Bipolar disorder (BPD) is a devastating illness that is characterized by recurrent episodes of mania and depression. In addition to these cyclic episodes, individuals with BPD exhibit changes in psychovigilative function, cognitive performance, and general health and well being. In this article we draw from neuroimaging findings in humans, postmortem data, and human genetic and pharmacological studies as well as data from animal models of behavior to discuss the neurobiology of BPD. We conclude with a synthesis of where the field stands and with suggestions and strategies for future areas of study to further increase our conceptual understanding of this complex illness.

Introduction
Bipolar disorder (BPD), classified as a mood disorder in the Diagnostic and statistical manual of mental disorders (4th edition), is a common, chronic, and recurring medical disorder that is characterized by episodes of mania — extremely elevated mood, energy, unusual thought patterns, and sometimes psychosis — and depression. Although these episodes are usually interspersed with periods of relatively normal mood, BPD is the cause of significant suffering for both patients and their families. BPD leads to limited functioning, which often results in decreased productivity in both the personal and the professional arenas of the patient’s life. The prognosis for patients with BPD is poor, with high rates of relapse, lingering residual symptoms, cognitive impairments, and diminished well being (1). Moreover, individuals with BPD frequently have coexisting medical conditions, such as obesity, cardiovascular disease, diabetes mellitus, and thyroid dysfunction, all of which are exacerbated by their BPD symptoms (2).

The prevalence of BPD was thought to be around 1%, but current reported diagnoses indicate that this figure may be closer to 5%. This increased prevalence is mainly accounted for not by an increase in diagnosis of full-blown BPD (which is known as BPD I), but by various suffer (i.e., less severe) conditions that fall under the BPD spectrum. Disorders under the BPD spectrum have been grouped based on some overlap in clinical manifestations; however, whether they share the same underlying genetics and pathophysiology is uncertain. Therefore, we focus on the emerging neurobiology surrounding BPD I, referred to herein simply as BPD (3).

Because of the elevated morbidity and mortality suffered by individuals with the disorder, BPD has been increasingly recognized as a major health problem. Despite advances in its diagnosis and recognition, the underlying neurobiology of BPD remains largely unknown. It is thought that BPD is a multifactorial disease that results from a combination of different genetic profiles, characterized by the presence of various protective and/or preventive genes relative to susceptibility and/or risk genes as well as environmental influences, including chronic stressors and traumatic experiences.

Historically, the brain systems receiving the greatest attention in neurobiological studies of BPD have been the monoaminergic neurotransmitter systems, i.e., the serotonergic, noradrenergic, and dopaminergic neurotransmitter systems. These neurotransmitter systems are extensively distributed throughout the brain’s network of limbic, striatal, and prefrontal cortical neuronal circuits and are thought to support the behavioral and visceral manifestations of mood disorders (Figure 1). Despite evidence showing that many of these circuits are likely to be involved, no obvious degeneration or complete dysfunction of any single neurotransmitter system has been identified. In this regard, the biological underpinnings of BPD appear to differ from classic neurodegenerative disorders such as Parkinson disease and Alzheimer disease, where clear deficits can be traced to the dopaminergic and cholinergic pathways, respectively. However, as we discuss here, many researchers believe that BPD arises from modulation of synaptic and neural plasticity in critical circuits mediating affective and cognitive function. Thus, BPD may represent a disorder of altered synapses and circuits, rather than being the result of imbalances in individual neurotransmitters.

In this Review, we discuss how the currently employed research strategies have furthered our understanding of the underlying neurobiology of BPD. We have broadly categorized these strategies into three main areas: patient-based research including neuroimaging, postmortem tissue analyses, and genetic association studies; analysis of pre-clinical animal models of BPD, such as models of stress and/or depression, models of mania, and, more recently, genetic models that resemble specific facets of BPD symptomatology; and molecular and cellular pharmacologic studies that attempt to identify the cellular and molecular effects of validated mood-stabilizing drugs (MSDs). We end with a conclusion about where the field stands, both clinically and pre-clinically, and discuss what we believe is necessary to make the next steps forward. Clearly, we are at the early stages of understanding the neurobiology of this complicated and multi-factorial disease. However, better understanding of the underlying biology is critical for the future development of targeted therapies that are both more effective as well as free from harmful and/or intolerable side effects.

Patient-based research
Neuroimaging. Functional neuroimaging studies have identified abnormalities in the brains of individuals with BPD, which may be indicative of dysfunction in key neural circuits (4). These circuits are distributed throughout a wide array of neuronal structures, including the amygdala and related limbic nuclei, the orbital and medial prefrontal cortex (PFC), the anterior cingulate, the medial thalamus, and related regions of the basal ganglia (Figure 2) (5).
It is plausible that dysfunction or modulation of these circuits may predispose an individual toward manifestation of the symptoms of BPD. For example, studies have shown reduced activity in the right PFC during episodes of mania (6), and dysfunction of the right PFC is thought to contribute to a disinhibited profile, including poor impulse control, risk taking, distractibility, poor sustained attention, and delusions, all of which resemble the symptoms of mania (4, 7).

Researchers using CT and MRI have identified structural changes in the brains of patients with mood disorders, including patients with BPD. Overall, gray matter volume is not substantially different in patients with BPD compared with normal healthy individuals (8–10). However, several studies have found region-specific reductions in gray matter volume, including increased ventricular size and decreased frontal cortical volume (11, 12). Specifically, volumetric decreases have been identified in the anterior cingulate cortex (13), and other studies have indicated gray matter loss in the left dorsolateral PFC (DL-PFC) (14), the ventral PFC, and the orbital PFC (15). Temporal lobe regions, including the hippocampus and the amygdala, have not been as well studied. However, one study found more prevalent volumetric decreases in the right hippocampal formation of the affected twin in monozygotic twin sets discordant for BPD (16).

White matter hyperintensities (WMHs) around the ventricles and in the subcortical white matter have consistently been found in the brains of depressed elderly persons as well as in the brains of patients with BPD (17, 18). Although the functional and pathological relevance of WMHs has not been fully determined, WMHs have been associated with cerebrovascular accidents (stroke and transient ischemic attacks), ischemia, axonal loss, increased perivascular space, minute brain cysts, and necrosis (19). Recent data suggest that a substantial proportion of individuals with BPD, including children diagnosed with BPD, exhibit WMHs more frequently than the general population. The incidence of WMHs is higher in children with neuropsychiatric disorders, including children with BPD (20, 21). In addition, WMHs are associated with poor treatment outcome in patients with mood disorders, particularly when localized to the subcortical rather than the periventricular areas (22, 23). Combined studies on the increased incidence of WMHs in affective disorders suggest that these lesions could indicate some type of damage to brain tissue, which could result in disruption of the neuronal connections necessary for normal behavioral functioning (24).

Imaging studies using magnetic resonance spectroscopy (MRS) led to the notion that BPD may be associated with mitochondrial dysfunction (25, 26). In particular, high-resolution 1H-MRS imaging studies conducted to quantitatively assess concentrations of N-acetyl-aspartate (NAA), a predominant neurochemical in the human brain that is localized to mature neurons and synthesized within mitochondria, found decreased concentrations of NAA in various patient populations, including patients with BPD (27–32). In addition, studies using 31P-MRS, which allows examination of energy metabolism in the brain, showed a decrease in phosphocreatine and/or ATP levels in patients with a mood disorder, including patients with BPD (33–35). Consistent with these findings, low brain pH levels, as measured indirectly by 31P-MRS, have also been identified in patients with a mood disorder (33–35).

While it is clear that BPD is not a mitochondrial disorder per se, some data suggest that cellular and molecular abnormalities
MRS study of children with BPD, patients were shown to have increased levels of glutamate in the frontal lobes and basal ganglia, which is responsible for the regulation of synaptic strength (40). Mitochondrial dysfunction may also be involved in the calcium signaling abnormality found in BPD (37). Regulation of intracellular calcium levels may be particularly important in the context of the CNS, since fast changes in the levels of calcium are responsible for mediating actions associated with the release of, and response to, neurotransmitters. Changes in calcium levels are also crucial for initiating transcription events at several genes whose transcription is dependent on increases in neuronal activity. One of these neuronal activity–dependent genes, brain-derived neurotrophic factor (BDNF), has been implicated in the etiology of several mood disorders (38). Furthermore, it is known that mitochondria are involved in the initiation of apoptotic processes (39), and newer evidence has suggested that proper mitochondrial function may be important for the regulation of synaptic plasticity. For example, increased neural activity has been shown to induce transcription of genes encoded by mitochondrial-related genes in the occipital cortex of depressed adult patients (42). Although current data from neuroimaging studies are interesting, their interpretation remains incomplete and often controversial. Despite finding differences in specific locations and deficits in biochemical markers, it is not yet understood what these findings represent or how they may affect the function of various circuits in the brain.

In addition, it is not yet known whether the abnormalities that have been discovered represent developmental problems that confer vulnerability to severe mood disorders, compensatory mechanisms for other related pathogenetic processes, or the continual recurrence of affective episodes.

Analysis of postmortem human tissue. Studies of postmortem brain tissue from patients with recurrent mood disorders, including some with BPD, have found reduced subcortical nuclei volumes (43). Various other studies have shown differential neuronal densities and morphologies that appear to be layer and cell-type specific in both patients with BPD and patients with major depressive disorder. For example, the density of neurons containing large cell somas, which are likely to correspond to glutamatergic pyramidal neurons, was decreased in layers III and V of the DL-PFC of individuals with recurrent mood disorders (44). Furthermore, the size of DL-PFC neurons in layers V and VI was also reduced (45). In patients with BPD, the size of the cell soma of pyramidal neurons in the CA1 region of the hippocampus has been found to be decreased (46), as have cell densities in layers III–VI in different regions of the anterior cingulate (24, 36). Interestingly, cell densities of nonpyramidal neurons were decreased in layer II, and neuron size was increased in layers II and V of these anterior cingulate regions (24, 36).

In addition, decreased levels of calbindin- and parvalbumin-expressing neurons, both of which are subtypes of GABAergic interneurons, have been identified in the anterior cingulate cortex, the hippocampus, and the entorhinal cortex of patients with BPD (47, 48). These data, in combination with studies showing decreased hippocampal expression of glutamic acid decarboxylase 67 (GAD67) and somatostatin in the hippocampus of patients with BPD, have led to the hypothesis that a subset of hippocampal interneurons may be abnormal in BPD (49).

In addition to changes in the size and density of neuronal cell types, changes in glial cell biology have also been observed in studies of postmortem tissue from patients with BPD. The density of glial cells seems to be decreased in frontal cortical areas, while the nucleus size increases (44, 50–52). Moreover, a proteomics study using brains from individuals with BPD and individuals with major depressive disorder found disease-specific alterations in levels of glial-fibrillary acidic protein (GFAP), an abundantly expressed astrocyte-specific protein (53). Additional studies have found reductions in oligodendrocyte number and in the expression of genes that are related to oligodendrocyte differentiation and myelin production in the DL-PFC of individuals with BPD (54). The recurrent theme of decreased cell density and number may represent cell loss and atrophy in these patients over the course of disease progression. It is presently unknown whether this type of brain atrophy is one of the underlying causes of the disease, or whether it contributes to illness pathology by disrupting the normal circuitry that is key to normal affective and cognitive functioning.

As discussed above, neuroimaging studies have provided suggestive evidence for abnormalities in mitochondrial function and energy production in BPD. Analysis of expression levels of key mitochondrial-related genes in the postmortem human brain has provided additional evidence to support this idea. A recent microarray study comparing postmortem hippocampal tissue from individuals with BPD and individuals with schizophrenia as well as normal healthy controls revealed that the expression of 43 genes was decreased in patients with BPD compared with those with schizophrenia (49). Furthermore, 42% of these genes encoded mitochondrial proteins.
In addition to the prominent findings of altered expression of genes encoding mitochondrial proteins, substantial changes have been observed in the level of expression of proteins implicated in synaptic function. These findings suggest alterations in synaptic plasticity mechanisms in individuals with BPD. The neuronal plasticity marker GAP-43 is highly expressed in axonal growth cones during development and is implicated in regulation of axonal morphology and synaptic plasticity in the mature brain (55). In patients with BPD, it has been reported that levels of GAP-43 are reduced in both the cingulate cortex and the hippocampus (56, 57). The synapsin family of proteins binds synaptic vesicles to the cytoskeleton, preventing their transport to the presynaptic membrane and subsequent neurotransmitter release (58). Docking of synaptic vesicles and neurotransmitter release are regulated by a complex of proteins that includes SNAP-25, syntaxin, synaptobrevin, and synaptophysin. In postmortem brains from individuals with BPD, a reduction in the levels of synapsin family members has been found in the hippocampus (59). However, increases in the levels of SNARE complex proteins, which are responsible for mediating the fusion of synaptic vesicles with the cell membrane, have been observed in the DL-PFC (60). An additional study showed decreased levels of synaptobrevin and synaptophysin in the visual association cortex of patients with BPD (61). The levels of mRNAs encoding netrins, a family of proteins important in regulating axon guidance, have also been shown to be reduced in the CA3 region of the hippocampus and entorhinal cortex of patients with BPD (62).

Genetic studies. Combined evidence from family, twin, and adoption studies suggests that BPD has a strong genetic component. Twin studies show a highly elevated concordance rate in monozygotic twins when compared with dizygotic twins (63, 64), and BPD is more likely to occur in biological parents of adopted children than in the adoptive parents (65). Strategies for detecting the genetics of BPD include both linkage and association studies. Linkage methods test the loci of vulnerability genes by studying co-inheritance of chromosomal fragments in specific illnesses, whereas association studies test whether a specific gene variant is associated with a given disorder. A list of genes, including their known functions and potential role in the etiology of BPD, that have been implicated via both linkage and association studies is lengthy and beyond the scope of this review (see ref. 66 for an in-depth discussion). Of the implicated genes, most attention has been given to those that encode proteins known to interact with signaling pathways previously implicated in BPD. For example, several of the risk genes that have been identified are known to interact with the PKC and glycogen synthase kinase 3β (GSK3β) signaling pathways (67, 68). In addition, other putative susceptibility genes, including glutamate receptor, metabotropic 3 (GRM3) and GRM4; glutamate receptor, ionotropic, N-methyl-d-aspartate 2B (GRIN2B); D-amino acid oxidase (DAO); and DAO activator (DAOA, also known as G72), encode proteins involved in glutamatergic signaling (68, 69). In addition, a number of genes encoding proteins involved in circadian biology, including ARNT-like protein 1, brain and muscle (Bmal1), TIMELESS, and PERIOD3, have been implicated as susceptibility genes for BPD (68). These genes are of interest because virtually all patients with BPD have alterations in circadian function, including alterations in sleep patterns, activity, hormonal secretions, and appetite.

Beyond the older linkage and association studies, recent advances in technology have allowed researchers to study the genetic component of BPD using genome-wide association studies (GWASs). Four groups have recently performed independent GWASs of BPD (67, 70–72). However, the significance of these findings is unclear, since very few findings have been replicated from sample to sample and there is the possibility of multiple false positives due to the substantial number of comparisons. In one study using 1,233 patients with BPD and 1,439 control subjects, Baum and colleagues identified a SNP in diacylglycerol kinase η (DGKH) as being associated with BPD (67). The Wellcome Trust Case Control Consortium (WTCC) analyzed 1,868 individuals with BPD and 2,938 control subjects and identified a locus in a gene rich region of high linkage disequilibrium on chromosome 16p12 as being associated with BPD (70). Sklar et al.’s study of 1,461 patients with BPD and 2,008 control subjects found their strongest result in myosin 5B (MYOS5B) (71). The most recent GWAS, by Ferreira et al., used a new sample with 1,098 individuals with BPD and 1,267 control subjects and found the strongest results in ankyrin G (ANK3) and calcium channel, voltage-dependent, L-type, α 1C subunit (CACNA1C) (72). A broad comparison of the WTCC and the Sklar study also confirmed the CACNA1C results, identifying CACNA1C as showing a consistently strong signal in individuals with BPD (72). More studies are clearly necessary before any consensus can be made as to the significance of these recent findings.

Analysis of pre-clinical animal models of BPD
While considerable caution undoubtedly needs to be taken in applying animal models to complex neuropsychiatric disorders, animal models can be valuable tools for exploring the underlying pathologies of human diseases and developing better therapies. An ideal animal model for BPD should show face validity, i.e., it should include spontaneous and progressive behavior that oscillates between increased and decreased manifestations of the behavior being modeled, which should be similar to a behavior characteristic of either human mania or depression. Moreover, the modeled behavior should show predictive validity, i.e., it can be normalized by treatment with MSDs (36). The progressive and cyclic nature of BPD presents a unique challenge for modeling in rodents. Indeed, as discussed in detail below, most models have tended to focus on either mania or depression, rather than modeling both behaviors, which both occur in individuals with BPD (36, 73). Currently, more models of depressive illness exist than of mania or of the cyclical nature of BPD (73). Increasing the difficulty of developing animal models of BPD is the hypothesis that an interaction between both genes and the environment are needed in order to manifest symptoms of the illness. Determining what types of environmental disturbances to pair with the various genetic alterations is difficult, but may be critical in effecting useful models.

Models of stress and/or depression. A commonly used line of research for those studying BPD using animal models is to examine the relationship between cellular and molecular characteristics of animals and behavior in laboratory situations that replicate environments thought to predispose humans toward depression. Some of the more commonly used models have been maternal separation and various chronic stress paradigms including psychosocial stress, mild unpredicted stress, and restraint/imobilization stress (36, 74). Another model used to induce a depression-like state is the learned helplessness model (73, 75). In this model, animals are administered an inescapable aversive stimulus, frequently an electric foot shock. Following the inescapable portion of the paradigm, animals are again administered the aversive stimulus, but in an environment where they are capable of escaping the aver-
sive stimulus. Longer latencies or a lack of any response is seen as indicative of “learned helplessness” (73, 75). The forced swim test is the most commonly used animal model in depression research, and more specifically has been widely used as a screen for antidepressant treatments (74, 76). These models have been useful thus far in helping to screen for potential therapeutic candidates. Further, studies with genetically modified animals using these paradigms have helped to identify candidate molecular and genetic signaling pathways that may be involved in the development of behavioral attributes that mirror disease symptomatology. However, we have yet to be able to adequately model the diseases well enough to determine anything conclusive about the overall etiology of the disorder.

Models of mania. Hyperactivity is one of the more simple behaviors to detect and quantify in animals. Thus, many models of mania that have been developed have focused on this aspect as the core of their modeling (36, 77). Acute treatment with psychostimulants such as methamphetamine and cocaine can produce a wide range of mania-like behaviors, including hyperactivity, heightened awareness and alertness, insomnia, and changes in sleep patterns (78, 79). Psychostimulant-induced hyperactivity is sensitive to lithium treatment and possibly to other anticonvulsants that have been used as MSDs. Sleep deprivation has also been used with some success as a model for mania. Rats exposed to 72 hours of sleep deprivation exhibit a variety of behaviors that show face validity for modeling human mania. These behaviors include insomnia, hyperactivity, irritability, aggressive behavior, and hypersexuality (78, 79). Furthermore, in terms of predictive validity, lithium has been shown to alleviate some of these mania-like behaviors.

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<td>Important genetically engineered mice that model BPD behaviors</td>
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Open arm time is the time an animal spends on the open arm of an elevated plus or zero maze; anxiolytic drugs increase open arm time. Open arm entries are the number of times during the EPM test that the animal enters into one of the open (i.e., unenclosed) arms of the EPM. Open field center time is the amount of time during the entire open field test that the animal spends within a defined center portion of the open field arena. EPM, elevated plus maze; FST, forced swim test; Glur6, glutamate receptor 6; Glura, glutamate receptor A; GR, glucocorticoid receptor; 5-Hht, 5-hydroxytryptamine transporter; ICSS, intracranial self-stimulation.
antidepressants (Table 2). These models have been useful in helping to identify various genes and signaling pathways that may be involved in the development of depressive symptoms and, importantly, in the response to antidepressant drugs. Here we highlight several findings that have produced much interest and stimulated continued research in the field.

The neurotrophic hypothesis of depression posits that growth factors, in particular BDNF, play a key role in regulating mood (38). This hypothesis arose from several observations: first, that the amount of BDNF is decreased in the hippocampus of animals showing abnormalities in depressive- and anxiety-like behavior (80); second, that chronic but not acute antidepressant usage increases BDNF levels (81); and third, that acute application of BDNF, via intrahippocampal injection, has an antidepressant effect (82). However, the neurotrophic hypothesis of depression has been complicated by various genetic manipulations within the BDNF signaling pathway, which show complex, region-specific effects. For example, it has now been shown that contrary to its antidepressant effect in the hippocampus, administration of BDNF in the ventral tegmental area (VTA) results in pro-depressive behaviors (83). Conditional deletion of BDNF in the forebrain does not seem to confer depression per se but does seem to be important in the ability to respond to antidepressants (84). Interestingly, the inability to respond to antidepressants has also been observed in mice that lack adult hippocampal neurogenesis (85).

A number of investigators have produced results suggesting that creation of new neurons in the dentate gyrus of the hippocampus may be important in the ability to respond to antidepressants and that this pathway may be uniquely involved in the regulation of mood-related behavior (86). However, it is clear that the story is not completely straightforward, and further research is necessary to elucidate the effects of BDNF in the etiology of depression, the role of BDNF in the response to antidepressant treatment, and how these effects may be related to adult hippocampal neurogenesis.

Two recent reports of mutant mice resembling behavioral facets of mania have reinvigorated interest in animal models of BPD (Table 1). In the first report, McClung and colleagues found that mutant mice carrying a Clock gene mutation that encodes a dominant-negative CLOCK protein are hyperactive, need less sleep, and show an increased propensity for stimulants and reward (87). These mutant mice exhibit less depressive- and anxiety-like behavior and, remarkably, chronic administration of lithium reduced many of the behavioral features reminiscent of mania. These results are particularly interesting since circadian rhythms and the

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<td><em>Bdnf</em>&lt;sup&gt;+/−&lt;/sup&gt;</td>
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<td><em>Bcl2</em>&lt;sup&gt;−/+&lt;/sup&gt;</td>
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<td><em>Il6</em>&lt;sup&gt;−/−&lt;/sup&gt;</td>
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<td>Transgenic overexpression of TrkB</td>
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Creb, cAMP response element-binding protein; CUS, chronic unpredictable stress; Gnaz, guanine nucleotide binding protein (G protein), αz polypeptide; *Il1r1*, IL-1 receptor, type 1; NIH, novelty-induced hypophagia; TrkB, neurotrophic tyrosine kinase, receptor, type 2; TST, tail suspension test.
genes that underlie the molecular clock have been highly implicated in BPD and because patients with BPD show marked behavioral impairments in their circadian rhythms. A second report of interest concerns the gene encoding glutamate receptor 6 (GLUR6) (101). As GLUR6 resides in a genetic linkage region (6q21) associated with BPD, the demonstration that GLUR6-deficient mice exhibit behavior related to symptoms of mania is very exciting (88). Specifically, GLUR6-deficient mice exhibit less anxious behavior, less risk-taking behavior, less despair-type manifestations, and increased aggressive displays. Further, chronic lithium treatment in these mice reduces hyperactivity, aggression, and some risk-taking behavior (88). Future studies similar to these should begin to shed light on the etiology of BPD as well as provide various genetic and behavioral models that can be used to dissect the underlying neurobiological mechanisms associated with particular deficits in BPD.

Recent studies have identified a molecule that may be associated with affective resilience and enhanced recovery from depressive-like and manic-like states (89). The molecule, Bcl-2-associated athanogene (Bag1), interacts with Hsp70, glucocorticoid receptors (GRs), Bcl-2, and Raf, thereby regulating intracellular signal transduction, nuclear hormone receptors, gene transcription, and cell survival. Microarray studies identified Bag1 as a target for the actions of MSDs (90), and analysis of neuron-selective Bag1 transgenic mice indicated a beneficial role for the protein in depressive-like and manic-like behaviors (102). Specifically, Bag1 transgenic mice showed less anxious-like behavior and had higher spontaneous recovery rates following analysis in the learned helplessness model of stress and/or depression. On mania-related tests, Bag1 transgenic mice recovered much faster in the amphetamine-induced hyperlocomotion test and displayed a clear resistance to cocaine-induced behavioral sensitization.

**Molecular and cellular pharmacologic studies**

Despite substantial advances in pharmacotherapeutics, lithium remains the gold standard therapy for BPD. Valproic acid (VPA), which was first used therapeutically as an anticonvulsive, has become an additional mainstay for treating BPD (36). Current research on understanding the actions of lithium and VPA assumes that these drugs do not have major direct interactions with cell surface receptors and are more likely to exert their effects directly or indirectly through modulation of intracellular targets. Although several targets and mechanisms of these drugs have been identified, precisely which effects are responsible for specific facets of the therapeutic response remains to be fully elucidated. Since the chemical structures of lithium, which is a monovalent cation, and VPA, which is a fatty acid, are highly dissimilar, identifying common targets for these two effective MSDs could be potentially important, as shared biochemical targets would be more likely to represent targets underlying their therapeutic value. It is beyond the scope of this article to review all the molecular and cellular pharmacologic studies; the reader is referred to other reviews (91, 92). Here, we evaluate the most extensively studied and highly prominent targets.

Lithium at therapeutically relevant concentrations functions as a direct inhibitor of a limited number of enzymes because it acts as a competitor for magnesium (93, 94). The enzymes known to be inhibited by lithium include inositol monophosphatase (IMPase); inositol polyphosphate 1-phosphatase (IPPase); bisphosphate 3’-nucleotidase; fructose 1,6-bisphosphatase; GSK3β; and phosphoglucomutase (reviewed in refs. 95–97). The signaling pathways that involve IMPase and GSK3, the phosphoinositol signaling cascade and the Wnt signaling pathway, respectively, have received the greatest attention and interest (Figure 3).

The phosphoinositol signaling cascade mediated by G protein activation of PLC-B is one of the most extensively studied pathways in BPD research (Figure 3A). The direct inhibition of IMPase and IPPase by lithium suggests that it may decrease the bioavailability of myoinositol. This inositol depletion hypothesis postulates that lithium interferes with the regeneration of inositol and, under conditions where inositol limits phosphatidylinositol (PI) synthesis, depletes the cell of PI. Because PI is an obligate precursor of PI biphosphate (PIP2), the hypothesis suggests that inhibition of IMPase could disrupt PIP2/PIP3-mediated signaling (98, 99). Although the inositol depletion hypothesis provides an elegant potential mechanism to explain the actions of lithium, several important issues remain unresolved, and recent data have challenged some of the predilections to the model. For example, therapeutic levels of lithium do not decrease EGF2 in vivo, and this phenomenon is an essential component of the hypothesis (100). Moreover, although the in vivo inhibition of IMPase is incomplete, it is not clear whether this modest reduction in inositol is sufficient to impair PI or PIP2 synthesis (101). Research has therefore moved to more downstream targets in the PI signaling cascade. It has been demonstrated that lithium reduces the levels of specific PKC isozymes in limbic and limbic-related areas of the brain. Notably, VPA also exerts similar attenuating effects on PKC signaling, whereas agents capable of triggering mania exert the opposite effects. One of the genes emerging as a risk factor for BPD in GWASs, DGKH, encodes a protein that is an upstream regulator of PKC signaling. These findings on the PKC pathway have led to the use of tamoxifen, as high concentrations directly inhibit PKC, with promising preliminary results in the treatment of mania (102). These data suggest that direct, centrally acting PKC inhibitors may represent novel therapeutics for the treatment of mania (92, 103).

GSK3 antagonizes both the insulin and Wnt signaling pathways and has recently been demonstrated to regulate synaptic plasticity (104–112). Many of the known effects of lithium can theoretically be explained in terms of GSK3 inhibition. For example, lithium increases neuronal growth cone area (104, 105), alters synaptogenesis (106, 107), and stimulates hippocampal neurogenesis (108). All of these effects mimic Wnt signaling (107, 109, 110). However, inhibition of GSK3 is achieved at the higher end of the therapeutic range of lithium, raising some questions about whether this level of inhibition is sufficient to have strong biological effects. However, secondary inhibition modes have been proposed, which may enhance direct inhibition of GSK3 by lithium (111, 112). Pharmacologic and genetic approaches that regulate GSK3 have behavioral effects that suggest utility in the treatment of BPD. For example, two structurally dissimilar GSK3 small molecule inhibitors and a peptide inhibitor reduce immobility in the forced swim test and attenuate amphetamine-induced hyperactivity (113, 114).

In addition to its direct effects on GSK3, chronic lithium administration at therapeutically relevant concentrations induces prominent neuroprotective and neurotrophic proteins, including Bcl-2 (115) and BDNF (116), in rodents and cultured neurons. Bcl-2 is not only a major antiapoptotic protein, but also stimulates axonal regeneration following injury (reviewed in ref. 117). Consistent with its effects on GSK3, BDNF, and Bcl-2, lithium has been dem-
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onstrated to exert robust neuroprotective properties against various insults both in vitro and in vivo (reviewed in ref. 118). Notably, several independent studies demonstrated that lithium has neuroprotective effects in animal and cellular models of Alzheimer disease, Huntington disease, Parkinson disease, retinal degeneration, spinal cord injury, and HIV infection (119). Finally, human neuroimaging data have shown that chronic lithium treatment in patients with BPD increases NAA levels (a putative marker of neuronal viability) as well as regional gray matter volumes (120, 121). These human data suggest that activation of neurotrophic cascades may underlie the mechanisms by which this simple monovalent cation exerts major effects in the treatment of BPD.

VPA has clear effects on high-frequency sodium channel firing and GABA function, effects that are likely to contribute to

Figure 3
Overview of the PI and Wnt/GSK3 signaling pathways in the neuron. (A) Lithium directly inhibits key enzymes including IPPase and IMPase that regulate inositol-1,4,5-triphosphate (IP3) recycling. IMPase-mediated catalysis represents the final step in the conversion of inositol monophosphate (IMP) to myoinositol. Thus, inhibiting IMPase can reduce myoinositol levels. The myoinositol depletion hypothesis proposes that IMPase inhibition interferes with PI synthesis. The PI signaling cascade starts with surface receptor activation, shown here as GPCR-mediated activation of PLC. Activated PLC catalyzes the hydrolysis of PI biphosphate (PIP2) to diacylglycerol (DAG) and IP3. DAG activates PKC, which, among many other functions, activates myristoylated, alanine-rich C-kinase substrate (MARCKS). The antimanic drugs lithium and VPA both decrease levels of phosphorylated and total MARCKS. Molecules in red depict enzymes, while molecules in blue depict second messengers. PI signaling modulates other second messenger proteins including the small GTPase, Ras, and phosphatidylinositol-4-phosphate (PIP4). (B) Overview of the Wnt and GSK3 signaling pathways. Both lithium and Wnt signaling can inhibit GSK3. In the Wnt signaling pathway, Wnt glycoproteins interact with the frizzled family of receptors to stimulate the dishevelled-mediated (not shown) inactivation of GSK3. Inhibition of GSK3 prevents β-catenin phosphorylation, which inhibits its degradation and allows it to act as a transcription factor. Wnt proteins have been implicated in the regulation of neuron morphology, neurotransmission, and synaptogenesis. GSK3 is also inhibited by Akt (also known as PKB), which is activated downstream of the tropomyosin receptor B (TrkB), whose ligand is BDNF. In addition to inhibiting β-catenin, GSK3 has numerous other targets including microtubule-associated proteins, tau, GSK, and rev-erb α. Adapted with permission from Neuropsychopharmacology (24).
its effects as an antidepressant and potentially to its effects as a mood stabilizer (91). Its ability to function as a histone deacetylase (HDAC) inhibitor has received a great deal of attention, as the role of epigenetic modification and transcription regulation in the control of mood behaviors has garnered increased visibility (122). Recent research has fostered the idea that epigenetic mechanisms, exerting long-lasting control over gene expression, could mediate stable changes in brain function in individuals with BPD. Acetylation of histone tail lysines is a major epigenetic mechanism that is generally associated with transcriptional activation. Recently, Weaver and colleagues showed that the epigenomic state of the GR gene could be established via behavioral programming in rats (123). Remarkably, this epigenomic state could be reversed with an HDAC inhibitor (124). Thus, it has been postulated that early-life environmental stressors contributing to the development of or predisposition to BPD, might be responsive to HDAC inhibitors.

Conclusions and strategies for the future

The large body of literature about the development and underlying neurobiology of BPD indicates a potential role for a multitude of different signaling pathways and neuroendocrine systems. As we have reviewed here, there is an emerging body of evidence suggesting that abnormalities in the regulation of signaling and neural plasticity are integral to the underlying neuropathology of BPD. A correction of dysregulated trans-synaptic signaling by MSDs represents a physiological process that curtails the major oscillations in behavioral states associated with BPD. However, determining which abnormalities actually mediate predisposition to the disease and which are merely factors that are along for the ride seems difficult with so many different pathways and genes being implicated. Although recent research has made great strides toward a better understanding of the illness, continued technological progression in genetic studies and animal models of behavior hold promise for extending our knowledge in the future.

An important question to address in future research concerns the distinction between acute, subchronic, and chronic effects of MSDs and the environmental stressors that serve to predispose to or contribute to illness development. A clear understanding of the timeline under which stressors and MSDs mediate their effects will help guide future studies, enabling researchers to develop hypotheses to test the mechanisms underlying known effects. For example, acute and subchronic effects are most likely to be mediated by regulation of neurotransmission, kinase activity, and enzyme inhibition or facilitation. On the other hand, chronic effects are most likely to be mediated by more stable, long-lasting changes rooted in changes in gene transcription, likely as a result of epigenetics and chromatin remodeling or changes to signaling loops as a result of continual feedback pressure.

As discussed above, a wealth of neuroimaging, neuropathological, biochemical, and behavioral studies over the course of the past decade have suggested numerous candidates for impairments in BPD, including a preponderance of components of various cellular plasticity cascades. These results highlight the potential importance of neuroprotective and neurotrophic effects in the etiology and treatment of BPD. A growing list of genes that confer susceptibility to BPD have been identified, a number of which have been implicated in the cellular or molecular pathophysiology of the disease. These data and the promise of future technological advances hold much promise for the potential discovery of novel therapeutics for BPD.

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