated biomarker abnormalities has been an elevation in peripheral blood IL-6 concentrations [92]. Notably during IFN- α treatment the magnitude of the increase in plasma and CSF IL-6 levels correlates positively with depressive symptom severity. Conversely, the above-mentioned functional polymorphism in the promoter region of the IL-6 gene (rs1800795) that results in decreased IL-6 expression is associated with a significantly lower risk for developing major depressive episodes during IFN- α treatment [77]. The relationship to IL-6 function is compatible with findings that, in patients with primary mood disorders higher IL-6 levels in the CSF correlated with suicidality, and elevated IL-6 levels in the plasma correlated with non-responsiveness to conventional antidepressant drugs [93]. In contrast, during the euthymic (i.e., asymptomatic) phase of BD, the CSF concentration of IL-6 was decreased with respect to healthy controls, despite the same BD subjects showing an abnormal elevation in the CSF levels of IL-1 β [65].

Although IL-6 can be released by immune cells in the CNS as well as in the periphery, preclinical evidence suggests that elevated IL-6 release from peripheral immune cells is sufficient to induce depressive behaviors, irrespective of central immune system activation. In studies conducted by Hodes and colleagues [15] to elucidate the biological basis of susceptibility to depression-like behaviors under stress, mice that developed a persistent depression-like phenotype in response to social defeat stress (SDS) were compared to genetically identical mice that did not develop depression-like behaviors under SDS. The susceptible animals differed from the resilient animals by showing elevated basal IL-6 levels in the pre-SDS condition and higher IL-6 release in response to the stressed condition. In addition, white blood cells sampled pre-SDS from susceptible mice showed higher LPS-induced IL-6 release ex vivo compared to cells from resilient mice. Crucially, the susceptibility to the depression-like phenotype could be altered toward either susceptibility or resilience by generating bone marrow chimeras that had hemopoetic stem cells transplanted from high IL-6 expressing mice or IL-6 knockout mice, respectively. The bone marrow recipients in these studies had received radiation to their bodies while the head was shielded, so the hemopoetic stem cells in periphery conferred the susceptibility to the depression-like phenotype under stress.

<u>IL-1β</u>: In contrast to the therapeutic potential offered by neutralizing IL-6 predominantly in the periphery, the extant data suggest that for the pro-inflammatory cytokine IL-1 β , reducing signaling in the brain may prove critical to achieving antidepressant effects. IL-1 β is probably the most potent pro-inflammatory cytokine released from microglia in the brain. Clinical studies found that IL-1 β is present at abnormally higher levels in plasma, CSF, and postmortem brain tissue of individuals with mood disorders, and that IL-1 β levels correlated positively with depression severity [63, 65]. Anisman and colleagues reported increased IL-1 β production from lymphocytes in patients with dysthymic disorder and a modest correlation existed between the cytokine and depressive symptoms [94]. In studies of primary mood disordered subgroups, IL-1 β has been linked with both geriatric depression and postpartum depression [95, 96].

In animal models of stress-induced depression-like behaviors, several groups showed that IL-1 β signaling is critical to the acquisition of the depression-like phenotype [97, 98]. The development of the depressive behavioral phenotype during chronic stress can be blocked by IL-1 receptor antagonists, and is absent in IL-1R receptor knockout mice [99]. In addition, manipulation of central IL-1 β , either by exogenous administration of IL-1 β directly into the brain, or by selective ablation of signaling via pharmacology or genetics, produced behavioral analogues of depression when IL-1 β was increased, or antidepressant-like effects when IL-1 β was decreased [100]. The IL-1 β driven changes in the brain resulted in decreased neurogenesis in the hippocampus [99] and increased corticosterone response to stress in the periphery [101], suggesting an interplay between stress-induced-IL-1 β release and HPA axis function.

Recently, it was shown that both acute and chronic stress increase brain IL-1 β release [102] [100]. Stress-induced IL-1 β release appears to be driven by

ATP-induced activation of the P2X purinoceptor 7 ion channel (P2X7), and genetic deletion of P2X7 receptors results in antidepressant-like reversal of stress-induced depressogenic behaviors in rodents [102, 103]. Other experimental evidence has similarly demonstrated that P2X7 activation causes release of IL-1 β ([104]). The initiation of transcription and translation of the pro-form of IL-1 β is induced by activation of Toll-like receptor (TLR), but it is the second signal from P2X7 (due to ATP activating the ion channel) that results in maturation and release of the pro-inflammatory IL-1 β cytokine; this process has been referred as a "two-hit" model of IL-1 β release. Priming of the TLRs is achieved by factors such as cellular debris, by endotoxins, by damage- and pathogen-associated molecular pattern molecules (DAMPs and PAMPs, respectively). Since P2X7 is abundantly expressed in blood cells, IL-1 β release in the blood has been used as a biomarker of P2X7 activity in both preclinical and clinical assessment of target engagement.

Based on robust microglial expression of P2X7, and IL-1ß signaling leading to neuroinflammation, CNS penetrable P2X7 antagonists would be potentially beneficial for treating mood disorders, and there is growing evidence that strengthens the role of P2X7 in MDD and BD. Several human genetic studies have associated the highly polymorphic P2RX7 gene with the risk for developing both BD and MDD, and some of these mutations have been linked to a modulation of P2X7 channel function in vitro [105, 106]. The rs2230912-G allele exhibits a gain-of-function and human monocytes expressing this variant secreted more IL-1 β in response to activation of P2X7 than monocytes expressing a wild-type variant [107]. It is conceivable that such a variant in P2X7 receptors based in human microglia enhanced IL-1 β release (or production), would lead to leading to neuroinflammation over time. Nevertheless, several other GWAS studies have not confirmed the association between P2RX7 variants and the risk for mood disorders [108], so the relationship between the variation in P2RX7 and depression is not yet established. The lack of clarity for a genetic association of P2RX7 variation in the risk for mood disorders (or any disease phenotype) is perhaps not surprising as the underlying factors of such pathologies are often a result of interplay between genetic (often many genes), environmental, and developmental factors.

In addition to the human genetic literature, several laboratories have demonstrated that P2X7 knockout mice manifest a protective phenotype in models of depression and mania, strengthening the hypothesis that P2X7 antagonism may be therapeutically beneficial in mood disorders. Consistent with the antidepressant phenotype observed in P2X7 knockout mice, emerging data suggest that P2X7 antagonists can reverse depressogenic behaviors in animal models. For example, pharmacological antagonism of P2X7 (by AZ-10606120 and A-804598) restored the deficit observed in the preference for a sucrose solution (a putative behavioral analogue of anhedonia) induced by either chronic stress or systemic administration of lipopolysaccharide (LPS) administration [103]. Recently, it was shown that a P2X7 selective, brain-penetrant antagonist was efficacious in chronic stress models in rats [102]. In addition, a large corpus of evidence suggests that manipulation of central IL-1 β (by either exogenous administration or selective ablation of signaling by pharmacological or genetic manipulation) results in depression-like behaviors when IL-1 β is increased, or in resilience against the development of depression-like behaviors when IL-1 β is decreased [97, 99, 109]. These observations appear consistent with the above-mentioned findings that IL-1 β levels are abnormally elevated in the plasma, cerebrospinal fluid (CSF), and postmortem brain tissue obtained from MDD and BD patients [63, 65, 110].

Preclinical data also suggest that P2X7 antagonism may produce anti-manic or mood-stabilizing effects in BD [111]. For example, P2X7 antagonism produced attenuation of amphetamine-induced sensitization of hyperactivity [112, 113], a putative rodent model of mania-like behavior, and similar phenotypes were observed in P2X7 knockout mice [111] suggesting a potential therapeutic role of P2X7 antagonism in the manic phase of BD. Taken together, it remains plausible that a selective and brain-penetrant P2X7 antagonist may be therapeutically beneficial in mood disorders, especially targeting treatment resistant patient populations.

5 Conclusion

Neuroimmunology stands at the interface of emerging biology and breakthrough therapeutics for mood disorders. Taken together, the extant data support the hypothesis that elevated cytokine levels contribute to the pathophysiology of depression and the neurobiological mechanisms underlying resistance to conventional antidepressant drugs, in at least a subpopulation of depressed patients. They also suggest that specific cytokines (such as TNF- α , IL-6, and IL-1 β) and their effectors and regulators (such as P2X7) may constitute novel therapeutic targets for depression. However, the extant postmortem data also indicate that mood disorders are not associated with classical neuroinflammation, and in vivo blood-based biomarker studies suggest that not all patients suffering from mood disorders manifest an inflammatory component. Consequently, for clinical proof-of-concept studies with compounds that target signaling of microglia, astrocytes, or cytokines/ chemokines, it may prove necessary to discriminate the patient population suffering from concomitant depression and neuroinflammation through the aid of immunological biomarkers.

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