

A Review of Biomarkers in Mood and Psychotic Disorders: A Dissection of Clinical vs. Preclinical Correlates

Sarel J. Brand¹, Marisa Möller¹ and Brian H. Harvey^{1,2,*}

¹Division of Pharmacology and ²Center of Excellence for Pharmaceutical Sciences, School of Pharmacy, North-West University, Potchefstroom, South Africa



Abstract: Despite significant research efforts aimed at understanding the neurobiological underpinnings of mood (depression, bipolar disorder) and psychotic disorders, the diagnosis and evaluation of treatment of these disorders are still based solely on relatively subjective assessment of symptoms as well as psychometric evaluations. Therefore, biological markers aimed at improving the current classification of psychotic and mood-related disorders, and that will enable patients to be stratified on a biological basis into more homogeneous clinically distinct subgroups, are urgently needed. The attainment of this goal can be facilitated by identifying biomarkers that accurately reflect pathophysiologic processes in these disorders. This review postulates that the field of psychotic and mood disorder research has advanced sufficiently to develop biochemical hypotheses of the etiopathology of the particular illness and to target the same for more effective disease modifying therapy. This implies that a “one-size fits all” paradigm in the treatment of psychotic and mood disorders is not a viable approach, but that a customized regime based on individual biological abnormalities would pave the way forward to more effective treatment. In reviewing the clinical and pre-clinical literature, this paper discusses the most highly regarded pathophysiologic processes in mood and psychotic disorders, thereby providing a scaffold for the selection of suitable biomarkers for future studies in this field, to develop biomarker panels, as well as to improve diagnosis and to customize treatment regimens for better therapeutic outcomes.

Keywords: Antidepressant, biomarker panel, GABA-glutamate, genomics-proteomics, immune-inflammation-redox, kynurenine-cytokine, neurotransmitters, nitric oxide, schizophrenia.

INTRODUCTION

Major depression (MD), bipolar disorder (BPD) and psychotic disorders (e.g. schizophrenia) are often misdiagnosed, leading to inadequate treatment and devastating consequences [1, 2]. MD is among the most debilitating diseases worldwide, with a life-time prevalence of up to 20% [3], and even though major advances have been made in developing new drugs, less than 50% of patients achieve remission after antidepressant treatment [4]. Bipolar disorder affects approximately 1.2% of the population worldwide [5], and differs from MD with a unique hallmark of mania (elevated mood or euphoria, hyper-activity with a lack of need for sleep, and an increased optimism) which frequently leads to a deficit in the patient’s judgment [6]. On the other hand, schizophrenia is a debilitating neuropsychiatric disorder, typically emerging during adolescence or early adulthood and continuing to plague patients suffering from the disease to varying degrees throughout their lifetime [7]. Approximately 1% of the general population worldwide is affected by the disorder and the life expectancy of patients with schizophrenia has been demonstrated to be nearly 20% shorter than that of the general population [8].

Despite an abundance of research, the pathogenesis and aetiology of mood and psychotic disorders remain unclear, challenging the diagnosis and treatment of these disorders [9], mainly for the following reasons:

- Diagnoses of typical psychiatric disorders are primarily based on operationalized behavioural diagnostic systems either as self-reported symptoms by patients or observations by clinicians, being confirmed against diagnostic criteria set out in the *Diagnostic and Statistical Manual of Mental Disorders 4th/ 5th ed. (DSMIV/V)* and *International Statistical Classification of Diseases, 10th Revision* [10].
- Laboratory diagnostic and screening tools, such as a non-invasive blood-based test, remain elusive [11], while mood and psychotic symptoms may overlap with other neurological and psychiatric problems [9].

Clinically useful biomarkers in these disorders could therefore significantly improve diagnosis and treatment and has been one of the holy grails of MD, BPD and schizophrenia research [12]. However, the likelihood of any single biomarker achieving a high enough degree of sensitivity and specificity for mood and psychotic disorders is relatively low. Biomarker panels may represent an attainable alternate to a single-biomarker approach [13]. Common features of this method include correlates attributed to the individual which may determine the

*Address correspondence to this author at the Division of Pharmacology and Center of Excellence for Pharmaceutical Sciences, School of Pharmacy, North-West University, Potchefstroom, South Africa; Tel: 018 299 2238; E-mail: Brian.Harvey@nwu.ac.za

presence or absence of a state of sickness or that may even predict response to treatment [13]. Biomarkers may also indicate the presence of a pathophysiological process that can be addressed with a preventive treatment [14], as well as highlight “state” and/or “trait” markers. Therefore, the identification of biomarkers prior to onset of depressive and bipolar symptoms or psychosis has enormous potential importance for the design of future preventive strategies.

The Biomarkers Definitions Working Group of the National Institutes of Health Group [15] (2001) defined a biomarker as “a characteristic that is objectively measured and evaluated as an indicator of normal biological processes, pathogenic processes, or pharmacologic responses to a therapeutic intervention” [15]. Biomarkers should also provide high levels of sensitivity and specificity (>80%) in the detection and classification of MD, BPD and psychotic disorders [16], in order to be clinically useful.

There are many potential biomarkers for mood and psychotic disorders and previous studies have tested specific biomarkers based on the hypotheses of monoamine dysfunction, altered immune-inflammatory processes, neuroendocrine dysfunction and disturbances in neuroplasticity [17]. However, considering the most recent neuroanatomical basis for these illnesses, as well as relevant hypotheses of their aetiology, this review will provide an overview of potential biomarkers that could contribute to an initial multi-analyte biomarker panel of mood and psychotic disorders. Such biomarkers include molecules and/or processes directly or indirectly connected with growth factors, neurotransmitters, oxidative stress, inflammation, neuro-imaging, genetic, proteomic and neuronal resilience markers. In order to do this, we have collated neuroimaging and neurobiological findings from clinical studies as well as data from validated translational animal models, to assist in developing a putative, uniform biomarker panel for MD, BPD and schizophrenia.

There can be no doubt as to the value of translational animal models in drug discovery and in identifying novel neurobiological targets. Considering mood and psychotic disorders, these models include for example social isolation rearing (SIR) and the glutamate N-methyl D-aspartate (NMDA) antagonist models of schizophrenia [18, 19], and the Flinders Sensitive Line (FSL) or chronic mild stress (CMS) rat models of MD [20, 21]. Developing an appropriate animal model to mimic BPD has proven to be an arduous task, it being difficult to establish a model that combines symptoms of MD, mania and euthymia in an alternating manner as is observed in BPD. Instead, animal models have been developed to express features central to either MD or mania using pharmacologic (amphetamines and ouabain), environmental (e.g. behavioral despair; sleep deprivation), or genetic (e.g. FSL rat) models. Therefore in parallel with our analysis of the clinical scenario, we will also closely scrutinize appropriate animal models for correlation with clinical findings. This review will cover MD, BPD and schizophrenia with respect to the noted biomarkers and across clinical and pre-clinical correlates.

THE NEUROANATOMY AND NEUROCIRCUITRY OF PSYCHIATRIC ILLNESS

Clinical Correlates

Neuroimaging methods, such as structural magnetic resonance imaging (MRI), functional magnetic resonance imaging (fMRI), diffusion tensor imaging (DTI) and positron emission tomography (PET), provide important evidence for underlying biological factors of MD, BPD and psychotic disorders [22]. Generally, areas of the limbic system, the hippocampus and frontal cortical areas are under scrutiny with regard to structural and functional neuroimaging research in mood disorders [23]. Importantly, there is increasing awareness of the interplay between specific neurocircuitry of the brain and behavioural pathology.

Depression

Neuroimaging studies have been central in identifying the key structures involved in the pathophysiology of MD, showing decreases in hippocampal volume of up to 15% in depressed patients [24], as well as reductions in grey-matter volume and glial density in the prefrontal cortex and the hippocampus [25]. Other studies in MD indicated large volume reductions in the anterior cingulate cortex (ACC) and orbitofrontal cortex (OFC) accompanied by lesser reductions in the prefrontal cortex, along with moderate reductions in the hippocampus, the putamen, and the caudate nucleus [26]. Lorenzetti *et al.* [27] reported volume reductions of the hippocampus, basal ganglia, the OFC and subgenual prefrontal cortex in patients suffering from MD, while more persistent forms of MD (which may include recurrent episodes or relapses and extended illness duration) are accompanied by an increased effect on regional brain volumes [28]. While reductions in hippocampal volume in MD may have a genetic component [29], it is also a function of illness duration [30] as well as poor compliance [28]. Moreover, PET studies have revealed consistently increased regional blood flow and glucose metabolism in the amygdala, orbital cortex, and medial thalamus but decreased blood flow in the dorsomedial/dorsal-anterolateral prefrontal cortex and anterior cingulate cortex in un-medicated MD patients [31]. N-acetyl-aspartate (NAA), an indicator of neuronal viability, was also reduced in frontal cortex and in subcortical regions of MD patients [32, 33].

Regarding antidepressant therapy, Frodl *et al.* [34] demonstrated increases in hippocampal volume in patients who were subjected to continual treatment with antidepressants for three years, while Mayberg *et al.* reported that patients who responded to antidepressant treatment presented with increased anterior cingulate metabolism at baseline when compared to non-responders and to healthy controls [35]. Importantly, Macqueen *et al.* noted a greater degree of hippocampal shrinkage in patients with a prior history of switching antidepressants, indicating the possible deleterious effect of non-compliance on hippocampal integrity [28]. Furthermore, it has been proposed by structural neuroimaging studies that the volume of regional structures, for example the ACC and hippocampus, may provide an estimate of response to treatment [36]. Clinical response has also been demonstrated

to be predicted by activity in the rostral ACC region as measured by electroencephalography (EEG) studies which identified activity localised to the rostral ACC region as being predictive of a clinical response to antidepressant medication [37]. Finally, favourable treatment outcomes have repeatedly been demonstrated to be associated with integrity of perigenual anterior cingulate volume [38]. Moreover, direct electrical stimulation of the striatum has been found to elicit a positive response in patients suffering from resistant MD and is bolstered by reports proposing the striatum as an important relay system between limbic and cortical structures [39]. ACC activity has also been positively related to a variety of treatment responses, including antidepressant pharmacotherapy and sub-chronic and experimental treatment strategies, including sleep deprivation [40], which suggests that ACC response is generalised across different treatment types.

Bipolar Disorder

In patients with BPD, neuroimaging studies have found enlargement of the amygdala [41] and reductions in the dorsal and ventral prefrontal cortices [42] while PET studies have found decreases in cortical metabolism and increased normalized subcortical metabolism in depressed patients with BPD [43]. Two meta-analyses of neuroimaging studies concluded that patients with BPD suffered from hypo-activation and gray matter reductions in cortical-cognitive brain structures and increased activation of the parahippocampal gyrus and amygdala [44].

Schizophrenia

In schizophrenia and other psychotic disorders, regions such as the ACC and dorsolateral prefrontal cortex (DLPC) have been emphasised [45]. Using cognitive paradigms, fMRI studies have demonstrated alterations in cerebellar activity in patients with schizophrenia, anxiety disorders and dementia (see review [46]). However, Farrow *et al.* [47] found that lateral and medial frontal regions and bilateral posterior temporal lobe regions feature structural losses in schizophrenia, whereas alterations in patients suffering from BPD were limited to bilateral inferior temporal gyri while deficits observed subsequently were limited to the ACC. Temporal lobe regions present with decreased activation in patients suffering from schizophrenia [48]. Additionally, EEG studies have demonstrated a reduction in the P300 wave amplitude, elicited in the process of decision making, in BPD and schizophrenia patients compared to control subjects [49]. Studies utilizing structural MRI have consistently observed temporal lobe abnormalities in schizophrenia, although results in BPD are less dependable [50]. Previous fMRI studies have also consistently reported anomalies in the prefrontal cortex in patients suffering either from a first episode or established schizophrenia [51, 52]. However, some of the evidence points to dorsolateral hyper-frontality, and especially for tasks which demand working memory, as well as increased activity in parietal regions [53]. Considering the progression from the prodromal phase to established chronic illness, patients with first episode and established schizophrenia show a gradual deterioration in frontal and striatal activation [54]. The most consistent findings in schizophrenia relating to cognition are detriments

in executive tasks requiring prefrontal cortical function, eg. a self-ordered working memory task [55] or anti-saccade eye movements [56], olfactory identification [57], and tasks that rely on rapid processing of information (eg. story recall) [58]. A recent ¹H-MRS study in schizophrenia patients measuring NAA and N-acetylaspartylglutamate (NAAG) found a significant increase in NAAG/NAA ratio in the ACC but no difference in the left frontal lobe, although an inverse correlation between frontal lobe NAAG and negative symptoms was observed [59].

Pre-Clinical Correlates

Depression

Reductions in hippocampal volume have been observed in FSL rats, a genetic model of MD, when compared to Flinders Resistant Line (FRL) controls and is associated with a decrease in the number of neurons and synapses in the hippocampus – these alterations are reversed after chronic imipramine therapy [60].

Bipolar Disorder

In the ouabain-induced rat model of bipolar mania, PET imaging suggests reduced cerebral glucose metabolism, and is prevented by pre-treatment with lithium which concurs with similar decreases in cerebral metabolism noted in BPD patients [61]. Furthermore, lithium prevented stress-induced alterations in the amygdala by preventing increases in dendritic branching of pyramidal neurons in this structure [62]. Unfortunately, a paucity of MRI studies remains a shortcoming in animal models of BPD.

Schizophrenia

Previous studies indicated that the SIR model, a neurodevelopmental animal model of schizophrenia, presents with significantly reduced PFC volume, reduced accumbal dendritic length and spine density, cytoskeletal alterations and loss of parvalbumin (PV)-containing interneurons [18, 63, 64]. Among the most robust pathologies observed in schizophrenia is a decrease in gamma-aminobutyric acid (GABA) signaling (discussed in section 3.1.4), deficits of which are limited to the class of GABAergic interneurons containing the calcium binding protein PV [65]. These neurons synapse on the cell body or axon initial segment of glutamatergic neurons and thus are positioned to potently regulate pyramidal cell output. Furthermore a decrease in PV interneuron functionality may lead to reduced inhibitory control over pyramidal cell activity and also reduce coordination in activity of large brain networks [66].

SIR rats without an enriched environment also present with a decrease in dendritic spine density in the dorsolateral striatum when compared to rats from an enriched environment [67, 68]. Moreover, a NMDA receptor antagonist model of schizophrenia, viz. the phencyclidine (PCP) model, presents with decreased synaptic spine density on frontal cortical neurones [64]. Interestingly, rats treated chronically with MK-801 (another NMDA receptor antagonist model) also show a reduction in the amount of PV-containing neurones in the dentate gyrus and CA1 region of the hippocampus, although this is not accompanied by alterations in the PFC [69]. Furthermore, chronic intermittent exposure to PCP

decreases NAA and NAAG levels in the temporal cortex, while it raises hippocampal NAAG levels [70]. Similarly, SIR reduces NAA in the temporal cortex without changes observed in the hippocampus, striatum or frontal cortex [71]. These changes may indicate neuronal dysfunction that mirrors alterations observed in schizophrenia, as discussed in the clinical section [59].

In order to determine which regions exhibit the most disease-relevant information as well as the most potential for predictive capacity, these neuroanatomical correlates need to be linked to biological and genetic markers to more accurately predict both the pathology underlying the disease and the clinical outcomes. Predicting clinical response will assist in early identification and to further stratify patients who may benefit from more intensive, alternative, or combined therapies.

BIOLOGICAL, GENETIC AND PROTEIN BIOMARKERS

Neuroendocrine and Circadian Rhythms

Various hormones, especially the HPA-axis, thyroid hormones, insulin, as well as altered circadian rhythm, have a pronounced influence on neurodevelopment and the neurobiology of MD, BPD and schizophrenia [72]. They also influence other hormones such as sex steroids, orexin, arginine vasopressin etc. that are also implicated in these disorders but for reasons of space cannot be covered here.

Circadian Rhythms

Clinical Correlates

Altered circadian rhythms occupy a critical role in how the brain copes with stressful experiences and ultimately in regulating behavioural responses [73, 74]. The influence of circadian rhythm on mood and behaviour has received much attention in recent years and implicates not only hormonal dysregulation in these disorders, but also includes disturbances in neurochemical, redox and inflammatory cascades in its sphere of influence [75, 76]. Indeed, these processes will be discussed in subsequent sections of this review. Output from the suprachiasmatic nucleus (SCN) of the hypothalamus, the master biological clock, is under regulation by serotonergic (5-HT_{2c}) and melatonergic (MT_{1/2}) receptors, the expression of which are regulated by various clock genes. Indeed, melatonin-mediated regulation of hippocampal plasticity as well as clock gene expression in hippocampal neurons suggests a hitherto poorly recognized aspect in our understanding and treatment of these disorders [78]. Altered SCN output to other hypothalamic centres, but also monoaminergic cell bodies in the brain stem, will lead to wide-spread disturbances in neuroendocrine as well as monoaminergic function [78]. It is therefore not surprising that a significant amount of preclinical and clinical data has described the association between altered circadian rhythms with genetic, environmental and developmental abnormalities precedent to the development of MD, BPD (mania symptoms) and schizophrenia (see recent reviews by Wulff *et al.* [75] and Karatsoreos [76]). It is of relevance that agomelatine, a recently available antidepressant that acts *via* re-entrainment of circadian rhythms, may also have a therapeutic role in disorders other than MD, in particular BPD and schizophrenia [77].

Pre-Clinical Correlates

Biomarkers of circadian rhythms in animals remain a shortcoming, with such studies relying heavily on endocrine markers (as will be clear in the following section). Nevertheless, alterations in this system have on numerous occasions been shown to not only be of importance in humans suffering from MD, BPD and schizophrenia, but also in translational animal models for these disorders. Most animal models selectively bred to display characteristics of MD feature disturbed diurnal rhythms, eg. the FSL rat [79], Wistar Kyoto rat [80] and mice bred for spontaneous helplessness [81]. Also, depriving animals of REM sleep has been suggested to model mania [82]. In animal models of schizophrenia, blind-drunk (Bdr) mice demonstrate fragmented rest and activity rhythms under a light/dark cycle – which is reminiscent of altered sleeping patterns in schizophrenic patients [83].

Cortisol

Clinical Correlates

A dysfunctional hypothalamic-pituitary-adrenal (HPA) axis has been implicated in MD, BPD and also schizophrenia [84-86], affecting adrenocorticotrophic hormone (ACTH) release and cortisol secretion from the adrenal cortex [87]. Elevated salivary levels of cortisol after waking may represent a biomarker for depression in adolescence [88]. Importantly, an abnormal cortisol response, such as a flatter diurnal cortisol pattern, implies an abnormal stress reactivity that correlates with a greater severity of depression [73, 74], suggesting that altered circadian rhythms occupies a critical role in how the brain copes with stressful experiences and ultimately in regulating mood. Although the dexamethasone suppression test has attracted interest as a promising diagnostic test for MD, there has not been a consistent approach to evaluate its clinical usefulness [89].

Steen *et al.* [90] found no significant difference in cortisol release during a mental challenge in schizophrenia and BPD patients, although blunted cortisol release was observed in male patients compared to controls in both disorders [90]. A significant increase in systemic cortisol metabolism in both schizophrenia and BPD patients has been described, with results in patients with schizophrenia vs. controls being most consistent [87]. Interestingly, studies in children at risk for developing psychosis lend further support to the suggestion that illness onset is predated by a degree of HPA axis abnormalities, rather than being a subsequent epiphenomenon [91]. A blunted cortisol awakening response may embody an early marker of susceptibility to develop psychosis which may even be genetically mediated, whilst increases in diurnal cortisol levels may develop only proximate to disease onset [91]. These studies reaffirm the status of the HPA-axis, particularly cortisol levels and metabolism, as a putative biomarker in MD, BPD and schizophrenia, and warrants further study.

Pre-Clinical Correlates

FSL rats have been found to be hypocortisolemic, while Wistar Kyoto rats present with increased levels of corticotropin releasing hormone (CRH) and ACTH. Brain levels of dehydroepiandrosterone (DHEA), an adrenal androgen

known to have antidepressant-like effects, has been demonstrated to be decreased in both FSL and Wistar Kyoto rats vs. healthy controls, Sprague Dawley and Wistar rats respectively [92]. Mice showing high reactivity to stress also present with symptoms resembling that in depressed patients and were demonstrated to have a flattened diurnal rhythm of glucocorticoid secretion [93]. Similarly, mice exposed to 6 weeks CMS presented with high plasma corticosterone levels and decreased hippocampal expression of glucocorticoid receptors [94]. Sleep deprivation, which has been used to induce an animal model of mania, leads to a marked increase in CRH [95]. Cortisol levels have also been found to be increased both in the frontal cortex and periphery of rats exposed to prenatal stress, which has relevance in that prenatal stress may predict the development of these disorders. In addition to increased corticosterone release, Ward *et al.* [96] found rats exposed to prenatal stress to also have adrenal hypertrophy with increased expression of CRF-1 receptors in the amygdala [97]. Furthermore, olanzapine treatment was able to reverse the increased cortisol observed in the prefrontal cortex following prenatal stress [98].

Thyroid Hormones

Clinical Correlates

The relation between thyroid dysfunction and mental disorders has long been recognized, ranging from depression [99], anxiety [100] and schizophrenia [101]. A recent study explored thyroid-stimulating hormone (TSH) in patients with acute schizophrenia, unipolar depression, bipolar depression and bipolar mania and, apart from measuring TSH disturbances in all the disorders, observed a definite higher prevalence of thyroid dysfunction in patients with both unipolar and bipolar mood disorders vs. controls [102]. Another study observed significant thyroid dysfunction (hypothyroidism and hyperthyroidism) in schizophrenia as well as BPD patients [88]. Interestingly, autoimmune thyroid disease was more common in schizophrenia [103], emphasising an immune-inflammatory basis for the illness. Santos *et al.* [104] reviewed research on thyroid function in schizophrenia, relating interrelations between the pituitary-thyroid axis and major neuro-signaling systems involved in schizophrenia (including serotonin (5-HT), dopamine (DA), glutamate and GABA networks), as well as myelination and inflammatory processes. These processes are all convergent on the pathology of this disorder, as will be discussed. The authors conclude that thyroid hormone deregulation is a common feature in schizophrenia. Together, these studies emphasize the relevance of thyroid hormonal status as possible biomarkers in MD, BPD and schizophrenia, although further work in this regard is required to establish its putative role as a biomarker.

Pre-Clinical Correlates

Wistar Kyoto rats, an animal model of depression, have increased TSH levels that, together with ACTH, remains elevated after the diurnal peak [80]. To the best of our knowledge, the current body of literature on preclinical BPD and schizophrenia research does not contain significant data on thyroid hormones in animal models of mood and psychotic disorders.

Neurochemical Markers

The majority of drugs used clinically to treat MD, BPD and psychotic disorders such as schizophrenia target monoamine (DA, 5-HT and noradrenaline (NA)) receptors, reuptake transporters and monoamine metabolism [105, 106]. DA-ergic, 5-HT-ergic and/or NA-ergic neurotransmission affects behaviour by regulating motivation, reward seeking, aggression, and activity level – all symptoms that play an important role in the pathophysiology of these disorders [107, 108]. However, the cause of mood disorders is far from being a simple dysregulation of central monoamines. For example, monoamine oxidase inhibitors and monoamine reuptake inhibitors produce immediate increases in monoamine transmission [109], whereas their mood-enhancing properties are only fully realised following 4-6 weeks of sustained treatment. In fact, some patients do not show adequate improvement even after many months of treatment [110]. This indicates that enhanced serotonergic or noradrenergic neurotransmission *per se* is not immediately responsible for the clinical actions of these drugs [111]. Indeed, neurotrophins, neurogenesis and the concepts of neuroplasticity has now taken centre stage in our understanding of mood and psychiatric disorders and the mechanism of action of antidepressants [109, 112, 112a]. Thus, and apart from some data based on NA (see later), selecting an antidepressant based on its monoamine selectivity remains to be substantiated. The same can be said for antipsychotic drugs. Furthermore, a realization that neuroendocrine and metabolic dysfunction also contribute to the eventual development of these disorders, has provided a new framework for understanding their neurobiology and treatment. Nevertheless, their contribution towards the understanding and treatment of these disorders warrants closer scrutiny with respect to viable clinical biomarkers.

Dopamine

Clinical Correlates

Depression

Depressive symptoms (e.g. avolition, guilt, suicidality and social withdrawal) are ascribed to frontal cortical hypodopaminergia [109]. Striatal DA levels in MD are also reduced [113], being linked to symptoms such as anhedonia, reduced motivation and decreased energy levels. Patients presenting with MD episodes have been demonstrated to have significantly decreased dopamine transporter (DAT) binding potential, with binding potential correlating to receptor density and affinity [114]. Anhedonic MD patients exhibit significantly decreased levels of DAT in basal ganglia which are in accordance with the hypothesis linking impaired DA transmission to an impaired reward system [115]. It has also been suggested that decreased striatal D₂ receptor density may underlie depressive symptoms, while increased striatal D₂ receptor density/affinity has been observed in patients after successful SSRI treatment [116] which coincides with evidence of 5-HT modulating DA pathways [117].

Bipolar Disorder

Pharmacological evidence supports evidence that excessive DA neurotransmission mediates manic symptoms in BPD patients [118], while DA receptor D₂ antagonists are robust anti-manic agents [119].

Schizophrenia

The DA hypothesis of schizophrenia proposes that a hyper-dopaminergic state in the striatum mediates positive symptom expression, while a hypo-dopaminergic state in the frontal cortex mediates cognitive and negative symptoms [120]. In line with this hypothesis, post-mortem studies in schizophrenia patients have described frontal cortical hypo-dopaminergia [121] and elevated DA levels in the striatum [120]. However, a previous study reviewing clinical evidence for DA involvement in schizophrenia came to the conclusion that multiple “hits” (i.e. adverse environment, infection, chronic substance abuse etc.) interact to result in DA dysregulation, thereby producing the final common pathway to psychosis in schizophrenia [122]. In fact, MD [123, 124] and BPD [125, 126] are also correlated to early life trauma. It is pathways related to the latter that are deemed critical prodromal events in early life adversity, such as neurotoxicity, oxidative stress and inflammation that may hold the clue to identifying more appropriate biomarkers for these illnesses. Investigators have noted a strong correlation between D₂ receptor binding and response to an antipsychotic, with a minimum 70% receptor occupancy necessary for antipsychotic action [127]. In fact, the success of treatment with antipsychotic agents depend on dopamine D₂ receptor blockade, while Howes and Kapur [122] recently suggested various genetic and environmental factors to be implicated in compromising the brain and ultimately leading to dysregulation of DA.

Preclinical Correlates

Depression

CMS decreases *in vivo* DA release [129] and leads to a decrease in D₂ and D₃ receptors in the limbic forebrain which is reversed by chronic treatment with imipramine [130]. A decrease in DA release in the nucleus accumbens has been observed as well as increased DA levels in limbic regions in FSL rats (together with elevated 5-HT and NA; see below) – possibly due to an increased synthesis and decreased release of DA [131].

Bipolar Disorder

Models of mania which incorporate dopaminergic agents, eg. amphetamine, have been demonstrated to be superior to other similar models [132]. Alpha-methyl-para-tyrosine (AMPT) mediated catecholamine depletion mitigates some mania-related characteristics in DAT knockdown (KD) mice [133], while treatment with valproate reverses locomotor hyperactivity in these animals [134]. Also, treatment with lithium and valproate reverses increased extracellular DA and oxidative damage in a dextro-amphetamine-induced rat model of mania [135]. Furthermore, hyperactive rats exposed to CMS display significantly reduced HVA (homovanillic acid – DA metabolite) compared to DA in the nucleus accumbens, indicating decreased DA release in this brain region [136].

Schizophrenia

Previous evidence on the SIR model has indicated elevated striatal and decreased frontal cortical DA, dihydroxyphenylacetic acid (Dopac) and homovanillic acid

(HVA) levels [137]; increased or decreased frontal cortical DA and unchanged striatal DA [138]; as well as decreased frontal cortical and elevated nucleus accumbens DA, Dopac and HVA [139, 140]. Another study in the SIR model observed reduced PFC D₁ receptor density [141]. However, changes in mesolimbic dopamine D₂ receptor expression are inconsistent, describing down-regulation in striatum [142], but no change in mesolimbic [143], hippocampal, PFC or amygdala areas [144]. SIR also induces a hyper-responsiveness in DA release in the PFC in response to systemic administration of the atypical antipsychotics clozapine and olanzapine, but not haloperidol [145]. Moreover, microdialysis data show that both basal and stress-induced PFC DA levels are reduced in rats chronically treated with PCP [146, 147].

Serotonin (5-HT)

Clinical Correlates

Depression

Although 5HT_{2c} antagonists are ineffective alone in the treatment of MD, they do show benefit when combined with other mood-regulating mechanisms, such as 5-HT reuptake inhibition (SRIs) or melatonin agonism (e.g., agomelatine) [78]. Since 5-HT_{2c} receptor activation inhibits NA and DA release [78], the suppression of these monoamines by elevated 5-HT contradicts traditional views that antidepressant response typically involves an increase in brain 5-HT, as well as NA and DA. Indeed elevated 5-HT-mediated suppression of DA and NA release will be counter-productive [148], such as causing emotional detachment and failure to address the anhedonic symptoms of MD [78]. Clearly there are valid reasons to doubt whether an elevation in brain 5-HT is in any way essential for antidepressant response. In fact, a sustained increase in 5-HT does not appear to be a requirement for anxiolytic/antidepressant effects of an SRI [149]. Further on this point, 5-HT agonists are ineffective as antidepressants while the 5-HT reuptake enhancer, tianeptine, is an effective antidepressant despite having the exact *opposite* effect on synaptic levels of 5-HT than SRIs [150]. This evidence contradicts the simplistic view that brain 5-HT needs to be elevated to improve mood, and has been instrumental in fueling the search for new generation antidepressants.

Post-mortem studies have indicated that suicidal patients with MD present with low cerebrospinal fluid (CSF) levels of 5-hydroxyindole-acetic acid (5-HIAA), the metabolite of 5-HT [151, 152], along with increased 5-HT_{2A} receptor binding sites in platelets [153] and prefrontal cortical sites [154] as well as increased limbic and decreased frontal cortical 5-HT_{1A} receptors (reviewed in [155]). MD patients also present with reduced 5-HT_{2A} receptor density in the frontal cortex [155a]. Interestingly, limbic density and activity of monoamine oxidase (MAO) is elevated in MD [156, 157] which in turn will influence a number of components of monoamine signalling.

Bipolar Disorder

The role of 5-HT in the pathogenesis of BPD is less studied, although a post-mortem study has indicated that

subjects with DSM-III-R diagnoses of BPD who died while depressed had significantly reduced levels of 5-HIAA in frontal (−54%) and parietal cortex (−64%) [158]. A deficit in 5-HT uptake sites has also been observed in the brains of depressed BPD patients after death [159]. Furthermore, 5-HIAA levels were found to be decreased in the CSF of depressed BPD patients [160] and elevated in manic BPD patients [161].

Schizophrenia

Post-mortem studies in patients with schizophrenia [162, 163] as well as psychotic patients [164] have observed reduced frontal cortex 5-HT_{2A} and increased 5-HT_{1A} receptor density. Another study also indicated increased striatal but diminished frontal cortical 5-HT uptake sites in schizophrenia patients [165]. In line with these findings, CSF, genetic and neuroimaging studies have demonstrated an increase in central 5-HT-ergic neurotransmission in schizophrenia [166, 167] and typified by the serotonergic psychedelics such as lysergide. A previous review suggested that the positive symptoms observed in schizophrenia (delusions, hallucinations etc.) could be associated with an excess of 5-HT in the striatum [168]. Despite the above evidence for 5-HT involvement in schizophrenia, clinical studies have found selective 5HT_{2A/2C} antagonists to be ineffective as antipsychotics (reviewed in [169]) and that alterations of brainstem 5-HT transporters are generally not associated with schizophrenia [170].

Preclinical Correlates

Depression

Excessive activation of the 5-HT_{2C} receptor is anxiogenic [171] while 5-HT_{2C} receptor antagonists are rapid acting with sustained anxiolytic actions [172]. 5HT and 5HIAA levels were noted to be higher in limbic structures in the brains of FSL rats compared to normal Sprague Dawley rats [173], while 5-HT_{2/3} receptor density is compromised in the nucleus accumbens leading to a lack of DA-5-HT interaction [174]. Furthermore, CMS leads to an increase in 5HT_{2a} receptors in the cortex which is reversed by imipramine treatment [20]. However, contrary to human subjects, FSL rats present with decreased 5-HT synthesis [175] and SERT-expression [176].

Bipolar Disorder

5-HT-related data are limited in animal models of BPD and mania. However, a mutPOLG transgenic (Tg) mouse model of BPD has been demonstrated to have enhanced 5-HT turnover, accompanied by reduced 5-HT levels, in the amygdala and hippocampus when compared to non-Tg animals [177].

Schizophrenia

Studies on the SIR model has observed decreased cortical (or striatal) 5-HT/5-HIAA [138, 178], decreased frontal cortical and elevated nucleus accumbens and striatal 5-HT and 5-HIAA levels [140, 179]. Deficits in prefrontal 5-HT following SIR is also linked to the behavioural impairments associated with schizophrenia [180]. Evidence of altered 5-HT levels in the NMDA receptor antagonist model is limited

with only one study indicating that 5-HT₃ receptor antagonists can attenuate the behavioural hyperactivity caused by PCP [181].

Noradrenaline (NA)

Clinical Correlates

Depression

NA is of major importance in MD (reviewed in [182]). Previous studies have observed reduced levels of NA transporters in the locus coeruleus [183], altered density and sensitivity of frontal cortical α_{2A} -adrenoceptors [184, 185], and a reduction of NA levels in non-compliant MD patients [186]. Further, a positive relationship between urine NA levels and MD has been confirmed [187]. Symptoms of anxiety were also associated with increased NA excretion in the urine [187]. Moreover, studies have demonstrated that low urinary excretion of the NA metabolite, 3-methoxy-4-hydroxyphenylglycol (MHPG), predict a positive response to NA-selective drugs such as imipramine, nortriptyline, desipramine, or maprotiline [188, 189]. These studies illustrate the significance of urinary noradrenergic measurements as a biomarker in guiding treatment selection and predicting efficacy. Expression of adrenoceptor density has also been investigated in individuals suffering from MD. However, even though dysregulation in alpha and beta-adrenoceptor systems have been noted, it remains unclear whether alterations in the expression of these receptors are causative in the pathology of MD. Considering the heterogeneity of the disorder, the value of adrenoceptor dysregulation as a biomarker is unclear [190].

Bipolar Disorder

NA studies in BPD are limited although an increased turnover of NA has been shown to be central to the pathology of the disorder [158]. Furthermore, post-mortem studies in schizophrenia associated with the positive symptoms of the illness describe elevated brain NA levels as mentioned above [191]. NA has also been shown to be one of the primary neurotransmitters targeted during carbamazepine therapy in BPD patients [192].

Schizophrenia

An earlier review found consistent evidence that the positive and negative symptoms observed in schizophrenia are associated with over-activity and under-activity of central NA, respectively [191]. Moreover, increased NA reactivity and/or tone have been linked to anxiety observed in schizophrenia [193].

Preclinical Correlates

Depression

Data relating to NA as a biomarker in preclinical models of MD are limited, although increased catecholamines, including NA, has been reported in limbic regions in FSL rats [194].

Bipolar Disorder

Interestingly, very little data is currently available in animal models of BPD to support the role of NA as a

biomarker, although a preclinical study has suggested a noradrenergic role for lamotrigine, producing an anti-immobility effect in the mouse forced swimming test (FST) while investigating the depressive facet of the disorder [195].

Schizophrenia

Similarly, evidence in support of NA in a schizophrenia animal model is extremely limited. However, a recent study on the SIR model in our laboratory has demonstrated elevated frontal cortical NA as well as striatal NA and MHPG, with decreased frontal cortical MHPG levels, in SIR rats [179]. Earlier SIR studies found an increase in NA turnover in the hippocampus, cerebellum and cortex of Wistar rats [196].

Glutamate and Gamma-Aminobutyric Acid (GABA)

Glutamate and aspartate, and GABA and glycine, are the preeminent excitatory and inhibitory amino acids respectively, in the brain. Their diffuse presence in interneurons (GABA) or as relay neurons and interneurons (glutamate) allows them to play a profound role in regulating the function of most neurotransmitter systems in the brain. As a result of their ubiquitous presence they are implicated in the neurobiology of probably all central nervous system disorders, in particular MD, BPD and schizophrenia. GABA-glutamate interactions have importance in kindling, a mechanism suggested to underlie the development of rapid cycling of mood or psychotic episodes, and how stressful life events adversely impact long-term outcome. GABA pathways exert a permissive role on the kindling action of glutamate, with excessive glutamatergic activity associated with synaptic remodeling and neurodegeneration.

Clinical Correlates

Depression

Abnormalities resulting in an increase in glutamate transmission have been reported in patients with MD [197]. Elevated levels of glutamate act on extrasynaptic NMDA receptors leading to an influx of Ca^{2+} into the neurons, which results in the toxic accumulation of reactive oxygen species (ROS) [198], with increased nitric oxide (NO) production playing a key role in MD pathology and treatment response [199, 200]. We have earlier proposed that the NO pathway may play an important role in relapse and treatment resistance [201] as well as influencing the effect of non-compliance on treatment outcome [202]. In MD, glutamatergic hyper-function seems to be closely related to the lack of 5-HT-ergic and noradrenergic neurotransmission noted to underlie the core symptoms of MD. Indeed, studies examining peripheral blood of MD patients have demonstrated the glutamatergic system to be overly activated [203, 204]. Elevated glutamate levels have also been found in the occipital cortex of un-medicated subjects with MD [205]. Accordingly, reduced glycine binding (where it acts to abrogate NMDA receptor activity) has been described in the frontal and temporal cortex of suicide victims and MD patients [206, 207], leading to hyperglutamatergia. Glutamate, in combination with quinolinic acid (QA), a glutamate agonist derived from the kynurenic pathway (see later), may

contribute to excitotoxicity in the central nervous system [208]. While several factors may influence the levels of kynurenic acid and its metabolites (eg. inflammation), a decrease in tryptophan (TRP) may generally be observed in patients suffering from MD resulting in reduced 5-HT levels. In general, depression is associated with lowered TRP, increased indoleamine 2,3 dioxygenase (IDO) activity as well as reduced levels of kynurenic acid [209, 210]. Furthermore, microglial levels of QA have been demonstrated to be upregulated in MD [211].

Bipolar Disorder

A recent review of magnetic resonance spectroscopy (MRS) studies in patients with BPD observed the cingulate and prefrontal cortices to contain higher glutamate levels, and possibly associated with illness state [212], while a decrease in NMDA receptor binding has also been noted in the CA3 region of the hippocampus [213]. In a post-mortem morphological study, an increase in QA positive microglia has been observed in the subgenual anterior cingulate cortex of BPD patients, commensurate with increased glutamatergic activity [211]. Drug therapy with the pyrimidine compound, cytidine, reduces glutamine/glutamate levels in BPD and possibly related to symptom severity, suggesting that the presence of glutamatergic dysfunction is an important factor in the underlying pathology of BPD [214]. Furthermore, the presence of genetic mutations affecting the glutamate pathway has also been suggested to be implicated in BPD [215].

Schizophrenia

Release of DA from cortical and limbic striatal structures are controlled by glutamate-GABA-glutamate feedback loops situated on pyramidal cells of the frontal cortex, the disturbances of which underlie the behavioral manifestations of schizophrenia [216, 217]. GABA'ergic interneurons in the brain stem monoaminergic nuclei, viz. raphe nuclei, locus coeruleus, ventral tegmentum, also modulate ascending serotonergic, noradrenergic and dopaminergic pathways, resulting in tonic *inhibition* of NA and DA release in the PFC, resulting in the emotional, mood and cognitive deficits associated with MD and schizophrenia [78].

The "glutamate" hypothesis of schizophrenia emerged from the observation that NMDA receptor inhibition induces schizophrenia-like behaviors in humans. Cortical hypoglutamatergia compromises DA release in the ventral tegmentum leading to meso-limbic hyperdopaminergic (positive symptoms) and meso-cortical hypodopaminergia (negative symptoms) [217]. Mitochondrial dysfunction, pro-inflammatory cytokines and increased IDO-mediated conversion of tryptophan to QA (supported by clinical evidence for elevated QA [218]), the latter an NMDA receptor agonist, may be directly or indirectly implicated in eliciting glutamate hyperactivity thereby increasing NMDA receptor activation, altered redox balance and oxidative stress [217]. Schizophrenia has also been likened to the kindling phenomenon, a process of increased excitatory glutamatergic activity coupled with a relative loss of inhibitory GABA'ergic tone [217].

In post-mortem schizophrenia studies, deficits of glutamate systems have been described in the temporal cortex, medial temporal lobe and striatal regions [219, 220], together with losses of glutamate uptake sites [221] and increases in NMDA receptors in the same brain regions [207]. Previous studies also emphasize the impact of NO metabolism *via* glutamate and GABA on NMDA receptor mediated neurotransmission in schizophrenia [222, 223]. NO is an important second messenger for the glutamate NMDA receptor pathway, and its overproduction is implicated in schizophrenia. Excessive NO release include impairment of NMDA-receptor mediated neurotransmission, disturbed DA metabolism, excessive ROS generation and mitochondrial dysfunction with cell-death (reviewed in [222, 223]). However, altered NO metabolism is not unique to, or indicative of, schizophrenia as disturbances in this signalling cascade has been noted in MD and BPD [222], as noted earlier. A recent clinical study also indicated elevated GABA and glutamate levels in the medial prefrontal cortex of un-medicated patients, with no alterations in medicated schizophrenia patients, suggesting possible normalization of GABA and glutamate with antipsychotic treatment [224].

Preclinical Correlates

Depression

Under conditions of chronic stress, elevated glucocorticoid levels enhance glutamatergic transmission by increasing the expression of the glutamate ionotropic NMDA receptors, as well as increasing the synthesis and extracellular concentrations of glutamate [225]. Abnormalities resulting in an increase in glutamate-NO transmission have been reported in FSL rats [226]. The possible mechanisms whereby NO can contribute to mood disorders is obscure, although persistent research has highlighted various possibilities including the actions of the NO/cyclic guanosine 3'5'-monophosphate (cGMP) pathway. Modulators of the NO-pathway have also gained relevance in MD research due to NO-inhibitors demonstrating antidepressant effects in models predictive of antidepressant activity [227, 228]. By activating soluble guanylate cyclase (sGC) which converts guanosine 5'-triphosphate (GTP) to the intracellular messenger cGMP [229, 230], NO is enabled to mediate many cellular processes, particularly the regulation of ion channels, activation and inhibition of cyclic nucleotide hydrolysis by phosphodiesterase, activation of G-kinase and modulation of neurotransmitter release [229, 231]. Interestingly, neuronal nitric oxide synthase (nNOS) inhibitors (eg. methylene blue) [232] as well as PDE₅ inhibitors (e.g. sildenafil) [233] have antidepressant and anxiolytic effects in animal models [234, 235] (Fig. 1), as do clinically relevant antidepressants [228]. These effects however are due to interference with constitutive nNOS-mediated effects and *not* inducible NOS (iNOS), which rather plays an important role in chronic stress associated with inflammation [231] (Fig. 1). Stressed vs. naive FSL/FRL rats show elevated hippocampal glutamate-NO signalling [224], indicating that a chronic environmental stressor is required in order to demonstrate altered glutamate-NO signalling in FSL rats [236]. This is congruent with the fact that MD involves a

prior and/or ongoing chronic stressor [237, 238]. Considering these findings, it is not surprising that NMDA antagonists such as dizocilpine (MK-801) [239], ketamine [240], memantine [241] and others [242] exert antidepressant effects, while disinhibition of glutamate-NO signalling follows antidepressant discontinuation after chronic treatment [243, 244]. A possible explanation could be that NMDA receptor antagonists increase 5-HT levels in the brain [245], while also having a modulatory effect on pathways involved in neuroplasticity and cellular resilience [202].

Preliminary evidence also supports the use of NMDA antagonists such as ketamine in treatment-resistant MD [246]. How this happens still needs illumination, although animal studies have begun to delve into the possible mechanisms involved [247]. The latter work has indicated a mutual cooperation with glutamate AMPA receptors [247], resulting in activation of mammalian target of rapamycin (mTOR) [248] and inhibition of glycogen synthase kinase-3 β (GSK-3 β) [249] signalling (Fig. 2). The mTOR pathway plays a pivotal role in protein synthesis by stimulating mRNA translation *via* interaction with its downstream targets [248], and leads to prolonged elevation of synapse-associated proteins in the prefrontal cortex [248]. Diminished activity of the mTOR pathway could underlie synaptic deficits in the PFC as previously reported in MD [250]. Furthermore, this evidence is supported by the behavioural responses to ketamine being blocked in mice which express constitutively active GSK-3 β [251]. Considering the contributory role of oxidative stress in MD, inactivation of GSK-3 β is linked to the regulation of redox homeostasis *via* stress responsive genes that protect cells against inflammation and oxidative stress [252, 253] (Fig. 2).

Bipolar Disorder

The standard treatment for BPD, lithium salts, target the glutamate-NO system [254, 255]. Unfortunately, current literature lacks sufficient data to elaborate on the role of glutamate/GABA as a biomarker in preclinical models of mania and BPD.

Schizophrenia

Decreased glutamate release has been observed in the frontal cortices of Homer1 mutant mice, a putative animal model of schizophrenia [256], while chronic phencyclidine (PCP) administration in rats is associated with a decreased expression of glutamate receptors in the prefrontal cortex [257] and a reduced number of cortical and hippocampal PV-immunoreactive neurones [64]. Confirming this, partial deletion of the NMDA receptor in mice is associated with behavioural alterations akin to that observed in PCP treated mice [258], while increased NMDA receptor binding has been described in the frontal cortex of SIR animals [141].

Neuronal Growth Factors

Growth factors are intricately involved in the survival, growth and differentiation of specific groups of neurons. Their relevance is gaining in importance in the light of

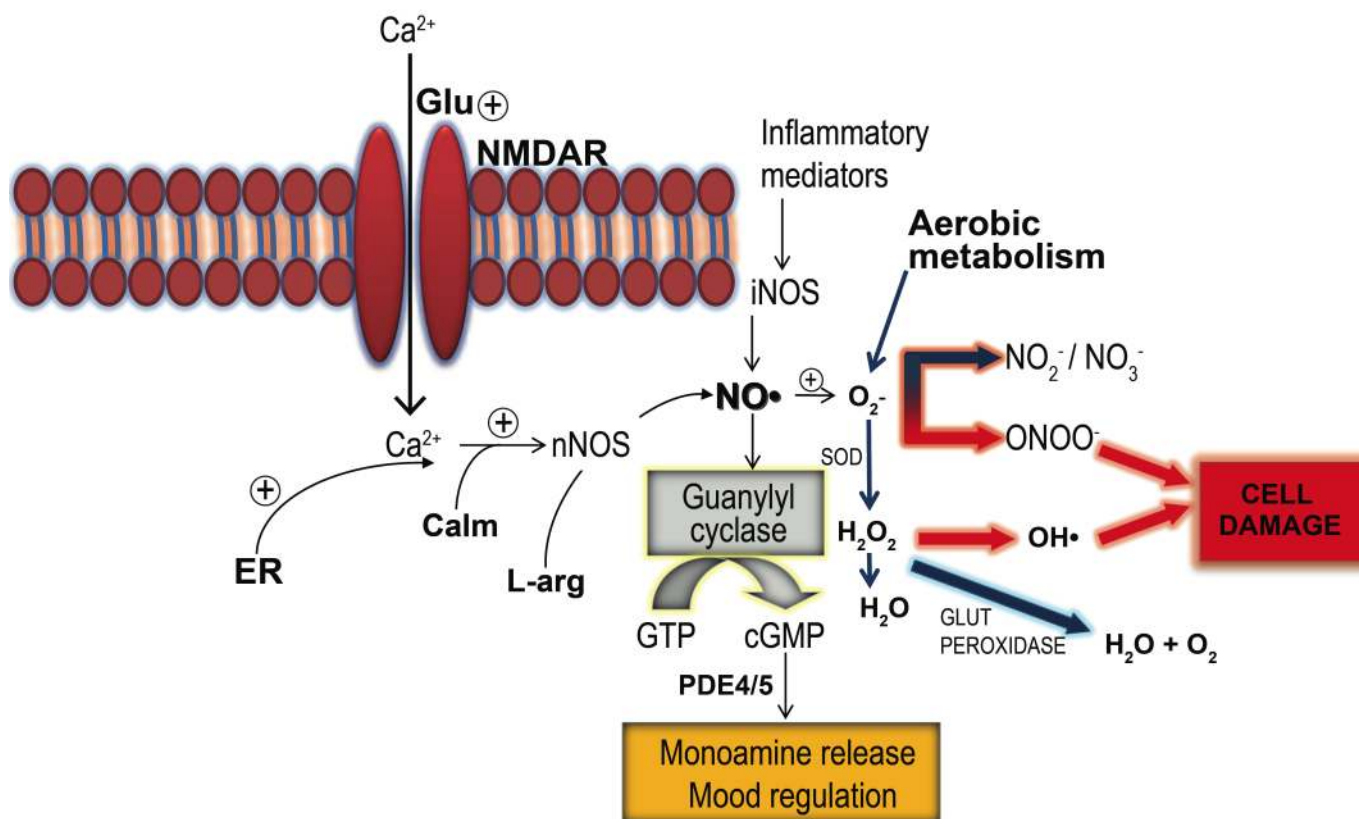


Fig. (1). Glutamate-mediated effects on the cGMP-NO system leading to monoamine release that in turn can be targeted by pharmacological means, eg. PDE5 and NOS inhibitors, as well as known antidepressants. However this pathway can also lead to oxidative stress if excessive glutamate-mediated NO synthesis combines with O_2^- from aerobic metabolism. Also depicted is the effect of inflammatory mediators that promote iNOS-mediated NO synthesis thereby promoting the formation of cell-damaging reactive oxygen and nitrogen species. These pro-oxidative mechanisms can be abrogated by endogenous antioxidant systems such as superoxide dismutase (SOD) and glutathione that act as a sink to quench excessive NO and/or O_2^- .

Abbreviations: calmodulin (Calm); cyclic guanosine monophosphate (cGMP); endoplasmic reticulum (ER); glutamate (Glu); glutathione peroxidase (glut peroxidase); guanosine triphosphate (GTP); inducible nitric oxide synthase (iNOS); l-arginine (L-arg); neuronal nitric oxide synthase (nNOS); nitric oxide (NO); NMDA receptor (NMDAR); phosphodiesterase (PDE); superoxide dismutase (SOD); superoxide (O_2^-); hydrogen peroxide (H_2O_2).

increasing evidence that mood and psychotic disorders are associated with structural brain changes and that alterations in growth factors may precipitate or exacerbate depressive, BPD and psychotic episodes [13, 259, 260].

Brain-Derived Neurotrophic Factor (BDNF)

Clinical Correlates

Depression

Extensive studies have established that altered BDNF plays a pivotal role in MD. BDNF and the transcription factor, cyclic adenosine monophosphate (cAMP) response element binding protein (CREB), are intimately linked biochemically (see Fig. 2), playing a critical role in cellular resilience and neuroplasticity. Antidepressant treatment up-regulates CREB in the cortex and hippocampus of humans [261]. Both serum BDNF levels and CREB phosphorylation and protein levels are reduced in depressed individuals [262]. Moreover, an inverse relationship exists between serum levels of BDNF and the severity of MD [263], while

antidepressant treatment is able to reverse the deficit in BDNF observed in MD [264, 265] and to increase phosphorylation and binding of CREB [266, 267]. BDNF is expressed throughout the body [268], but the exact origins of circulating BDNF remain elusive. BDNF has been shown to originate from several sources including brain neurons, vascular endothelial cells and platelets. It has also been shown to cross the blood-brain barrier [269] so that plasma BDNF levels may reflect central BDNF levels [270]. BDNF regulates synaptic plasticity in neuronal networks and appears to be a particularly relevant factor for mood disorders with associated cognitive dysfunction [271-273].

CREB is responsible for regulating BDNF expression [274]. Activation of CREB is associated with the regulation of synaptic plasticity as well as transcription of specific target genes involved in the production of proteins, BDNF being one example [275, 276]. Post-mortem studies have reported decreased hippocampal BDNF in MD patients who committed suicide, but elevated levels in patients who were being treated with antidepressant agents at the time of death

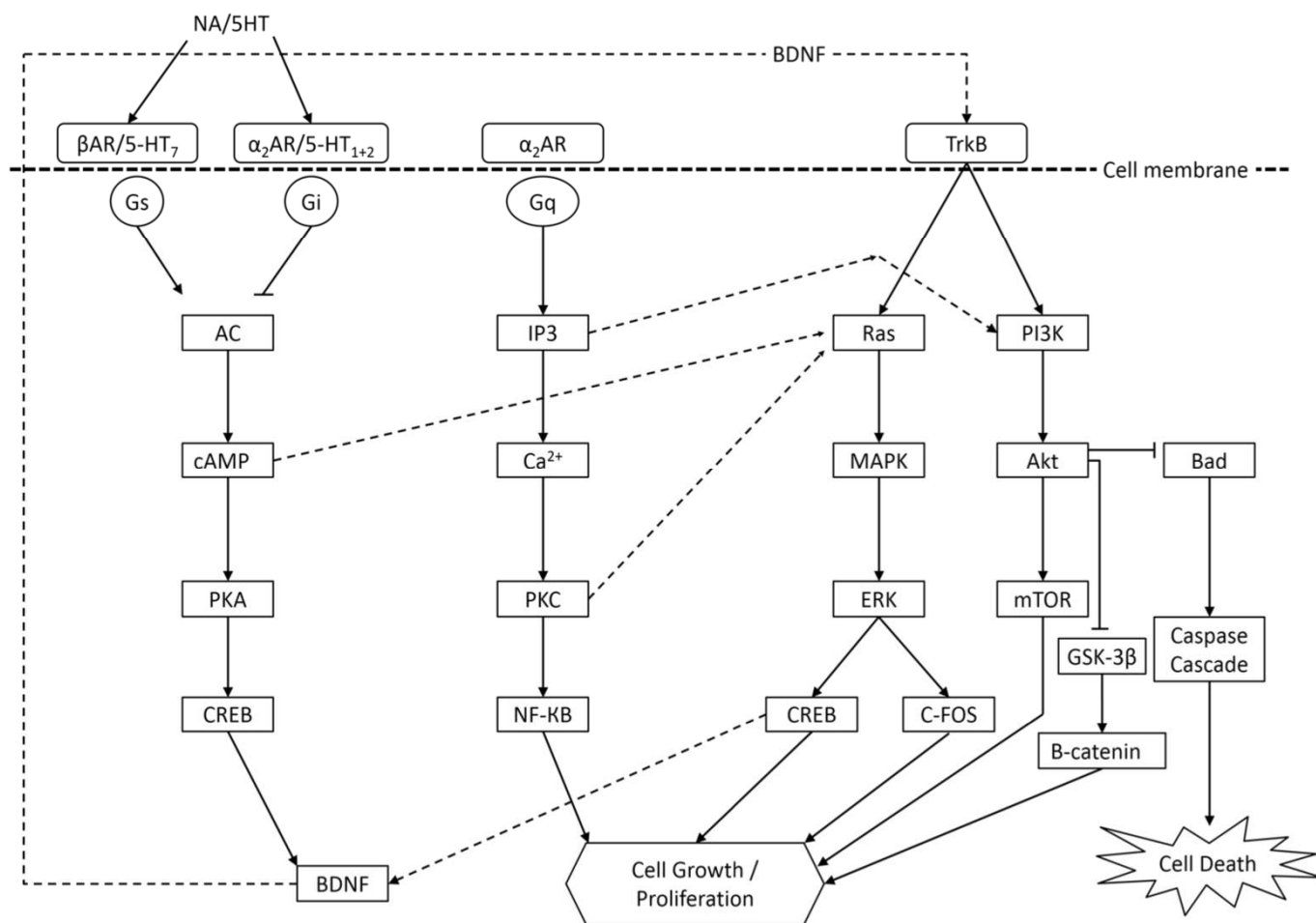


Fig. (2). An overview of key neuroprotective and neurotoxic molecules involved in drug (antidepressant, lithium etc.)-induced neuroplasticity. The monoaminergic system mainly exerts its effect on BDNF expression *via* the cAMP cascade, while BDNF in turn exerts its effect on monoaminergic neurons through the TrkB-receptor *via* the MAPK/ERK cascade and the phospholipase C (PLC) signalling system. Adapted from [342]. Abbreviations: beta adrenoceptor (β AR); alpha-2 adrenoceptor (α 2AR); BDNF receptor (TrkB), see text for further details/ abbreviations not noted here.

[268, 277]. Considering the growing evidence for an interaction between MD and metabolic and redox-related conditions [278-280], our group recently showed that altered serum BDNF may be linked to metabolic and redox factors, with BDNF levels indicating either a counter-regulatory action on the effects of glutathione oxidation or that BDNF may mediate the redox effects itself, leading to the development of a mood disorder [281].

Various factors associated with an increased risk of developing MD, e.g. smoking [282, 283] and type II diabetes mellitus [284], have been linked to BDNF deficits. Thus decreased levels of BDNF have been found in smoking individuals when compared to non-smokers [285, 286], while smoking cessation leads to improved BDNF levels [285, 286]. Likewise, serum levels of BDNF have been shown to be significantly lower in subjects with Type II diabetes when compared to healthy controls [287, 288], while cerebral output of BDNF is inhibited in the presence of high blood glucose levels [287]. These findings reiterate the

causal link between metabolic diseases, altered BDNF and the development of MD noted earlier. On the other hand, for example, physical exercise has been shown to increase BDNF [289, 290], to be neuroprotective, to improve mood and to have antidepressant effects [291, 292].

Bipolar Disorder

Decreased peripheral BDNF levels have been observed in BPD patients, possibly associated with the pathophysiology and severity of manic symptoms [293]. Exercise has also been shown to decrease depressive symptoms in BPD patients and to even increase the frequency of mania [294]. The latter would indicate that elevated levels of BDNF may not always be beneficial, as has been proposed in a study in patients with MD [281]. The authors suggest that, by adversely affecting resilience, BDNF facilitates activity-dependent plasticity that may translate to a variable effect on mood and other plasticity-dependent functions. In fact, BDNF has been noted to induce paradoxical depressogenic effects [281]. As a mediator of synaptic plasticity, maladaptive

secretion of BDNF (eg. a response to environmental adversity) may set in motion counterregulatory responses that are counterproductive.

Schizophrenia

Decreased peripheral BDNF levels have been observed in schizophrenia [295]. Importantly, a recent study indicated that clinically stable schizophrenia patients present with significantly increased serum levels of BDNF after exposure to cognitive training targeted at improving neuroplasticity [296]. Furthermore, post-mortem studies reported a decreased concentration of BDNF-positive neurons [297] and BDNF concentrations in brain tissue of schizophrenic patients, which include the cortical areas and the hippocampi [298].

Preclinical Correlates

Depression

Antidepressant treatment up-regulates CREB in the cortex and hippocampus of rats [299], while an overexpression of CREB in the dentate gyrus results in antidepressant effects in the FST and a learned helplessness paradigm – both animal models of MD [300]. In the latter, decreased hippocampal BDNF levels were described. Although contrary to that in human subjects, some studies have also noted increased serum levels of BDNF [301, 302]. In FSL rats, serum and whole blood BDNF levels have been found to be significantly increased compared to control but significantly decreased in the hippocampus, with no differences noted in the frontal cortex and CSF [303], suggesting that BDNF is differentially regulated in hippocampus, serum, and whole blood in these animals. The latter is not unlike similar paradoxical findings in humans, where BDNF has been suggested to play a counter-regulatory role [281]. Preclinical studies have indicated that BDNF administration produces antidepressant-like behaviour [304], while antidepressants and electroconvulsive therapy increase BDNF levels [305]. After animals were subjected to repeated stress, they constantly presented with decreased BDNF levels as measured in the hippocampus and serum, while corticosterone levels returned to normal levels, suggesting that changes in brain plasticity occur following a second stressful event [306]. The presence of decreased serum BDNF levels accompanied by normal serum cortisol levels may therefore represent a relevant biomarker for identifying individuals who are more likely to develop depressive symptoms in the subset of a population which may be predisposed to developing affective disorders. These alterations may even be expanded to other disorders provoked by stressful life events, for instance schizophrenia [307].

With smoking having been found to affect BDNF levels in humans, decreased levels of BDNF have also been found in rats repeatedly exposed to nicotine [308]. Similarly, physical exercise also increases BDNF in animals [289, 290].

Bipolar Disorder

BDNF levels are decreased in both the amygdala and hippocampus of rats in the ouabain model of mania, and reversed by lithium [309]. Moreover, in an amphetamine-induced model of mania, BDNF was also decreased in the hippocampus and increased by valproate and lithium [119].

Schizophrenia

Neonatal PCP administration produced a sustained elevation of BDNF in the hippocampus and the entorhinal cortex of 8-week-old rats [310]. However, studies in the SIR model observed significantly reduced medial PFC BDNF levels [311] as well as decreased hippocampal BDNF [312].

Insulin-Like Growth Factor

Insulin-like growth factor-1 (IGF-1) is involved in regulating peripheral cell growth and metabolism [313]; and plays a crucial role in the growth and differentiation of nerves and also in the synthesis and release of neurotransmitters [314].

Clinical Correlates

Depression

Unfortunately, there have not been sufficient clinical studies to determine whether peripheral IGF-1 is altered in MD patients or following antidepressant administration.

Bipolar Disorder

In BPD patients, a previous study observed altered IGF signalling in post-mortem brain tissue [315].

Schizophrenia

Antipsychotic-free schizophrenia patients have been found to present with a decrease in plasma IGF levels [316].

Preclinical Correlates

Depression

Unfortunately, we are not aware of any extensive IGF-related data in established animal models of MD. Nevertheless, preclinical studies have indicated that peripheral IGF-1 administration reduces immobility in the FST [317], increases central BDNF mRNA [318] and produces antidepressant-like behavioural responses in mice exposed to CMS [317]. Moreover, after chronically treating rats with antidepressants, elevated IGF-1 expression was observed in the brains of these animals [319]. Finally, IGF-1 has been found to regulate adult hippocampal neurogenesis in rats [320].

Bipolar Disorder, Schizophrenia

To the best of our knowledge there is no pre-clinical data in established animal models of IGF-1 as a preclinical biomarker in BPD or schizophrenia.

Vascular Endothelial Growth Factor

Vascular endothelial growth factor (VEGF) acts as a neurotrophic factor, is a cytokine implicated in angiogenesis [321] and has been related to the vascular niche hypothesis of adult neurogenesis [322]. This hypothesis attributes increases in the proliferation of neurons in the adult hippocampus to VEGF-induced angiogenesis. VEGF is purported to play a role in several features associated with neuronal growth, including neuronal regeneration and differentiation as well as axonal outgrowth [323].

Clinical Correlates

Depression

MD patients present with higher plasma VEGF levels which can be reversed with antidepressant treatment [324], while earlier studies have confirmed said increase in MD [325]. Furthermore, remitted MD patients have significantly elevated VEGF levels, while MD patients with a family history of psychiatric disorders also have higher baseline levels of VEGF, compared to MD patients without a family history and healthy controls [326]. This may be indicative of a role for VEGF in the *pathology* of MD, possibly hinting of a neuroprotective role to counter reduced neurogenesis in MD.

Bipolar Disorder

BPD patients present with higher plasma VEGF levels during acute episodes vs. healthy controls [327], emphasizing that a depressive and manic episode in mood disorders may be associated with the neuroprotective role of VEGF. Interestingly, a recent study indicated that VEGF mRNA levels were significantly decreased in BPD patients treated with lithium vs. healthy controls [328], suggesting that VEGF may be a useful marker in BPD and as an indicator of lithium response.

Schizophrenia

VEGF data in schizophrenia patients are limited, although previous studies have not observed any differences in serum VEGF in schizophrenia vs. healthy individuals [285]. However, significantly reduced levels of VEGF mRNA have been observed in the DLPFC of patients with schizophrenia [329].

Preclinical Correlates

Depression

The relationship between central and peripheral levels of VEGF still needs clarification. In a genetic rat model of MD, Elfving and colleagues found decreased levels of VEGF in the brain but no variations in serum VEGF levels [301].

Bipolar Disorder

To the best of our knowledge, there is no data available on VEGF as a biomarker in preclinical models of BPD.

Schizophrenia

Similarly, no pre-clinical data is currently available on VEGF in the SIR or NMDA antagonist models of schizophrenia. However, a pre-clinical study did observe that VEGF levels are increased in rat hippocampi following 14 days haloperidol or olanzapine treatment [303]. Interestingly, in the case of haloperidol treatment this increase was lost 45 days later, while olanzapine treatment bolstered the initial increase in VEGF [330], reaffirming that first and second generation antipsychotics are not therapeutically equivalent. This underlines VEGF as a possible marker in schizophrenia treatment but not diagnosis *per se*.

Neuronal Resilience Markers

Several neurochemical markers have been associated with neuroprotective effects and positive antidepressant

treatment response. With the increased evidence for a neurodegenerative profile for MD, BPD and schizophrenia and the progressive nature of these illnesses, identifying neuroresilience markers is gaining in relevance. In this regard, resilience markers linked to the BDNF pathway are especially attractive.

Stress and environmental adversity is a common thread throughout all three illnesses under review [252, 331, 332]. Stress-induced increases in glucocorticoid levels have been shown to decrease the synthesis of neurotrophic factors, particularly BDNF, which is an effective neuroprotective factor and protagonist of neurogenesis [333]. These neurotrophic effects are mainly mediated through inhibition of cell death pathways and activation of mitogen-activated protein kinases (originally extracellular signal-regulated kinases or MAPK/ERK) signalling pathways and phosphatidylinositol-3 kinase (PI-3K)/Akt (protein kinase B) pathways (see Fig. 2) [334]. As noted earlier, BDNF expression is decreased during MD, BPD and schizophrenia, a response that is reversed by effective pharmacological treatment [295, 335-337]. Furthermore, increased structural atrophy observed in treatment resistant MD has been correlated with greater decreases in BDNF levels [338] in patients failing to respond to SRI treatment compared to treatment responsive patients.

The cAMP cascade is activated following increased serotonergic and adrenergic receptor activity which results in downstream activation of CREB [339]. The ensuing elevation in cAMP ultimately leads to increased BDNF expression which subsequently activates the MAPK/ERK pathway, a major pathway involved in cell growth and proliferation [340, 341] (see Fig. 2). Monoaminergic neurons experience an increase in growth following MAPK/ERK pathway activation thereby accounting for how BDNF modulates the monoaminergic system [342].

Activation of the PI-3K cascade by BDNF leads to phosphorylation of Akt, a molecule at the crossroad of cell survival and cell death [343] (see Fig. 2). Activation of Akt following phosphorylation leads to enhanced activity of mTOR which is responsible for regulating the expression of several genes involved in cell growth, particularly a group of synapse-associated genes that have been directly linked to neuroplastic events [248]. Conversely, inactivation of Akt by dephosphorylation leads to a decrease in phosphorylation and subsequent activation of Bcl-xL/Bcl-2-associated death promoter (Bad), a pro-apoptotic molecule [343]. It is therefore evident that in combination, the MAPK/ERK and PI-3K pathways are largely accountable for the neuroplastic events occurring during antidepressant response and, furthermore, directly links the actions of BDNF to these processes.

Clinical Correlates

Depression

Decreased cAMP levels and lower MAPK/ERK pathway activity has been associated with MD, which has been shown to be reversed by increasing BDNF levels [344].

Bipolar Disorder

GSK-3 β has been demonstrated to be an important role-player in BPD with lithium, an inhibitor of GSK-3 β , having served as a mainstay in the treatment of BPD. It also regulates various proteins and is involved in neuroplasticity and neurotransmission [345]. Therefore, agents involved in the modulation of GSK-3 β and its downstream pathways may serve as valuable biomarkers in the diagnosis and treatment of BPD – e.g., several molecules involved in both cell survival and apoptosis, such as CREB [346] and p53 [347], respectively. GSK-3 β also plays an important role in the regulation of the Wnt [347] and PI-3K [348] signalling pathways linked to cellular resilience [348, 349] (Fig. 2). It has also been suggested that progranulin (PGRN) may serve as a neurotrophic factor- modulating neurite outgrowth as well as neuronal differentiation and survival [350]. Furthermore, plasma levels of PGRN are decreased in BPD patients [351, 352] and GSK-3 β has been implicated in mediating PGRN activity [351]. GSK-3 β protein expression is decreased in the platelets of BPD patients [349]. Even though previous studies could not find alterations in brain expression of GSK-3 β , decreased protein expression in platelets can be reversed by mood stabilizers – but not antidepressants – thereby emphasizing a valuable role for GSK-3 β as a peripheral biomarker and even a state – rather than trait – marker of BPD [353].

Schizophrenia

A recent post-mortem study in patients with schizophrenia found increased levels of various proteins involved in the MAPK- and cAMP-associated pathways, as expressed in frontal cortical structures [354]. In line with these observations are studies indicating alterations in several proteins in the MAPK-associated pathway: extracellular signal-regulated kinase (ERK)-2, immediate early genes c-fos and c-Jun levels were elevated in the thalamus on both protein and transcription level, whereas c-Jun protein and Elk-1, CREB, and ATF-2 protein levels were elevated in the cerebellar vermis [355, 356]. Moreover, other proteins involved in the MAPK pathway, including MEK1, MEK2, RSK1, B-Raf, and CREB were found to be reduced in the frontal cortex of schizophrenia patients [357]. With regards to the cAMP pathways, decreased DA- and cAMP-regulated phosphoprotein Mr 32 kDa (DARPP-32) was observed in the frontal cortex and thalamus of schizophrenia patients [358, 359]. Furthermore, a recent review highlights numerous evidence and theories in support of a novel mTOR based hypothesis of the neuropathology of schizophrenia [360]. Control of protein synthesis is the primary role of this signalling cascade while it is also regulated by known extracellular and environmental factors implicated in the pathology of schizophrenia [360].

Preclinical Correlates

Depression

Blocking MAPK signalling leads to depressive-like behaviour in the FST in rats and inhibits the antidepressant effects of ketamine [340]. These findings provide some insight on how glutamate-NMDA signalling interacts with

monoaminergic-cAMP pathways to mitigate a faster onset of action or to treat refractory MD. Furthermore, SIR in rats, a putative neurodevelopmental animal model of MD and schizophrenia [18], leads to an enhanced expression of mitogen-activated kinase phosphatase and apoptosis-related genes in the prefrontal cortices of Sprague-Dawley rats [361].

Bipolar Disorder

Transgenic mice that overexpress GSK-3 present with decreased habituation and an increase in activity that has been related to hyperactivity in mania [362]. β -catenin, a downstream molecule of GSK-3 (Fig. 2), was found to be decreased in the hippocampi of black Swiss mice, a putative model of mania [363], while the behaviour of transgenic mice overexpressing β -catenin was found to have a behavioural phenotype similar to that of lithium-treated animals [364]. Of significance is that lithium stabilizes β -catenin by inhibiting GSK-3 β thus reducing neuronal vulnerability to apoptosis [365].

Schizophrenia

Gururajan and Van den Buuse [360] have explained the involvement of mTOR in schizophrenia by referring to numerous animal models of schizophrenia. However, direct measurement of neuronal resilience markers in the SIR model and the NMDA antagonist model is limited, with only one study indicating that MK-801 administration elevates phosphorylation of MAPK in the frontal cortex of rats [19].

Oxidative Stress Markers

Normal oxidative metabolism in cells results in the production of various ROS. Oxidative stress occurs when cellular antioxidant defence mechanisms, such as SOD, catalase, glutathione peroxidase, fail to counterbalance and control endogenous production of ROS such as O₂⁻ and H₂O₂. This leads to a free radical attack on proteins, DNA and lipids [366, 367]. SOD is the primary defense against oxidative stress by converting O₂⁻ to H₂O₂ [368]. Hydrogen peroxide in turn is converted to water and glutathione (GSSG) by catalase and glutathione peroxidase [369], with GSSG rapidly being converted to reduced glutathione (GSH) by glutathione reductase [370]. The brain has relatively low levels of antioxidant defences, as well as a high lipid content that is highly susceptible to attack by ROS [371]. Thus, a reduction in GSH, and an increase in GSSG, is regarded as being indicative of increased oxidative stress.

Many of the changes in oxidative status may be directly related to increased inflammatory response due to the presence of other systemic illnesses, such as endocrine and metabolic disorders and cardiovascular disorders [281]. Furthermore, changes in certain neurotransmitter systems in the brain, especially glutamate and GABA, increase the risk of oxidative stress in the brain and subsequent neuronal oxidation and cell death [279]. Moreover, oxidative stress in its own right may mediate altered monoaminergic activity [372] that underlies the pathology of many neuropsychiatric illnesses associated with oxidative stress [373]. One of the more prominent redox active molecules released by changes in glutamate activity in the brain is NO, and which is well

described as being a contributing factor towards the development of MD [374], schizophrenia [223] and possibly BPD [375]. In this regard, both constitutive NOS-, such as nNOS, and iNOS-mediated NO synthesis needs to be considered, with nNOS being more involved in neurotransmission and iNOS in inflammation. Fig. 1 provides an outline of how glutamate, NO and redox systems interact to produce oxidative stress.

Clinical Correlates

Depression

In recent years MD has been associated with several changes in redox status, presenting as either an increase in oxidative stress and/or diminished oxidative defence systems [279]. Elevated plasma malondialdehyde (an indication of lipid peroxidation) levels and susceptibility of red blood cells to oxidation, as well as an increase in serum SOD activity, has been observed in MD patients [376, 377]. However, Srivastava *et al.* [378] found no alterations in the activities of SOD and glutathione peroxidation in polymorphonuclear leukocytes from patients with MD. In their clinical study, Berk and colleagues noted only limited support for the role of antioxidant and glutathione precursor, N-acetyl cysteine (NAC), as an adjunctive therapy for MD, although further such clinical studies are required [379]. A high incidence of co-morbid metabolic syndrome and MD have been observed [380] with inflammation a major mediator in the development of both MD and metabolic syndrome [381]. In support of this, substantial evidence exists linking insulin- and NO-mediated pathways in the brain. In fact, insulin upregulates expression of nNOS [382] while a role for increased NO and insulin/peroxisome proliferator-activated receptor (PPAR) signalling has been noted following stress, thus presenting as a susceptibility factor in the subsequent development of MD [383].

Bipolar Disorder

BPD patients present with changes in antioxidant enzymes, for example Andreazza *et al.* [384] reported manic and depressive phases to be associated with increased SOD activity, but unaltered activity in euthymia. This is corroborated in part by Machado-Vieira [260] who found untreated manic bipolar patients to present with increased activity of SOD. Furthermore, patients who were euthymic presented with decreased catalase activity [384], while activity was increased in manic patients who did not receive treatment [260]. Increases in lipid peroxidation due to oxidative stress unrelated to the phase of illness have also been reported [260, 384]. In addition, BPD patients were found to express increased lipid peroxidation in the cingulate cortex [385], while clinical studies have indicated that the antioxidant, NAC, is effective as adjunctive treatment in BPD [386-388].

Schizophrenia

Evidence has accumulated in recent years that antioxidant systems are impaired in schizophrenia [389]. Gawryluk *et al.* [390] reported reduced levels of GSH in post-mortem prefrontal cortices of patients with schizophrenia. Do and colleagues [391] found a 52% decrease in GSH

levels in the prefrontal cortex of patients with schizophrenia. Interestingly, a significant deficit in total antioxidant status was inversely associated with some domains of cognitive deficits in schizophrenia patients, such as attention and immediate memory [392]. Moreover, plasma SOD activity was negatively correlated with positive symptoms in first-episode schizophrenia patients [393]. Lower levels of total antioxidant status, catalase and glutathione peroxidation has been described in first episode schizophrenia patients, with GSH levels positively associated with executive function [394]. Furthermore, clinical studies have described the clinical utility of NAC as an adjunctive treatment in schizophrenia [387, 395], as well as the combination of ω -3 fatty acids with vitamins E and C [396]. However, we have been unable to demonstrate efficacy for ω -3 polyunsaturated fatty acids (PUFA) plus alpha-lipoic acid in preventing relapse in patients who had responded well to antipsychotic treatment after a single episode of psychosis [397]. Further studies in this regard are nevertheless warranted.

Preclinical Correlates

Depression

With regard to significant animal studies, SOD and catalase activities have been found to be decreased in rats exposed to a chronic stress model of MD, and could be reversed by tianeptine [398]. Similarly, animal studies have found NAC to be as effective an antidepressant as imipramine [399]. Of particular interest is that exposure to ozone worsens anxiety, cognitive and depressive-like behaviour in the FSL rat model of MD, suggesting that genetically susceptible individuals exposed to high levels of oxidative stress are at higher risk of developing mood and/or anxiety disorders [400]. Moreover, exacerbating levels of oxidative stress (eg. with ozone) can attenuate antidepressant action [400]. In fact, stress-related activation of the NMDA-NOS cascade has been proposed to be a vulnerability factor in stress-sensitive FSL rats [236].

Considering the connection between inflammation, NO and insulin/PPAR signalling in MD, it is not surprising that PPAR γ has been associated with suppression in immune response through its ability to inhibit the expression of inflammatory cytokines [401] and to have actions on pathways involved in apoptosis, cellular proliferation and cellular resilience [402]. Moreover, it has been demonstrated that metabolites of 5-HT act as PPAR γ -agonists in the periphery [403], which further indicates the possibility of an underlying link between biochemical pathways of mood disorders and metabolic syndrome. The recently discovered prostaglandin and endogenous PPAR γ ligand, 15d-PGJ₂, presents with anti-inflammatory properties [404], increases the neuronal metabolism of glucose, prevents stress-induced suppression of glutamate uptake [405] and has been suggested to be a possible marker for psychiatric diseases [406]. Indeed, animal studies have also described the antidepressant activity of PPAR γ agonists [407].

Bipolar Disorder

SOD was found to be increased and catalase (CAT) decreased in an ouabain-induced rat model of BPD [408], while lithium and valproate protect against amphetamine-

induced oxidative stress in the same model, thus further supporting a role for oxidative stress in BPD [119]. When considering the proposed role of oxidative stress and inflammatory process in BPD it is important to note that some mood stabilizers, eg. lithium, valproate (VPA), carbamazepine, lamotrigine, suppress (brain) pro-inflammatory mediators such as cyclooxygenase-2 (COX-2) and prostaglandins [409-411], indicating possible anti-inflammatory properties. Moreover, NAC reverses and protects against oxidative protein damage induced by d-amphetamine in a rat model of mania [412].

Schizophrenia

Animal studies, too, have confirmed that schizophrenia involves redox imbalance and oxidative stress. Using the ketamine challenge model, researchers have noted a decreased expression of PV-interneurons (relating to GABA; see section 2 and section 3.1.4) in the hippocampus [413] – a recent review by Bitanirwe and Woo, 2011 [366] explain that GABAergic pathways innervate primary neurons and could increase intracellular calcium levels and subsequently trigger oxidative damage. Elevations in nicotinamide adenosine dinucleotide phosphate (NADPH) oxidase 2 (Nox2) are observed in the prefrontal cortex of rats exposed to the SIR model of schizophrenia [414]. Nox2 is a major source of ROS and controls glutamate release in the prefrontal cortex ([415]; reviewed in [416]). We have also observed increased SOD activity, a decrease in the ratio of oxidized vs. reduced glutathione and an increase in lipid peroxidation in both the striatum and frontal cortex of SIR rats [373]. Importantly, all the latter alterations could be reversed with clozapine treatment [373]. The latter not only emphasizes the validity of these findings but also highlights the ability of contemporary treatments like clozapine to address disturbances in redox balance. Preclinical studies in the SIR model of schizophrenia also confirmed NAC's utility as adjunctive treatment to an anti-psychotic [137].

Inflammatory Markers

Increasing evidence indicates that inflammation may have a critical role in the pathophysiology of MD, BPD and schizophrenia [386, 417]. Inflammation is also a closely associated phenomenon with oxidative stress, discussed above. Cytokines play a crucial role as signalling molecules in the immune system and have the ability to cross the blood-brain barrier (BBB), granting it both central and peripheral activity [418]. Cytokines have been demonstrated to exert activity in almost every area relevant to the pathophysiology of MD, BPD and psychotic disorders including neurotransmitter metabolism, neuroendocrine function, and neural plasticity [386, 419-421]. The pro-inflammatory cytokines, such as interleukin (IL)-1, tumor necrosis factor (TNF)- α , and IL-6, can inhibit neurogenesis *in vivo* [422, 423], induce apoptosis [424, 425] and negatively affect synaptogenesis, synaptic plasticity and connectivity, and also the structure of synaptic membranes [426, 427]. On the other hand, anti-inflammatory cytokines such as IL-10 and IL-4 dampen the immune and inflammatory response [421] so that an inflammatory state is generally determined by an imbalance between pro- and anti-

inflammatory mediators. Fig. 3 provides an outline of how inflammatory mediators and oxidative stress are related to regulate immune response and redox status.

Clinical Correlates

Depression

A strong relationship has been demonstrated between MD and the presence of inflammation and its associated inflammatory mediators [428, 429]. These mediators include the pro-inflammatory cytokines, IL-1, -2, -6 and -8, interferon (IFN)- γ and TNF- α [430] that, when administered to a healthy individual, may induce sickness behaviour [431, 432]. Sickness behaviour describes a state in which many of the symptoms coincide with that of MD [420]. Furthermore, it has been proposed that constant elevation of cytokine levels may lead to neurotransmitter changes which are interpreted by the brain as stressors that further allow these molecules to contribute to the development of MD [429]. Not only do pro-inflammatory cytokines contribute towards altered neurotransmitter metabolism, neuroendocrine function, synaptic plasticity and behaviour characteristic of MD [419, 433-435], but also stimulate hypothalamic-pituitary-adrenal (HPA)-axis hormones as well as CRH in both the hypothalamus and the amygdala. The latter play an important role in fear responses and anxiety-related behaviour [419].

Cytokine-induced MD is associated with alterations in 5-HT metabolism through the activation of IDO [436], as well as alterations in CRH function [430, 436]. Importantly, MD induced by IFN- γ also involves the activation of iNOS [437], the latter known to play an important role in stress-related inflammation [438]. Peripheral levels of IL-1, IL-6 and TNF- α is increased in patients with MD [299, 439], and these effects are normalized following antidepressant treatment [440, 441]. Increased levels of IL-1 and IL-6 have also been measured in the CSF of patients suffering from MD [442]. Furthermore, an elevation in IL-6 levels has been proposed as an early marker for cognitive symptoms and has been found to correspond to the severity of MD as well as increased activity of the HPA axis [430, 443, 444]. Interestingly, IL-6 levels have been demonstrated to be increased in patients with treatment-resistant depression compared to treatment responders [445], which indicates that altered IL-6 levels in the blood of depressed individuals may serve as a marker of possible response to antidepressant treatment [208]. In line with these findings, a recent clinical study found that administration of the non-specific COX inhibitor acetylsalicylic acid leads to an improved onset of action during fluoxetine treatment and also increased the response rate to the drug when compared to patients receiving only fluoxetine [446].

Bipolar Disorder

BPD may be associated with moderately increased plasma levels of pro-inflammatory cytokines, such as IL-6 and TNF- α , along with increased IL-1, and IL-1 receptor antagonist protein, while elevated mRNA levels have been observed in post-mortem frontal cortex of BPD patients [447-449].

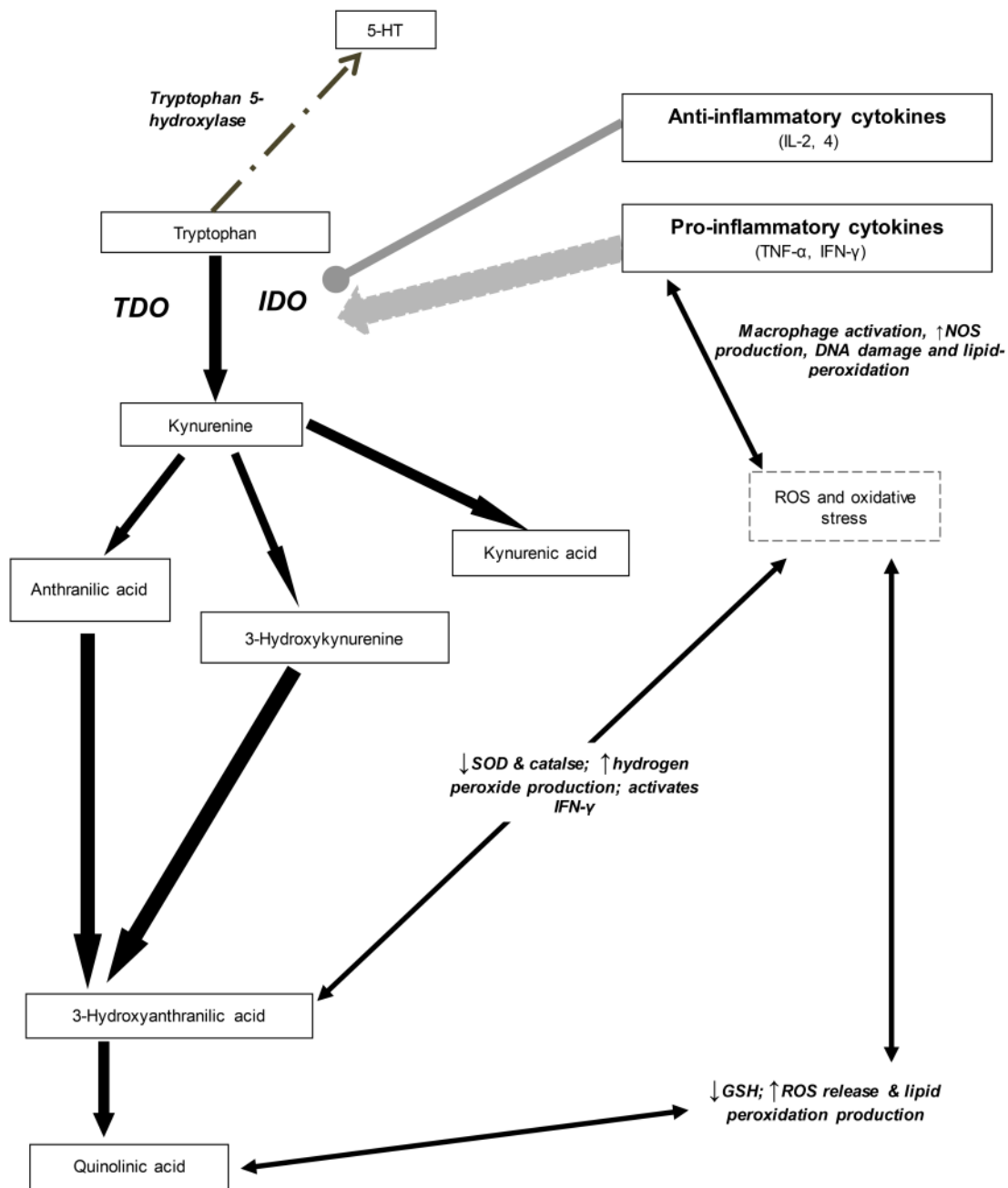


Fig. (3). A simplified diagram of the kynurenine pathway, indicating the principle enzymes, IDO and tryptophan-2,3-dioxygenase (TDO), and the subsequent formation of kynurenine and its metabolites from TRP. The diagram indicates the inter-relationship of kynurenine metabolites, particularly QA, kynurenic acid and 3-hydroxyanthranilic acid. This diagram also depicts the activation and inhibition of IDO *via* pro- and anti-inflammatory cytokines respectively, as well as the influence of oxidative stress processes that will eventually determine cellular resilience or susceptibility to neurotoxic insults. Increased activity of the tryptophan-kynurenine synthesis (by TDO/IDO) will also diminish the availability of tryptophan for serotonin synthesis *via* tryptophan-5-hydroxylase, with resulting effects on mood and behavior.

Abbreviations: glutathione (GSH); interferon (IFN)- γ ; interleukin (IL)-2 and -4; nitric oxide synthase (NOS); reactive oxygen species (ROS); serotonin (5-HT); tumour necrosis factor (TNF)- α . Adapted from [452, 453].

Schizophrenia

Schizophrenia patients and their first-degree relatives have been found to present with significantly elevated pro- vs. anti-inflammatory cytokines [450], as well as a significant elevation in pro-inflammatory cytokines in first episode psychosis patients, with a positive correlation

between IL-6 and duration of illness [418]. IL-6 has been found to be significantly increased in early and late stage schizophrenia, with IL-10 reported to decrease in the late stages [451].

The presence of elevated levels of IL-1 in the CSF has been suggested as a marker of acute psychotic relapse, while

immune-inflammatory dysfunctions may be implicated in the underlying processes mediating relapse, considering that the relationship between cytokine abnormalities and acute exacerbations of schizophrenia appear not to be related to treatment with antipsychotics. On this point, certain cytokines have been suggested to represent state markers for acute exacerbations of psychosis (see [217] for review). However, a number of studies have described inconsistent effects on plasma/serum cytokines, namely IL-4 [452], [453], IL-6 (reviewed in [454]), IFN- γ [452, 455]; reviewed in [456], and TNF- α (reviewed in [456]). Furthermore, a positive correlation between the severity of cognitive deficits and levels of IL-1 β , IL-6 and TNF- α have been described in schizophrenia [252].

Preclinical Correlates

Depression

Acetylsalicylic acid has proved to be an effective augmentation agent to fluoxetine in rats resistant to standard fluoxetine treatment using the CMS animal model of MD [457]. Furthermore, it has been demonstrated in gene-environment animal models of MD that peripheral levels of inflammatory markers and regulators of metabolic pathways, including glucocorticoids, MAPK's and cytokines are altered in these animals when compared to healthy controls [458].

Bipolar Disorder

Using the ouabain model of BPD, investigators have noted an increased activation of signalling pathways linked to inflammation in the brain [459].

Schizophrenia

Elevated pro-inflammatory cytokines (TNF- α , IFN- γ) and decreased anti-inflammatory (IL-4) and dual-action (IL-6) cytokines has been described in the SIR model, and which are for the most part reversed with clozapine treatment [137]. More interesting is that NAC is able to augment select behavioural, neurochemical and anti-inflammatory responses to clozapine in SIR animals [137]. Other pre-clinical studies have also indicated that IL-6 and TNF- α have direct inhibitory effects on adult hippocampal neurogenesis [422, 460], which may attenuate antidepressant and antipsychotic efficacy by decreasing hippocampal neurogenesis or interfering with the neurogenic properties of these drugs.

Kynurenine Pathway Markers

An important link with the inflammatory cascade is the kynurenine pathway whereby TRP is catabolized to kynurenine *via* hepatic TDO and *via* IDO in the central nervous system, lungs and placenta [461] (Fig. 3). Kynurenine is then metabolised to either kynurenic acid (KYNA) or 3-hydroxykynurine (3-OHK) and then to anthranilic acid, 3-hydroxyanthranilic acid (3-OHAA) and QA [462]. Conversion along this highly regulated pathway accounts for the metabolism of approximately 80% of non-protein bound TRP, an essential amino acid necessary for the synthesis of 5-HT (Fig. 3; [463]). QA, a NMDA receptor agonist and excitotoxin, along with

3OHK, a mediator of neuronal apoptosis, and 3-OHAA, a free radical, are all capable of inducing neurodegenerative changes in the brain [462, 464, 465]. KYNA, on the other hand, acts as an antagonist at the facilitatory glycine site on the NMDA receptor ion channel, thus having potential neuroprotective properties [466]. The activation of IDO by pro-inflammatory cytokines (e.g. INF- γ and TNF- α) in the CNS also leads to increased TRP degradation into kynurenine and QA, thereby reducing the bioavailability of TRP for 5-HT synthesis (Fig. 3) (reviewed in [467]). Hence, increased pro-inflammatory and decreased anti-inflammatory actions in the CNS can contribute to central 5-HT deficiency, which plays an important role in the pathogenesis of MD and BPD but also the negative symptoms of schizophrenia [386, 468, 469].

Clinical Correlates

Depression

Previous studies have observed decreased TRP and 5-HT along with increased kynurenine in the peripheral blood of depressed patients receiving the pro-inflammatory cytokine, IFN- γ [470, 471]. Significant decreases in the concentration of KYNA have also been observed in the plasma of depressed individuals [465]. Alterations in the symptoms of depressed patients were significantly positively correlated with kynurenine and negatively correlated with levels of 5-HT [471]. A previous study also indicated the correlation of increased IL-6 production *in vitro* with decreased TRP levels in depressed patients, emphasizing the influence of IL-6 on 5-HT metabolism *via* TRP in these patients [444]. A post-mortem study also indicated elevated QA levels in the brain of patients with MD [472]. This supports the hypothesis that the development of depressive symptoms may be mediated by increased TRP metabolism *via* IDO along the KYN pathway [436, 473].

Bipolar Disorder

Levels of TRP and kynurenine-dependent TRP index have been demonstrated to be diminished in bipolar mania [467]. Furthermore, increased kynurenine was found post-mortem in the ACC in patients with BPD, which corresponded with increased density and intensity of TDO positive glial cells [474].

Schizophrenia

In patients with schizophrenia clinical post-mortem studies found elevated levels of TRP, 3-OHAA, kynurenine and QA in various brain regions [475, 476]. Moreover, individuals suffering from schizophrenia present with increased levels of TRP in the CSF and plasma whether they received treatment or not [477, 478]. Although increases in KYNA levels have been observed in the post-mortem brain tissue of schizophrenia patients who received treatment [218], Myint and colleagues [479] have described a significant reduction in plasma levels of KYNA accompanied by a decrease in the neuroprotective ratio (a measure of the relationship between KYNA and kynurenine levels) in treated and non-treated patients suffering from schizophrenia.

Preclinical Correlates

Depression

In a stress-induced rat model of MD the kynurenine pathway was observed to be activated, leading to increased expression and activity of hepatic TDO as well as its expression in the cortex [480]. This increased TDO activity was associated with elevated circulating kynurenine concentration and a reduction in circulating TRP concentration [480]. Furthermore, induction of depressive and anxiety-like symptoms *via* the administration of a viral mimetic led to reductions in BDNF signalling and activation of the kynurenine pathway [481]. Recently, the potential for using the kynurenine 3-monooxygenase inhibitor, Ro 61-8048 (to elevate kynurenine levels), as an antidepressant has been suggested ([482], while the TDO inhibitor, allopurinol, has been noted for its antidepressant-like effects in rats [480].

Bipolar Disorder

Current animal models of BPD have not produced data related to the role of the kynurenine pathway, and data in this regard is eagerly awaited.

Schizophrenia

Recent pre-clinical studies in the SIR model of schizophrenia have described elevated plasma TRP, kynurenine, anthranilic acid, 3-OHAA and QA with reduced KYNA and neuroprotective ratio, with all alterations reversed with the antipsychotic, clozapine [137, 483].

GENETIC MARKERS

Full remission of a psychiatric illness is often impeded by variability in an individual's response to psychoactive drugs, so that being able to predict responses to psychotropic drugs holds great promise in improving treatment outcome. Genotyping, where proteins involved in neurotransmission and cellular signalling specific for illness pathology and drug response are identified and targeted, will allow the selection of an appropriate psychopharmacologic agent to best suit an individual's pathologic and metabolic characteristics.

Clinical Correlates

Depression

The development of MD has been demonstrated to be critically influenced by genetic factors which provide the opportunity to investigate mechanisms underlying the disorder [13]. Polymorphisms on genes encoding the 5-HT transporter, 5-HT_{2A}-receptor, BDNF, and tryptophan hydroxylase have been identified as candidate genes implicated in the pathology of MD [484], while several studies have suggested the Met/Met genotype (of the gene coding for COMT) to be predictive of antidepressant response rates [485, 486]. The Val/Met polymorphism in MD has not been consistent and has not been linked to any single effect [487]. Nevertheless, although contradictory findings have been presented, research into polymorphisms on the COMT variant may still hold promise [488].

Bipolar Disorder

Efforts to identify possible genetic markers and isolation of genes implicated in the pathogenesis of BPD have been challenging and at times contradictory [489]. Nevertheless, a recent study by Mahon *et al.* [490] suggested that abnormalities in lower white matter in the temporal lobe might be a marker for genetic risk of BPD. It has been suggested that the presence of an unknown gene on chromosome 12q22-q23.2 may predispose especially men to develop MD and perhaps even BPD [491]. Various studies have explored a possible link between variations in tryptophan hydroxylase II (TPH2) [492, 493] and the development of BPD, although its link with schizophrenia has been dismissed [494]. As in MD, the Val/Met polymorphism in BPD has not been consistent and has not been linked to any single effect [487].

Schizophrenia

Although the heritability for schizophrenia is estimated to be as high as 70%, the illness clearly does not have a pattern of inheritance in any population or even in single families that is consistent with the effect of a single gene [14]. Moreover, according to Riley and Kendler [495], genetics is not a determinant of schizophrenia but rather a means by which the disease is mediated. It is therefore necessary for several genes to interact and be influenced by environmental factors to lead to a patient presenting with the array of symptoms associated with schizophrenia. Like in MD and BPD, the Val/Met polymorphism is involved in predisposition to the development of schizophrenia [487] but further work is needed. The introduction of the SchizophreniaGene (SzGene) database by Allen *et al.* has made a substantial contribution to schizophrenia genomics [496]. Using the SzGene database, Sun *et al.* [497] described a collection of highly ranked genes that may be utilized as a working blueprint in the future. These genes, amongst others, include:

- i Disrupted in Schizophrenia 1 (DISC1). A gene that encodes a protein that has been implicated in neurite outgrowth and cortical development by interacting with other proteins.
- ii Dystrobrevin Binding Protein 1 (DTNBP1). A gene that encodes a protein purported to influence organelle biogenesis associated with melanosomes, platelet dense granules, and lysosomes.
- iii Catechol-O-Methyl Transferase (COMT). A catabolic enzyme involved in the degeneration of various molecules that are biologically active, including DA.
- iv D-Amino acid Oxidase (DAO). This is a NMDA-receptor-mediated signalling gene.
- v Regulator of G Protein Signalling 4 (RGS4). This is a regulatory molecule that acts as a GTPase activating protein.
- vi Neuregulin 1 (NRG1). This is a protein essential for normal development and function of the nervous system (reviewed in [498]).

vii Metabotropic Glutamate Receptor 3 (GRM3) gene. This gene, coding for the GRM3 glutamate receptor, is linked to inhibition of the cAMP cascade and has been associated with susceptibility to develop mood disorders.

Another possibly important genetic marker, and that links with oxidative stress, is NOS as discussed earlier. Weber *et al.* [499] indicated that a genetic variance in nitric oxide synthase NOS1 results in a reduction in the expression of the gene in the prefrontal brain region which adds to schizophrenia burden, and that NOS1 interacts with NOS1 adapter protein (NOS1AP) in doing so. The interaction observed in NOS1, NOS1AP and the PDZ binding domain may therefore establish a novel drug target for treating schizophrenia [499].

Preclinical Correlates

Depression

mRNA expression of neuropeptide Y have been demonstrated to be decreased in various brain regions of FSL rats [500, 501] while basal mRNA levels of several genes involved in the synthesis of neurotransmitters are similarly altered. These include tyrosine hydroxylase (TH), DA β -hydroxylase, phenylethanolamine *N*-methyltransferase (PNMT) and GTP cyclohydrolase I – all of which were elevated in FSL rats [502]. Also, protein expression of BDNF, CREB and Bcl-2 were reduced in rats after exposure to CMS [503].

Bipolar Disorder

The lack of viable genetic animal models of BPD has left a void in the search for genetic markers at pre-clinical level. However, black Swiss mice (a genetic model of mania) present with increased mRNA expression of β -catenin in the hippocampus, as opposed to other brain regions [363], while increased CSF levels of S100B (a neuronal trophic factor released by astrocytes) has been observed in the ouabain-induced model of mania [504]. Furthermore, transgenic mice expressing increased S100B have been demonstrated to be hyperactive [505].

Schizophrenia

The SIR model has shown increased expression of metabotropic glutamate receptor (mGluR)6 and ionotropic AMPA3 receptor subunit genes in the PFC [361], as well as reduced mGluR1 and mGluR5 expression [506]. These findings are consistent with the proposal that dysregulation of glutamatergic activity may contribute to the behavioural/cognitive deficits associated with SIR [64]. Furthermore, the PCP model produces up regulation of genes coding for frontal cortical NMDA receptors and produces differential expression of frontal cortical genes coding for BDNF [252]. Another clear indication of the involvement of genetic markers in schizophrenia is the numerous genetic animal models with validity for schizophrenia, such as: the DISC1 knockout (KO) model, the DA transporter KO model, the Homer1a KO model, insulin receptor KO model and the mGluR 1 and 5 KO models (reviewed in [507]).

PROTEOMIC MARKERS

The proteome is the entire collection of “proteins encoded by the genome of an organism at a specific point in

time, incorporating the set of isoforms, posttranslational modifications, covalent structures and complex protein-protein interactions present therein” [508]. Proteomics provides an insight into the character and interactions of proteins and thus of signalling pathways – understanding a proteome allows for development of effective predictive biomarkers [509].

Clinical Correlates

Depression

Ditzen *et al.* [510] observed several aspects in the CSF which were significantly altered in depressed patients when compared to controls, including 11 proteins and 144 peptide features. A recent large proteomic study in MD and schizophrenia patients observed that insulin was the marker with the highest statistical significance in MD patients compared to controls [2]. Their findings are consistent with the observation that MD is frequently linked with insulin resistance [511]. The increased comorbidity between type 2 diabetes and MD [512] and the strong link between MD and metabolic syndrome [513] further supports this hypothesis. Moreover, increased levels of chromogranin A, a secretory protein and precursor for functional peptides, are linked to dysregulated insulin levels in patients with schizophrenia [514].

Bipolar Disorder

Novikova *et al.* [515] identified several protein biomarkers which may possibly be involved in the pathology of BPD, e.g. MB-18.5, CBF2, DECR2, BYSL, ANKARD12, ALDOC and DKK2 (part of the Wnt signalling cascade) [516]. The Wnt cascade is a group of signal transduction pathways comprising proteins that pass signals from outside to inside of a cell through cell surface receptors. These proteins appear to be compromised in patients with BPD [517]. Importantly, these proteins interact with GSK3-B, BDNF, oxidative stress mediators and cytokines relevant to BPD [349].

Schizophrenia

Cyclophilin A is a protein biomarker that plays an important role in cerebral cortical plasticity [518]. Differing levels of this protein has been noted postmortem in patients with schizophrenia and BPD, suggesting contradictory influences on plasticity specifically in the DLPFC in these diseases [515]. We have earlier noted the importance of neurotrophins like BDNF and VEGF in the neurobiology of psychiatric disorders. Proteomic studies [2] have also observed significant differences in various growth factors and neurotrophins in patients with schizophrenia, influencing somatic and dendritic growth in the hippocampus and prefrontal cortex, such as BDNF, VEGF or stem cell factor. To a lesser extent, this may include members of the chemokine/cytokine family. In line with these findings, previous studies have indeed observed decreased VEGF in serum of patients with schizophrenia [259, 519], although this has not always been reproducible [520]. The current body of evidence relating to BDNF in schizophrenia provides little congruence, with increased, decreased, or no change in serum or plasma BDNF levels noted [259, 521].

Another study in schizophrenia patients identified distinctive profiles of peptides and proteins in the CSF [522] that are potentially specific for schizophrenia, including a VEGF-derived peptide sequence consisting of 40 amino acids, a transthyretin protein cluster (a serum and CSF carrier of the thyroid hormone thyroxin), and another smaller protein cluster also associated with transthyretin, with 95% specificity and 80 to 88% sensitivity [522]. Recent studies also observed that serum concentrations of insulin and chromogranin A were increased in schizophrenia patients [514, 523].

Preclinical Correlates

Depression

Yang *et al.* [524] recently published proteomic data from the CMS rat model of MD, noting decreases in glyoxalase-1 and dihydropyrimidinase-related protein-2 (DRP-2) in the prefrontal cortex, which translates to alterations in energy and glutathione pathways. Carboni and colleagues [458] detected significantly increased levels of leptin, IL-1 α and BDNF in FSL compared to FRL rats.

Bipolar Disorder

To the best of our knowledge there are no current proteomic data on animal models of BPD – this may be due to the lack of comprehensive animal models presenting with cyclic behaviour ranging from depressive to manic.

Schizophrenia

Proteomic studies in preclinical models of schizophrenia are limited. One study, however, describes the prohibitin protein, a potential marker of synaptic pathology, to be up-regulated in chronic schizophrenia patients as well as in the ketamine animal model of schizophrenia [525]. Another proteomics study describes the neuroprotective action of abrogated COX-II expression in insulin receptor KO mice as a validated animal model of schizophrenia [526]. Using the PCP animal model to access frontal cortical levels of chromogranin A, B and secretoneurin (large protein molecules of the chromogranin family), acute PCP treatment caused a decrease in secretoneurin, while chronic PCP treatment elevated this protein [527].

Micro-RNAs

Clinical Correlates

MicroRNAs (miRNAs), a large family of small non-coding RNAs, are potent regulators of gene expression with proposed roles in brain development and function [528]. These miRNAs may play a substantial role in the pathophysiology of mood and psychotic disorders and may even have an influence on the effect of drugs used to treat these disorders [529].

Depression

Bocchio-Chiavetto *et al.* [529] found 28 miRNAs to be up-regulated and 2 miRNAs to be down-regulated following antidepressant treatment – they further demonstrated that these differentiations could be associated with alteration of neuronal pathways and may be involved in the underlying mechanisms of MD.

Schizophrenia and BPD

Perkins *et al.* [530] observed that from 264 human miRNAs, 16 were differentially expressed, 15 were expressed at lower levels and 1 at a higher level in post mortem prefrontal cortex tissue of schizophrenia patients vs. controls. Similarly, post-mortem studies in schizophrenia or BPD found underexpression of several miRNAs of the adult prefrontal cortex [528], while in a study evaluating 667 miRNAs in postmortem prefrontal cortex tissue of schizophrenia and BPD patients, 22 miRNAs were found differentially expressed between cases and controls, 7 deregulated in schizophrenia and 15 in BPD [531]. Furthermore, these 22 miRNAs were found to target brain specific genes involved in neurodevelopment, behaviour as well as the development of schizophrenia and BPD [531]. It has also been observed that a considerable modification to post-transcriptional regulation may be defined by an overall increase in miRNA expression in two regions of the cerebral cortex in post-mortem brain tissue from patients suffering from schizophrenia [532].

These studies highlight miRNAs as possible biomarkers for mood and psychotic disorders, although it is clear that further research is needed to evaluate the relationship between miRNA alterations and disease development and progression. The fact that altered expression of miRNAs is reflected also in the blood of patients suffering from mood disorders potentially give rise to the idea that peripheral miRNAs may be screened as an aid in the diagnosis and treatment of these disorders. However, as with most peripheral biomarkers, the correlation between central and peripheral expression of miRNAs remains a topic of debate [533, 534].

Pre-Clinical Correlates

By referring to selected animal models of depression, recent reviews have evaluated the molecular biology of miRNAs in relation to the pathophysiology of clinical depression as well as the utility of targeting miRNAs for antidepressant treatment [533, 535]. They confirm the dysregulation of a large number of miRNAs in these depression models. Moreover, miRNAs to some extent may be associated with treatment and onset of BPD and schizophrenia. The reader is advised to consult these papers for further detail.

DEVELOPING A BIOMARKER PANEL

To date the diagnosis and treatment of patients suffering from mood and psychotic disorders have almost exclusively been based on behavioural symptomatology observed in these individuals, with laboratory testing and confirmation of diagnosis being absent due to the diverse aetiologies and underlying neurochemical abnormalities associated with these disorders. However, in recent years several biological markers have been linked to neuronal changes associated with the pathological processes and/or treatment response in these disorders. Being able to identify discrepancies in these markers in humans and using them in the diagnosis and treatment of mood and psychotic disorders will surely improve treatment efficacy and potentially even allow for provisional measures to be taken to counter neuronal deficits and prevent the onset and/or progression of symptoms and

structural brain changes associated with these disorders. Quantifying the underlying abnormalities in these disorders may also be helpful in understanding the pathogenesis of these disorders and mechanisms causing the delayed response to drugs observed in their treatment. But then again, as mentioned in the introduction the likelihood of identifying any single biomarker with sensitivity and specificity for MD, BPD and schizophrenia is relatively low so that a feasible alternative to the single-biomarker approach could be the development of biomarker panels.

From the data discussed in this review, we have attempted to distinguish between biomarkers that provide mostly substantial evidence in support of them being a strong or moderate biomarker of either MD, BPD or schizophrenia, as opposed to markers where there is less of an evidence base to link it to the neuroanatomy or pathophysiology of these disorders. Furthermore, we distinguish between markers observed in clinical vs preclinical studies in order to highlight affirmation as well as the possibility/importance of further investigation. Table 1 describes a putative biomarker panel based on the clinical and preclinical data described in this article, where strong, moderate and weak candidate biomarkers are color coded for easy identification and interpretation with the text.

Altered endocrine responses, indicated in the 2nd row of Table 1, are typical of a number of these illnesses, and can in many instances be related to altered circadian rhythm. Moreover, such changes, for example cortisol, can be instrumental in driving many of the behavioural and pathological changes evident in these disorders, eg. hyperglutamatergia, structural brain changes, etc. Furthermore, the recent introduction of the 5-HT_{2c} antagonist/M_{1/2} agonist, agomelatine, as the first antidepressant acting to re-establish altered circadian rhythms provides robust validation of the importance of these processes in the development and treatment of depression [78]. Moreover, its actions on frontal cortical DA function hints of possible value in the treatment of schizophrenia.

Regarding the neuroanatomy and neurocircuitry of MD, BPD and schizophrenia, it is evident (and quite expected) that there are distinct alterations in the volume of certain brain structures in mood and psychotic disorders, highlighting the importance of altered neuroplasticity and structural brain changes in these illnesses, indicated in the 1st row of Table 1. Decreased volumes have been recorded for various limbic structures in both MD and schizophrenia, specifically the prefrontal cortex. Although several observations have been made in BPD, current data is limited or of little clinical value and further research into the neuroanatomy of this disorder is necessary. Overall, there remains speculation as to whether the above-described neuroplastic changes are cause or effect.

Neuroimaging studies, indicated in the 1st row of Table 1, have also given insight into altered neurocircuitry, blood flow and metabolic rate in affected brain structures in these disorders and largely parallel the neurocircuitry described above. Of importance is the decreased activity in cerebellar, prefrontal, frontal and cortical structures in clinical studies of schizophrenia as well as decreases in cerebral glucose

metabolism in both preclinical and clinical studies of BPD and alterations in NAA in MD and schizophrenia.

Current pharmacotherapy of MD, BPD and schizophrenia in many ways provides robust construct and predictive validity for the importance of monoamines as biomarkers for these illnesses. At the neurochemical level, indicated in the 3rd row of Table 1, patients with MD present with a decrease in DA, 5-HIAA, NA and MHPG; BPD patients present with an increase in DA, associated with manic symptoms, and a decrease in 5-HIAA; schizophrenia patients present with an increase and decrease in DA in the striatum and prefrontal cortex respectively, along with increased 5-HT transmission and increased NA levels. Decreased levels of DA have been correlated with anhedonic behaviour in MD in both clinical and preclinical studies. Increases in DA as well as NA have, however, been linked to manic symptoms of BPD and models of mania as well as clinical and preclinical studies of schizophrenia. Glutamate on the other hand is increased peripherally and in the cortex of MD and BPD patients, is reversed with ketamine treatment, while in un-medicated schizophrenia patients the literature indicates an increase of glutamate and GABA in post-mortem prefrontal cortex and striatum tissue. However, as noted earlier, schizophrenia is proposed to involve cortical hypoglutamatergia that in turn drives meso-limbic hyperdopaminergia and meso-cortical hypodopaminergia. Decreases in the level of 5-HIAA, a 5-HT metabolite, have also been noted in both MD and BPD patients suffering from depressive symptoms, while 5HT transmission is increased in patients suffering from schizophrenia.

With regards to the growth factors, indicated in the 4th row of Table 1, the current literature supports a general decrease in BDNF in MD, BPD and schizophrenia; none to limited data of IGF-1 in MD, no changes of IGF-1 in BPD and decreased IGF-1 in schizophrenia; VEGF was increased in MD and BPD, but mRNA of VEGF was decreased in schizophrenia. While VEGF studies in humans report conflicting results and with preclinical data still lacking, it may still hold promise as a biomarker of psychiatric/mood disorders – especially considering the divergent results seen in MD vs schizophrenia patients. Recently published meta-analyses on BDNF are emphatic that altered serum BDNF is a state marker in MD, BPD and schizophrenia [536], although peripheral BDNF levels are not a sufficient measure of disease severity in MD [337] neither does it adequately discriminate between MD, BPD and schizophrenia. Animal studies tend to echo these sentiments, although serum BDNF levels may act as marker of predisposition to develop symptoms in rats used to model these disorders [306, 307]. Interesting enough, it does differentiate between mood states in BPD [336] and between acute and remitted states in MD [536]. In schizophrenia, a significant positive correlation between BDNF levels and positive and negative syndrome scale (PANSS) positive subscore has been described, as well as higher BDNF levels in the paranoid subtype of schizophrenia [537]. Low BDNF levels at the onset of psychosis may therefore contribute to the pathogenesis of schizophrenia and could perhaps be a candidate biological marker for positive symptoms. It may also play an important role as a marker of disease progression in BPD due to

Table 1. A putative biomarker panel, based on clinical and pre-clinical data for depression, bipolar disorder and schizophrenia.

	Depression	Bipolar disorder	Schizophrenia
Neuroanatomy Neurocircuitry	<p>↓ Hippocampal, prefrontal cortex, orbitofrontal cortex and basal ganglia volume [24; 25; 26; 27]* [60]#</p> <p>↑ Blood flow and glucose metabolism in amygdala, orbital cortex, and medial thalamus and ↓ Blood flow in the prefrontal cortex and anterior cingulate cortex [31]*</p> <p>↓ NAA in frontal cortex and subcortical regions [32; 33]*</p>	<p>Enlarged amygdala [41]*</p> <p>↓ Dorsal and ventral prefrontal cortical volume [42]*</p> <p>↔ Temporal lobe volume [50]#</p> <p>↓ Cortical activation, ↓ cortical gray matter and ↑ activation of the para-hippocampal gyrus and amygdala [44]*.</p> <p>↓ Cerebral glucose metabolism [61]*#</p> <p>↓ P300 wave amplitude [49]*</p>	<p>↓ Prefrontal cortex and temporal lobe volume [50; 51; 52]* [18]#</p> <p>Loss of PV-interneurons in hippocampus and ↓ frontal cortical spine density [64]#</p> <p>↓ NAAG and NAA in temporal cortex and ↑ NAAG in hippocampus [59]* [70; 71]#</p> <p>↓ Cerebellar and temporal lobe activity [46; 48]*</p> <p>↓ P300 wave amplitude [49]*</p> <p>↓ Frontal cortical and striatal activation [54]*</p> <p>↓ Prefrontal cortical function [55; 56; 58]*</p>
Neuroendocrine and circadian rhythms -Circadian rhythms -Cortisol -Thyroid hormones	<p>Circadian misalignment and changes in sleep-wake cycles [75, 76]##</p> <p>↑ Saliva cortisol [88]*</p> <p>↑ HPA-axis activation [86]*</p> <p>Thyroid and TSH dysfunction [99, 102]*</p> <p>↑ TSH [80]*</p>	<p>Altered sleep patterns, mutated core-clock-gene, Clock, leads to manic symptoms [75, 76]##</p> <p>↔ Saliva cortisol [90]*</p> <p>↑ Systemic cortisol metabolism [87]*</p> <p>Thyroid and TSH dysfunction [98, 102, 103]*</p>	<p>Sleep onset and sleep maintenance insomnia and altered melatonin rhythm [75, 76]##</p> <p>↔ Saliva cortisol [90]*</p> <p>↑ Systemic cortisol metabolism [87]*</p> <p>Thyroid and TSH dysfunction [101, 102, 103]*</p> <p>↑ Autoimmune thyroid disease [103]*</p> <p>Thyroid regulation involved in monoamine, myelination and inflammation networks [104]*</p>
Neurochemical markers -Monoamines -Glutamate and GABA	<p>↓ DA [109]* [130; 131]#</p> <p>↓ 5-HIAA [139]*</p> <p>↑ 5-HT & 5-HIAA in limbic structures [151]#</p> <p>↓ NA [186; 187]*</p> <p>↓ MHPG [188]*</p> <p>↑ Glutamate [204; 205]* [226]#</p>	<p>↑ DA (manic symptoms) [118]* [135]#</p> <p>↓ 5-HIAA (depressed) [158]* ↑ 5-HIAA (manic symptoms) [161]*</p> <p>↓ 5-HT in amygdala and hippocampus [177]#</p> <p>↑ NA [158]* [161]#</p> <p>↑ Glutamate [215]*</p>	<p>↑ DA (striatum) & ↓ DA (frontal cortex) [120; 121]* [137; 139; 146; 147]#</p> <p>↑ ↓ DA (frontal cortex) & ↔ DA (striatum) [138]#</p> <p>↑ 5-HT transmission at 5-HT_{2A} receptors [166; 167]*</p> <p>↓ 5-HT/5-HIAA in frontal cortex and ↑ 5-HT/5-HIAA in nucleus accumbens and striatum [140; 179]#</p> <p>↑ NA [191; 193]* [179]#</p> <p>↑ Glutamate & ↓ GABA [217]*</p> <p>↓ Glutamate [219; 220]* [256; 257; 141]#</p>
Neuronal growth factors -BDNF -IGF -1 -VEGF	<p>↓ BDNF [262; 263]* [304; 306]#</p> <p>↑ BDNF [281]* [301; 302]#</p> <p>↓ IGF-1 [317; 319]#</p> <p>↑ VEGF [324; 325]* ↓ or ↔ VEGF [301]#</p>	<p>↓ BDNF [504]* [119]#</p> <p>↔ IGF-1 [315]*</p> <p>↑ VEGF [327; 328]*</p>	<p>↓ BDNF [295; 298]* [311; 312]#</p> <p>↑ BDNF in paranoid schizophrenia [537]*</p> <p>↓ IGF-1 [316]*</p> <p>↓ or ↔ VEGF [285; 329]*</p>
Neuronal resilience markers	<p>↓ cAMP and MAPK/ERK pathway activity [344; 349]*</p> <p>↑ Expression of MKP [18]#</p>	<p>↓ PGRN plasma levels [351; 352]*</p> <p>↔ GSK-3 brain expression [353]* [363; 364]#</p>	<p>↑ Expression of MAPK [18; 19]#</p> <p>↑ MAPK-and cAMP proteins [354]*</p> <p>↓ DARPP-32 [359]*</p>

Table 1. contd....

	Depression	Bipolar disorder	Schizophrenia
Oxidative stress markers	<ul style="list-style-type: none"> ■ ↑SOD, lipid peroxidation [376; 377]*. □ ↓SOD and catalase [398]# 	<ul style="list-style-type: none"> ■ ↑SOD (manic and depressed symptoms) ↑catalase (manic symptoms) & ↓catalase (euthymic symptoms) ↑ lipid peroxidation [384; 260]* [408]# 	<ul style="list-style-type: none"> ■ ↑SOD, lipid peroxidation [393]* [373]# ■ ↓GSH [390; 391]* □ ↓PV-IR interneurons [413]# □ ↑Nox2 [414]#
Inflammatory markers	<ul style="list-style-type: none"> ■ ↑Pro-inflammatory cytokines (IL-1,6 ; IFN-γ; TNF-α) [299; 439; 443]* [458]# 	<ul style="list-style-type: none"> ■ ↑Pro-inflammatory cytokines (IL-1,6 ; TNF-α) [447; 448; 449]* 	<ul style="list-style-type: none"> ■ ↑Pro-inflammatory cytokines (IL-1,6 ; IFN-γ; TNF-α) [450; 418]* [104]# ■ ↓Anti-inflammatory cytokines (IL-10; IL-4) [442]* [137]#
Kynurenine pathway markers	<ul style="list-style-type: none"> ■ ↓Tryptophan & ↑Kynurenine [436; 471]* [480]# □ ↑QA [472]* □ ↓KYNA [465]* 	<ul style="list-style-type: none"> □ ↓Tryptophan & ↑Kynurenine [467; 474]* 	<ul style="list-style-type: none"> ■ ↑Tryptophan & ↑Kynurenine & ↑QA & ↑3-OHAA [476; 475]* [137; 483]# □ ↓ KYNA [479]* [137; 483]# or ↑KYNA [218]*
Genetic markers	<ul style="list-style-type: none"> ■ Polymorphisms in: 5-HT transporter, 5-HT receptor-2A, BDNF, and tryptophan hydroxylase [484]* [502]# ■ Val/Met polymorphism [487; 488]* □ ↓Neuropeptide Y expression [500; 501]# 	<ul style="list-style-type: none"> ■ Val/Met polymorphism [487]* ■ Tryptophan hydroxylase II gene variations [492; 493]* ■ ↑β-catenin expression (manic model) [363]# 	<ul style="list-style-type: none"> □ Val/Met polymorphism [485; 486]* □ NOS1 gene variance [499]* □ Possible SzGene database (DISC1, DTNBP1, COMT, DAO, RGS4, NRG1, GRM3 gene variations) [496]* ■ ↑mGluR6 and AMPA3 gene expression [361]# ■ ↓mGluR1 and mGluR5 expression [506]#
Proteomic markers	<ul style="list-style-type: none"> ■ Abnormal insulin secretion [2; 511; 512; 513]* ■ ↓Glyoxalase-1 and dihydropyrimidinase-related protein-2 [524]# ■ ↑Leptin, IL-1, BDNF proteins [458]# 	<ul style="list-style-type: none"> □ Proteins involved in the WNT cascade (MB-18.5, CBF2, DECR2, BYSL, ANKARD12, ALDOC and DKK2) [364]# 	<ul style="list-style-type: none"> ■ ↑Insulin and Cyclophilin A [514; 523]* □ VEGF-derived peptide sequence, a transthyretin protein cluster, and another smaller cluster related to transthyretin [522]* ■ ↔Secretoneurin (Chromogranin) [527]# □ ↑Prohibitin protein [525]*#
Micro-RNAs (miRNAs)	<ul style="list-style-type: none"> □ 28 miRNAs up-regulated and 2 miRNAs down-regulated in treatment [529]* 	<ul style="list-style-type: none"> □ Under expression of several miRNAs (prefrontal cortex) [530]* □ Deregulation of 15 miRNAs (involved in neurodevelopment and behaviour regulation) [531]* 	<ul style="list-style-type: none"> □ 16 miRNAs differentially expressed, 15 miRNAs down-regulated and 1 miRNA up-regulated [530; 532]* □ 22 miRNAs (involved in neurodevelopment and behaviour regulation) deregulated [533]* □ ↑miRNA expression in cerebral cortex [534]*

■ Strong marker □ Moderate marker ■ Not a strong marker/limited data

*Clinical data; #Pre-clinical data (Please refer to text for additional discussion)

Abbreviations: ↓, decrease; ↑, increase; ↔, Unchanged or differing results; PV, parvalbumin; NAA, N-acetyl aspartate; HPA, Hypothalamic-pituitary-adrenal; TSH, Thyroid stimulating hormone; DA, Dopamine; 5-HIAA, 5-Hydroxyindole-acetic acid; NA, Noradrenaline; MHPG, 3-Methoxy-4-hydroxyphenylglycol; GABA, Gamma-aminobutyric acid; BDNF, Brain-derived neurotrophic factor; IGF-1, Insulin growth factor-1; VEGF, Vascular endothelial growth factor; cAMP, Cyclic adenosine monophosphate; MAPK/ERK, Mitogen-activated protein kinases / extracellular signal-regulated kinases; MKP, Mitogen-activated kinase phosphatase; PGRN, Progranulin; GSK-3, Glycogen synthase kinase; DARPP-32, Dopamine-and cAMP-regulated neuronal phosphoprotein-32; SOD, Superoxide dismutase; GSH, Glutathione; PV-IR, Parvalbumin-interneurons; Nox2, Nicotinamide adenosine dinucleotide phosphate (NADPH) oxidase 2; QA, Quinolinic acid; KYNA, Kynurenic acid; 3-OHAA, 3-Hydroxyanthranilic acid; NOS-1, Nitric oxide synthase-1.

an observed association between peripheral BDNF levels and age and duration of illness [336, 538].

Markers of neuronal resilience, indicated in the 5th row of Table 1, have aroused considerable interest and play an important role in programmed cell death and plasticity. GSK-3 has surfaced as a possible marker in BPD *via* its regulation of several biochemical pathways, including the Wnt pathway. It has also been connected to the effects of

lithium treatment in BPD. Decreased cAMP and activity of the MAPK/ERK pathway has been demonstrated in both humans and animal models of MD and may serve as a valuable biomarker in the disorder. However, data in BPD and schizophrenia is less clear. Increases in MAPK have been noted in animal models of schizophrenia but need further investigation in a clinical setting as well.

Numerous findings in MD, BPD and schizophrenia patients indicate the presence of a prooxidative state, indicated in the 6th row of Table 1. However, elevations in SOD and lipid peroxidation are relatively non-specific, although BPD patients present with an increase and decrease in catalase in manic and euthymic symptoms respectively, making catalase a possible specific marker in the latter disorder. Furthermore, schizophrenia and BPD patients present with decreased levels of GSH [539]. Importantly, studies in translational animal models have provided evidence that the associated oxidative stress occurring with MD-related behaviors can be reversed by antidepressant treatment [400] and that the antioxidant and glutathione precursor, NAC, is antidepressant in rats [399]. Moreover, exacerbated levels of oxidative stress can attenuate antidepressant action [400]. However, preliminary clinical studies have only been able to provide limited support for the use of NAC as an adjunctive therapy for MD [379]. Similar studies in humans and animals have described reversal of redox changes by lithium and/or antipsychotics in BPD [119, 135] and schizophrenia [137, 373, 387, 395], respectively. This review is adamant that both oxidant and antioxidant systems and redox balance play a pathophysiological role in MD, BPD and psychotic disorders such as schizophrenia. This realisation has opened the door to the possible clinical utility of antioxidant drugs (e.g. NAC) in the treatment of these illnesses alone and as an adjunctive treatment [12, 118, 540].

As has been noted, inflammation and oxidative stress are closely linked. Significant increases in the levels of pro-inflammatory cytokines have been reported in humans and animal models of MD, BPD and schizophrenia, indicated in the 7th row of Table 1. Along with the above-mentioned oxidative stress observations is the evidence that MD, BPD and schizophrenia patients present with an elevation in pro-inflammatory cytokines (IL-1, 6, IFN- γ and TNF- α), that is reversed with antidepressant treatment in MD. A recent animal study in the SIR model of schizophrenia indicates that clozapine and NAC can also reverse this pro-inflammatory state [137]. This review suggests that the presence of a pro-inflammatory state is a non-specific pathological marker for MD, BPD and schizophrenia yet underscores the pathological role of inflammation in these disorders, especially given their reversal by typical drug treatment. Furthermore, anti-inflammatory cytokines have been found to be decreased in both humans and animal models of schizophrenia. However it remains unclear whether activation of inflammatory pathways in the CNS during MD, BPD and psychotic disorders is rooted in the periphery (e.g., as a function of overt or nascent medical illness or psychological stress) and/or whether stress or other yet to be identified processes (e.g., vascular insults in late life MD/psychosis) induce inflammatory responses directly within the brain. Strong evidence supports a prenatal inflammatory event as a prodromal event to the development of schizophrenia [541]. Prenatal immune challenge with either a systemic endotoxin or viral mimic vs. an inducer of local inflammation suggests that neurodevelopment of the fetus may rather be affected by circulating cytokines and/or fever as opposed to direct effects evoked on the fetus by the agent responsible for maternal infection [542]. What is evident from work in animals is that these pathological

processes seem to have their origin in a disturbance in the mitochondria [179], and explains why redox dysfunction is such a central feature of these illnesses [373]. Cytokine activity may elicit several effects on the brain, affecting the synthesis, release and reuptake of several neurotransmitters, including monamines, which have the ability to influence mood [417].

An interesting observation is that MD and BPD patients present with a decrease in TRP along with an increase in kynurenine and QA levels, while schizophrenia patients (corroborated by animal studies using the SIR model [137]), indicates an increase in TRP, 3OHAA, kynurenine and QA along with a decrease in KYNA [475, 476, 479]. These findings are indicated in the 8th row of Table 1. The decrease in TRP could therefore possibly be specific markers for MD and BPD, with the increase in TRP along with the decrease in KYNA specific to schizophrenia. The relative induction of KYNA versus QA may determine the effects of cytokines on the CNS and remains an important area for future investigation, including the therapeutic targeting of IDO and kynurenine enzymatic pathways in MD, BPD and schizophrenia [465, 467, 475].

Even though these disorders are known to be hereditary to some extent, the exact genetic basis still needs further elucidation, indicated in the 9th row of Table 1. Val/Met polymorphisms have been studied in MD, BPD and schizophrenia and even though various reports have been made, data remains inconclusive. Polymorphisms in 5-HT transporters and receptors, as well as BDNF and tryptophan hydroxylase hold promise in MD and warrants further research to pinpointing exact genetic markers involved in the development of MD [484]. DISC1 is a well researched candidate gene for schizophrenia and affective disorders with a range of functions relating to neurodevelopment, although studies into its role in these disorders remains promising albeit conflicting [543]. Gene variations in the SzGene database [496] also hold promise in schizophrenia research and also needs further research to clarify predisposition in developing schizophrenia. Tracking genetic variants in patient blood may therefore serve to compliment biomarker panels by providing more information relating genotypes to MD, BPD and psychotic disorders and their respective treatment responses.

With regards to proteomic markers, indicated in the 10th row of Table 1, it is clear that utilizing modern proteomic techniques, especially mass spectrometric approaches, may support attempts to understand the biochemical processes that accompany psychiatric disorders and may in turn lead to the development of diagnostics and better therapeutics. In MD, abnormalities in insulin secretion has been observed [511] and the disorder is also accompanied by decreased levels of glyoxalase-1 and dihydropyrimidinase-related protein-2 and increased leptin, IL-1 and BDNF protein levels in animal models [458, 524]. This supports the hypothesis for a shared etiopathology with an inflammatory underbuild in patients with co-morbid MD and metabolic syndrome and/or type II diabetes mellitus, which highlights the proposed utility of the PPAR γ -pathway in the treatment of MD [544]. Increased insulin levels have also been reported in patients suffering from schizophrenia and

may be accompanied by increases in cyclophilin A [518], suggesting increased support for the role of inflammation in the disorder with cyclophilin A already being linked to a variety of disorders with an inflammatory component – among others, type II diabetes [545]. Moreover, these findings may relate to the confounding observation of weight gain and metabolic syndrome in this disorder, and that may be worsened by certain antipsychotic drugs [546]. Various proteins involved in the WNT cascade may possibly serve as proteomic biomarkers of BPD which may lead to these markers aiding in the diagnosis and treatment of the disorder and therefore warrants further investigation. Thus, screening peripheral compartments, such as serum and CSF, in patients and controls for altered expression of proteins and metabolites known to be involved in the pathophysiology of the disease or associated with comorbid states could serve in developing a “fingerprint” for identifying persons at risk of developing MD, BPD or schizophrenia. However, it is critical that we bring together knowledge on the biology of these illnesses, co-morbid states, illness severity and treatment resistance to enable proteomic markers to realize this potential.

Finally, studies suggest that micro-mRNAs, indicated in the 11th row of Table 1, may play a valuable role as a biomarker in the diagnosis and treatment of mood and psychotic disorders, however further research is warranted and the relation between central and peripheral expression still needs elucidation.

This review has focussed primarily on suitable disease-specific biomarkers with especially predictive validity. However, we have on occasion made reference in the foregoing sections to *physiological markers*. It is maybe incumbent to mention that these markers are gaining in interest, with some recently been found to have value in predicting treatment response. Thus for example, we earlier described that clinical response to antidepressants can be predicted by assessing activity in the rostral ACC region *via* electroencephalography (EEG) [37]. Similarly, we noted that by studying P304 wave amplitude, EEG can be used to assess decision making in BPD and schizophrenia patients [49]. In a study based on the “disconnection hypothesis” of schizophrenia [547], and with accumulating evidence of abnormal functional connectivity in schizophrenia, Takahashi and colleagues argued that neurophysiologic signals may provide a retrospective window with which to view disordered neural dynamics in schizophrenia [548]. Using a novel entropy-based approach for measuring dynamical complexity in physiological systems, they observed abnormal dynamical EEG signal complexity in anterior brain areas in schizophrenia that normalized selectively in fronto-central areas following antipsychotic treatment. This approach has also been proven successful in MD [488]. Another promising neurophysiological marker is electroretinography (ERG), a specialized measure of retinal function, which has been studied in schizophrenia and BPD [549]. ERG abnormalities may reflect altered phospholipid metabolism and/or impaired dopaminergic transmission. With all patients receiving stable psychotropic medications at least for 2 weeks before the first assessment, the authors found that retinal dysfunctions are specific for schizophrenia,

as compared with BPD, and are confined to the acute stage of the illness. Another potential physiological marker is event-related-potentials (ERPs) (voltage fluctuations in an EEG depicting neural activity), which are specific for cognitive dysfunction in schizophrenia [550] and also studied in MD [551]. Although physiology markers have not been as extensively studied as biological markers, there is a literature describing their use in animals, such as employing EEG and related markers in translational animal models of MD [552] and schizophrenia [553]. For example, blind-drunk (Bdr) mice demonstrate fragmented rest and activity rhythms under a light/dark cycle, reminiscent of altered sleeping patterns in schizophrenic patients [83], while depriving animals of REM sleep, which can be studied as an EEG-related marker, has been suggested to model mania [82].

DISCUSSION AND CONCLUSION

The search for blood biomarkers can essentially be divided into screening putative markers inferred by our current knowledge of the given illness, e.g. BDNF, CREB etc, or through exploring candidate pathways through the use of “omic” procedures, such as proteomic or transcriptomic profiling that offer an unbiased view of these pathways. Either approach has its own set of advantages and disadvantages, yet studies deploying either still lack the critical requirement of reproducibility and selectivity [13]. Another important challenge in identifying possible biomarkers is the predictive efficacy of a specific biomarker in the treatment of MD, BPD and schizophrenia. Very few biomarkers of these illnesses have shown utility in regards to predictive efficacy following drug treatment. Having this in hand offers the possibility of introducing tailored pharmacotherapy. Furthermore, demonstrating a dose-dependent response of these markers under the conditions of treatment may aid in more accurately establishing an appropriate dose selection during clinical trials, thereby optimizing drug discovery and development. Knowing this brings the focus of future research to a more optimized translational approach [554]. Ideally, questions from a clinical situation should be translated into a valid animal model, where such animal data can be integrated with patients’ data in order to identify predictive biomarkers. Thereafter extensive validation should be performed on these biomarkers before diagnostic kits with predictive value can be developed and marketed [554]. In the end, the final requirement is that the foregoing process should allow clinicians to make evidence-based decisions that will reinforce the decision to treat, with what agent/s, and with a higher likelihood of success than that provided by current approaches. However, there are a number of obstacles to overcome before realising a biomarker panel that is sensitive and specific enough to be implemented as a reliable tool for diagnosis and treatment, which include:

- difficulties in translating findings observed in animal models to clinical studies and correlating markers measured in animal subjects with those measured in human patients [13, 555];
- employing clinical studies in a larger population in order to validate specific findings [13, 555, 556];

- attributing measured biomarkers to one specific pathway [2] as measured in a specific disorder, thus ensuring biomarker specificity [13] and the possible presence of underlying comorbid disorders [13, 547];
- the complexity of underlying pathophysiologic and etiologic origins of these disorders combined with demographics [556], and interpatient variables, eg. smoking [2];
- correlating data from observations made in different locations and utilizing different sampling techniques [2];
- accounting for disease state [13, 556] and previous/current drug therapy [2, 557];
- correlating levels of markers measured in different tissue samples, eg. plasma vs CSF [2, 13];
- the influence of the time at which biomarkers are measured, eg. the influence of circadian rhythm and disease state/progression [2, 558];
- and the inability to measure biological markers in brain tissue in live patients due to the invasive nature thereof [559].

Nevertheless recent studies using “omics” approaches have demonstrated that careful selection of appropriate biomarker panels can provide good separation between diseased and healthy states, as well as predict response to treatment. Two recent studies by Pajer [555] and Redei [556] set out to investigate the validity of potential biomarkers of MD and found that several of these markers may have possible use in discriminating between depressed and non-depressed individuals and may even predict response to therapy. Furthermore Redei *et al.* found several blood markers identified in animal models of depression to correlate with levels measured in depressed human individuals. Importantly, these studies affirm the approach taken in this review that clinical data and that obtained from validated translational animal models are supportive and should be used together when developing a biomarker panel [555]. Indeed, the latter studies corroborate that genes expressing transcripts belonging to processes related to transcription, neurodevelopment, neurodegeneration and redox are causally related to at least MD [555], which concurs with molecular mechanisms linked to these processes and highlighted in this review. Although metabolomics per se has not been covered in this review as it addresses much the same processes as do genomic and proteomic methods, studying the metabolome has great potential to map potential biomarkers in neuropsychiatric disorders. Indeed, as has been noted, mood and psychotic disorders are linked to a range of disturbances in metabolic pathways, eg. neurotransmitter systems, TRP-kynurenine metabolism, oxidative stress, etc, so that generating a metabolic signature for a specific disorder will aid in metabolic phenotyping and contribute to discovering disease-specific biomarkers as well as predicting treatment response [560].

Based on this review, the ever-increasing availability of new pre-clinical and clinical studies is beginning to forge a way through the neuropathologic complexity of illnesses like

MD, schizophrenia and BPD, so much so that we are in a position to portray how altered neuroendocrine, anatomical, neurochemical and other pathologies can be linked to a specific disorder. The role of the endocrine system has long been linked to mood and psychotic disorders with MD patients presenting with increased saliva cortisol as well as HPA-axis activation. Similarly, patients suffering from BPD and schizophrenia have been found to have increased systemic cortisol metabolism. Dysfunction of the hypothalamic-thyroid axis has been demonstrated in all three disorders.

In MD, decreased hippocampal volume as well as reductions in the size of the prefrontal cortex and basal ganglia is accompanied by reduced levels of monoamine neurotransmitters (NA, 5-HT and DA), decreased levels of the 5-HT metabolite, 5-HIAA, and an increase in the levels of glutamate. However, DA has been noted to be increased in BPD and increased in the striatum and decreased in the frontal cortex of both schizophrenic patients and most animal model studies of the disorder. In contrast with MD and BPD, increased 5-HT transmission and NA levels characterize neurotransmission in schizophrenia and is accompanied by decreased NAAG levels in the temporal cortex while increased in the hippocampus as well as reduced activity in various brain regions, including the cerebellar and temporal lobes, prefrontal cortex, cortex and striatum.

Although a variety of observations have been made regarding BDNF, VEGF and IGF, data currently available report conflicting findings. However, hypotheses and data surrounding these markers make a strong case for their involvement in these disorders – whether as a cause or result of underlying pathology. Continued investigation will, more than likely, eventually lead to pinpointing the exact roles of these markers in the pathophysiology and/or progression of mood and psychotic disorders and establish them as valid biomarkers in the diagnosis and/or treatment of MD, BPD and schizophrenia.

Inflammation has emerged as a central role player in the pathophysiology of all three disorders discussed in this review with levels of pro-inflammatory cytokines being observed to be markedly increased in MD, BPD and schizophrenia and a decrease in anti-inflammatory cytokines also contributing to the inflammatory component of schizophrenia. Inflammation is thus not illness specific but a residual marker of ongoing pathology. Increased levels of kynurenine add to the immune response in these disorders with TRP levels being decreased in MD and BPD but increased in schizophrenia. Closely associated with inflammation, nitrosative and oxidative stress deepens the extent of neuronal stress in these disorders – increased lipid peroxidation accompanied by raised levels of SOD are a feature in all three disorders, accompanied by decreased GSH in schizophrenia. The individual inflammatory components that characterize said inflammation in these disorders may in the end prove to be more illness-specific markers. Thus for example, we noted earlier that a decrease in TRP may be specific markers for MD and BPD, while the increase in TRP and decrease in KYNA is more specific to schizophrenia. Similarly, certain components of the inflammatory response such as NO may be pro- or anti-

inflammatory depending on the cellular milieu and/or pathways activated (eg see nNOS vs iNOS-mediated pathways in Fig. 1).

The current body of literature features data on a wide variety of possible biomarkers linked to mood and psychotic disorders. To improve diagnostic techniques and treatment strategies, it is of great importance that possible trait and state markers of these disorders are scrutinized to a point where they can be incorporated into an appropriate panel of biomarkers (as presented in Table 1) which may serve as adjunct to current diagnostic criteria. Furthermore, such a panel may assist in treatment strategies being tailored to the unique context in which mood and psychotic disorders present in each individual. In this manner we may move forward from the current “one-size-fits-all” approach to treating an individual to one that addresses the biological processes underlying the disorder and specific for that particular patient.

AUTHOR DISCLOSURES

Sources of Funding

The authors declare that this work has been funded by the South African Medical Research Council (BHH). The funder has no other role in this study.

AUTHOR CONTRIBUTIONS

BH Harvey devised the concept and lay-out of the manuscript, as well as finalized the pre-submission version of the manuscript. S Brand prepared the first draft as well as the final version of the manuscript; M Möller contributed to the initial and final versions of the manuscript.

CONFLICTS OF INTEREST

The authors declare that over the past three years Brian Harvey has participated in speakers/advisory boards and received honoraria from Servier, and has received research funding from Lundbeck and Servier. The authors declare that, except for income from the primary employer, research funding to BHH from the South African Medical Research Council as well as the honoraria described above, no other financial support or compensation has been received from any individual or corporate entity over the past three years for research or professional services, and there are no personal financial holdings that could be perceived as constituting a potential conflict of interest.

ACKNOWLEDGEMENTS

Declared none.

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