

Glutamate Neurotransmission in Psychotic Disorders and Substance Abuse

Berit Kerner*

Department of Psychiatry and Biobehavioral Sciences, University of California at Los Angeles, Los Angeles, California, USA

Abstract: Psychotropic substance abuse and addiction are very prevalent among individuals with major psychiatric disorders world wide. However, this significant association is poorly explained. The dopaminergic circuits have been implemented in addiction as well as in schizophrenia. Recently the important role of glutamatergic neurotransmission has gained attention and current theoretical models of psychosis and substance abuse support the role of interactions between glutamate and other neurotransmitters in the patho-physiology of both disorders. However, the identification of the underlying genetic risk factors remains challenging and not a single genomic variant has been identified with certainty, possibly due to important limitations of the methods used. Clinical trials with glutamatergic neurotransmission modulators, even though still controversial, support the role of glutamate in psychosis and justify further research.

INTRODUCTION

Psychotropic substance abuse is prevalent among individuals with severe psychiatric disorders, but this comorbidity is poorly explained. Genetic risk factors such as genomic variants, which carry specific vulnerabilities, and environmental risk factors, such as stress or exposure to substances of abuse at certain times during the life span have been proposed in both conditions. The fact that substance abuse is found more frequently among people with severe mental illness than in the general population has even led to the hypothesis that both conditions might be causally related [1]. Substances of abuse, such as phencyclidine (PCP), a non-competitive glutamate receptor antagonist, cocaine, a dopamine-transporter antagonist, or amphetamine, which reverses the transport of dopamine and blocks the dopamine transporter, produce psychotic symptoms often indistinguishable from symptoms seen in bipolar disorder (BPD) or schizophrenia (SZ) [2]. The overlapping and strikingly similar symptoms of these conditions suggest that substance induced psychosis and endogenous delusion in mental disorders may use the same neuro-physiological pathways and may share a common psychopathology. However, only a few studies have actually compared the pathological changes in specific brain regions across disorders [3].

THE DOPAMINE SYSTEM

Evidence for the involvement of the dopamine system in psychosis first came from the observation that dopamine production and release in the nucleus accumbens (NAc) were increased in some unmedicated patients with SZ during the acute stage of the disease [4]. However, variability of this finding suggested that the pathophysiology of SZ might be

heterogeneous. First-generation antipsychotic medications, such as chlorpromazine or haloperidol were found to bind to dopamine receptors, predominantly the dopamine receptor subtype 2 (D2), which is highly expressed in the NAc [5, 6].

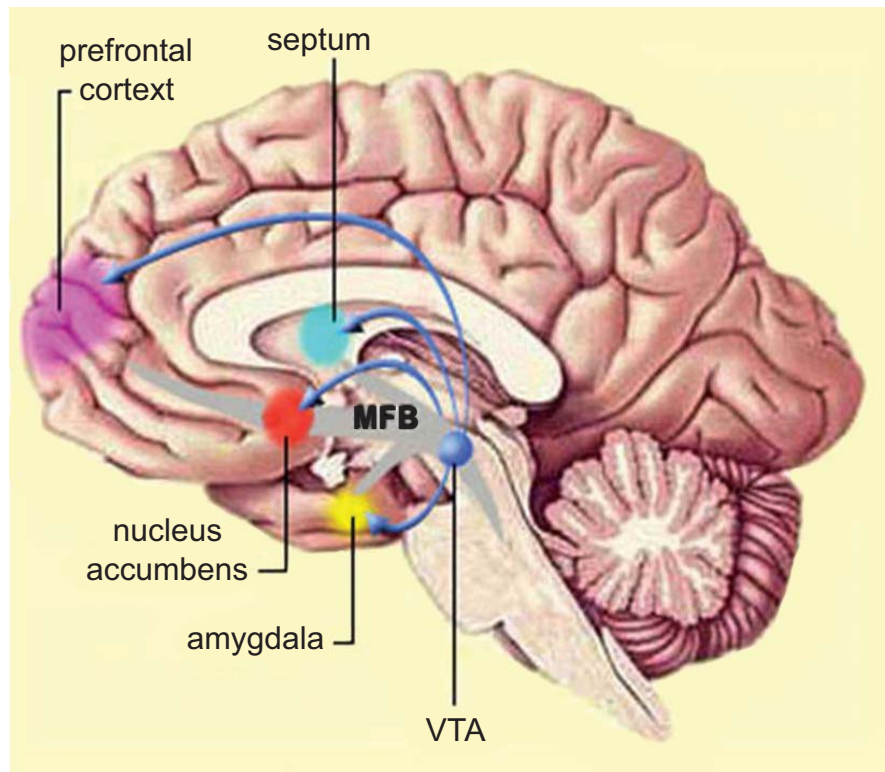
For several decades, the dopamine theory has been the dominant theory in the attempt to explain psychosis. It proclaims that psychotic symptoms arise from imbalances in dopaminergic neuronal pathways, also known as the mesolimbic pathways, which connect the ventral tegmental area (VTA) with the NAc and the prefrontal cortex (PFC) (Fig. 1). The majority of cells in the NAc are gamma-aminobutyric acid (GABA) producing inhibitory neurons that connect the NAc with the thalamus. The VTA is a brain structure involved in reward mechanisms and motivation, whereas the thalamus represents a filter for sensory input to the PFC. Inhibitory pathways from the NAc act on thalamo-cortical connections and protect the cortex from sensory overload and hyper-arousal. Therefore, increased dopamine release in the NAc secondary to imbalances in the mesolimbic pathways could affect GABAergic inhibition. A decrease in GABAergic neurotransmission could then lead to a breakdown of this gating system and subsequent manifestation of psychotic symptoms [7]. The important role of imbalances in the excitatory and inhibitory connections among those central brain regions has been the focus of excellent recent reviews [8] and should therefore not be the topic of this article.

The dopamine system has been extensively studied in addiction as well. Most psychotropic drugs increase dopamine release in the NAc and especially in the initial stages of drug abuse, activation of dopamine cells in the VTA seems to transmit the rewarding effects of drug use. Consequently this effect motivates the repeated use. Recent reviews have focused on dopamine in addiction and the interested reader is referred to these publications [9, 10].

THE GLUTAMATE SYSTEM

Glutamate is the major excitatory neurotransmitter in the brain. It is involved in synaptic plasticity during develop-

*Address correspondence to this author at the Department of Psychiatry and Biobehavioral Sciences, University of California at Los Angeles, 695 Charles E. Young Drive South, Room 3554A, Box 951761, Los Angeles, CA 90095, USA; Tel: (310) 794-9598; Fax: (310) 794-9613; E-mail: bkerner@mednet.ucla.edu



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Fig. (1). This graph of the human brain demonstrates the dopaminergic connections from the ventral tegmental area (VTA) to the amygdala and the nucleus accumbens, as well as to areas located in the septum and the prefrontal cortex. The medial forebrain bundle (MFB) contains the dopaminergic mesocorticolimbic pathway, as well as other ascending and descending neuronal tracts.

ment, memory, learning, and motor activity. The substrate of learning and memory is believed to be encoded in long term potentiations (LTP), which are excitatory currents that lead to changes in synaptic plasticity [10]. Major glutamatergic circuits connect the prefrontal cortex with the NAc and the VTA. The NAc receives also glutamatergic stimulation from the hippocampus and the amygdala, two limbic brain structures that are involved in long-term memory, especially spatial memory and memories of emotional reactions.

Attention to the glutamatergic neurotransmitter system in psychosis came from the observation that PCP, which is a glutamatergic N-methyl-D-aspartate (NMDA) receptor antagonist, could induce psychotic symptoms in humans and addiction-like behavior in animals [11, 12]. Functional magnetic resonance imaging (fMRI) studies in healthy human volunteers revealed that the systemic injection of the NMDA receptor antagonist ketamine led to rapid decrease in the regional blood oxygenation-level dependent (BOLD) signal in the ventromedial frontal cortex. Increased signals were detected in temporal cortical regions and in the thalamus. The glutamate-release inhibitor lamotrigine attenuated the signal [13].

The psychogenic effects of glutamate antagonists are not limited to the NMDA receptor alone. Non-competitive antagonists to the ionotropic (ion channel) NMDA glutamate receptor (such as PCP, MK-801 and ketamine), as well as the alpha-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA) receptor (such as GYKI52466), can induce psy-

chotic symptoms [14, 15]. In addition, kainate, an agonist to a third glutamate receptor subtype can induce imbalances in the dopamine system and psychotic symptoms in humans. The AMPA receptor is a fast, excitatory glutamate receptor and its activation results in depolarization of the synaptic terminal. The tetrameric ion channel consists of a pair of two homo/heteromer dimers, which are composed of GRIA1-4 subunits. The ratio of GRIA1 and GRIA2 receptor subunits determines the Ca^{2+} permeability of the channel. The AMPA receptor insertion into the postsynaptic membrane of excitatory synapses mediates both synaptic strength, as well as structural changes, such as increase in spine size [16]. The receptor is also involved in the homeostatic adaptation of glutamatergic synapses after stimulation for an extended period of time. This process is often referred to as homeostatic synaptic scaling. It involves an increase in the strength of synapses that have had experienced a decrease in activation. On the other hand, a decrease in the synaptic strength occurs after prolonged activation. Therefore, this mechanism ensures that the strength of the individual synapse is kept within a certain optimal range [17].

AMPA receptor trafficking and phosphorylation in the hippocampus have been linked to mania and depression in BPD. Pharmacological studies in mice have found strong effects of anti-manic and anti-depressive drugs on glutamate receptor expression [18, 19]. Chronic treatment of rats with lithium or valproate significantly reduced the GluR1 expression in rat hippocampus. In addition, the GluR1-specific in-

hibitor GYKI52466 reduced amphetamine-induced hyperactivity in these animals [20-22]. This model is often compared to manic symptoms in humans. In contrast, conventional anti-depressants or electroconvulsive therapy (ECT) up-regulate AMPA receptor expression [23, 24]. Psychostimulants, dopamine agonists, and sleep deprivation also increased phosphorylation and/or synaptic levels of GluR1 receptors in rat hippocampus [22]. It could be demonstrated that repeated administration of haloperidol, a first generation anti-psychotic drug, reduced the expression of NMDA receptor subunits, as well as the AMPA receptor subunit GRIA1. This effect was not observed with olanzapine, a second generation anti-psychotic, which might explain the cognitive side effects of first generation anti-psychotics. These experiments support the hypothesis that imbalances, particularly in the AMPA receptor expression might be involved in psychosis and mood disorders.

The NMDA and AMPA glutamate receptors have not only been linked to mania and psychosis, but also to cocaine seeking behavior and addiction in rats [25]. A single dose of cocaine could increase the AMPA receptor expression and change the receptor subunit composition in dopaminergic cells of the VTA [26]. AMPA receptor up-regulation was also observed after repeated administration of other addictive drugs, such as morphine, ethanol, nicotine, and even after stress [27]. On the other hand, the elevated expression of a particular AMPA receptor subunit gene GRIA1 in the NAc shell significantly increased the intracranial self-stimulation threshold in rats [14]. Based on these findings it was hypothesized that substances of abuse could act on the endogenous systems of synaptic plasticity, reward and learning and through re-setting and modification of these systems could increase the vulnerability to relapse [28].

Ionotropic glutamate receptors are not the only receptors that transmit glutamatergic signals in the brain. Metabotropic glutamate receptors (mGluR) that are coupled to G-proteins are located on the postsynaptic site as well as on the presynaptic site of glutamatergic synapses. The mGluRs are involved in the modulation of the glutamatergic signal through coupling with the NMDA receptor or through regulation of glutamate release in a negative feedback loop. The potential role of these receptors in mental disorders has been reviewed recently [29-31]. Even though evidence for the involvement of synaptic plasticity in psychosis and addiction is increasing [32], this view is not without challenges. Antagonists to the glutamate receptors also increase dopamine release in the striatum. Certain antagonists change the expression of D2 receptors in the VTA and even bind to these receptors [33]. Interactions between the glutamate and the dopamine system are prevalent and the specific effects of these neurotransmitter systems are difficult to deduce. The psychogenic actions of NMDA antagonists involve not only the dopamine system. Especially serotonin (5-HT) turnover and release have been detected more consistently than dopamine release after injection of NMDA receptor antagonists. On the other hand, glutamate release in the prefrontal cortex has been induced through the activation of 5-HT_{2A} receptors and antagonists to the metabotropic glutamate 2/3 (mGluR2/3) receptor could block

this release [34]. Selective 5-HT_{2A} receptor agonists, such as mescaline, psilocybin and lysergic acid diethylamide (LSD) can induce psychotic symptoms in humans [35]. The 5-HT_{2A} receptor can bind hallucinogenic and non-hallucinogenic substances. Until recently, it has been unclear, how the specific response induced by these two different ligands could be mediated by the same receptor. However, recent findings have demonstrated that hallucinogenic drugs can induce a unique cellular response mediated through the co-activation of the 5-HT_{2A} receptor and mGluR2/3. The binding of hallucinogenic drugs to the 5-HT_{2A} receptor led to the simultaneous activation of both receptors. This resulted in a particular pattern of G-protein activation and specific changes in gene expression. These changes were distinct from cellular responses after non-hallucinogenic drug binding, which did not involve mGluR2/3 activation [35, 36]. This study demonstrated that 5-HT_{2A} receptors and the mGluR2/3 receptor could form functional complexes in the membranes of cortical neurons. This interaction could then lead to specific changes in ligand affinity and second messenger activation. This finding was the first demonstration of a direct interaction between glutamatergic and serotonergic neurotransmitter systems. It illustrated how crosstalk between receptor subtypes could modify the response to particular ligands.

ANIMAL MODELS

Animal models have traditionally been used to study the biological effect of psychogenic substances and they have consistently provided evidence for the involvement of the glutamate neurotransmitter system in drug abuse and psychosis. A classic example is the injection of PCP, which causes hyperactivity and decreased social interaction in rats. Depression-like behavior, as indicated by immobility in the forced-swim test and impairment in certain cognitive tasks, especially learning and memory have been observed as well. After implantation of electrodes in the prefrontal cortex, ketamine diminished the transmission of the glutamate signal at synapses in the NAc, but it also decreased the neuronal activity of GABA interneurons in the prefrontal cortex. The overall neuronal activity of prefrontal pyramidal neurons increased, possibly because of disinhibition of these neurons [37]. A recent review summarized these results [38].

In addition, sophisticated genetic manipulation in animals has demonstrated the behavioral effect of mutations in some of the target molecules of psychotropic drugs. Knock-out (KO) mice for the AMPA receptor subunit GRIA1 showed normal neuro-anatomy, but abnormal receptor subunit composition in the hippocampus [39, 40]. KO mice for the entire AMPA receptor demonstrated disruption of stimulus-reward learning and deficits in neurocognitive tests, such as spatial working memory, an abnormality frequently found in psychotic patients. Knock-in mice for a phosphorylation site in the GRIA1 receptor subunit demonstrated deficits in synaptic currents, as well as memory deficits in spatial learning tasks [41].

POST MORTEM STUDIES

Region and disease specific alterations in glutamate receptor expression have been repeatedly observed in postmor-

tem brain studies in patients with SZ and to a lesser extent in BPD (Table 1) [42-44]. A recent study found abnormal NMDA receptor subunit composition and alterations in intracellular signaling proteins in the dorsolateral prefrontal cortex of individuals with SZ, BPD and major depressive disorder (MDD) [45]. These results have been replicated in a study, in which laser capture micro-dissection was used to explore gene expression of NMDA and AMPA receptor subunits in Layer II/III and Layer V pyramidal cells in this brain region [46]. In addition, region and disease specific alterations in subunit gene expression of the AMPA receptor subunit GRIA1 was found in medial temporal lobe, and especially in the hippocampus of SZ brains [44, 47]. Immunostaining confirmed reduced protein expression and binding of the AMPA receptor subunit GRIA1. However, studies on glutamate receptor expression and composition in other brain regions have not always been consistent [47-49]. Increased binding has been detected for the kainate receptor in multiple cortical areas and decreased expression was found in the hippocampus. In addition, receptor modifications have been observed affecting calcium influx in the PFC and the striatum. An increase in activity-dependent gene expression has been observed in glutamatergic neurons of the cerebellar cortex in patients with SZ [50, 51]. These studies have supported the role of glutamate neurotransmission in psychotic disorders.

GENETIC FACTORS

Psychosis has been found to aggregate in families [52]. Genetic contributions to the disease are supported by twin studies that found a higher concordance rate for psychotic symptoms in monozygotic twins compared to dizygotic twins [53]. These facts suggest that genetic factors may play a significant role in the psychopathology of psychotic disorders. However, the identification of genomic variants associated with these disorders has been challenging.

Genome-wide linkage studies in large extended families, nuclear families and sib pairs have attempted to identify chromosomal regions that might harbor genomic risk variants for mental disorders [54]. The failure to replicate most of the findings made the interpretation of these signals diffi-

cult. The identified regions might harbor rare genomic abnormalities that segregate in subsets of families and in these particular families, these variants might have a moderate to large effect. Nonetheless, the contributions to the overall phenotype and the frequency in patient populations remain unclear. Interestingly, fine mapping of some of the identified chromosomal regions discovered genomic variants that accumulate in certain pathways, such as the glutamatergic and dopaminergic pathways. They also affected genes involved in neuronal cell migration. The cumulative effect of rare mutations in individual families in certain pathways might explain the high and consistent prevalence of psychiatric disorders.

An alternative approach to the identification of genetic risk factors is the genome-wide case-control association study design with single nucleotide polymorphisms (SNPs) in large population samples. These studies aim to identify genomic regions that may contain common risk variants with small to moderate effects. Even though no single genomic variant has been identified and replicated with genome-wide significance in these studies, some interesting signals occurred in genomic regions harboring genes involved in glutamate neurotransmission [55]. The overall risk of developing a mental disorder associated with individual genomic variances was generally small.

Most recent trends in the hunt for genetic risk factors focus on the identification of small structural genomic variants, such as deletions, inversion, and duplications of genomic regions, as well as copy number variations (CNVs). Several CNVs have been identified in patients with SZ and BPD. Three out of four genes that contained these genomic variations were involved in glutamate signaling [56]. A recent study in a small sample of SZ patients found that the overall frequency of these structural variations was significantly higher in the patient group compared to the control group. The abnormalities affected several genes in neurotransmitter pathways involved in psychosis or drug abuse [57].

Studies on the genetics of substance abuse and addiction have had very similar results. Family, twin and adoption

Table 1. Neurotransmitter Receptor Distribution and Alterations in Postmortem Studies of SZ [44, 48]

Neurotransmitter Receptor	Location	Ligand	Changes in Postmortem Brains (SZ)
NMDA	cortex ventral striatum hippocampus amygdale	PCP ketamine (³ H)MK-801 (³ H)glycine	(↑) increased binding and expression in multiple cortical areas and possibly putamen
AMPA	cortex ventral striatum hippocampus amygdala	AMPA GYKI52466	↑ cortex ↓ hippocampus
Kainate	cortex hippocampus cerebellum	(³ H)kainate	↑ cortex ↓ hippocampus
mGluR2	Cortex Dentate gyrus	LY379268	
mGluR3	Glia	LY379268	

studies have provided strong evidence for heritability of addiction for all subclasses of abused substances, however, no genetic variant associated with substance abuse has been identified with certainty [58]. Well powered genome-wide case-control studies on SNPs and CNVs in addiction will reveal better insight into the patho-mechanisms of these disorders.

LIMITATIONS

The race for a biological explanation of the abnormalities seen in psychiatric disorders has many hurdles. Some technical limitations may explain part of the problem. The currently available genome-wide SNP arrays are designed to detect common genomic variants in high linkage disequilibrium with selected tagging SNPs relatively evenly distributed throughout the genome [59]. This approach can only point to genomic regions that might be associated with a disease, but often fails to identify functional variants. Genomic variants in highly polymorphic regions and in regions with low linkage disequilibrium are often missed. However, it appears to be likely that subtle differences in gene expression levels regulated by genomic variants in highly variable regulatory regions are the root cause of neurophysiologic abnormalities. These regulatory elements can be close to the genes (cis-regulatory elements) or even act over greater distances away from coding regions (trans-regulatory elements). Our knowledge about gene expression regulation is still incomplete. Regulation of gene expression might involve intronic sequences, micro-ribonucleic acids (microRNAs), and intergenic non-coding regions of the genome. The current technologies do not allow us to explore these issues completely. Direct sequencing of candidate gene regions might be required in order to detect genomic variations that might explain psychiatric symptoms.

Overall, genes involved in glutamatergic neurotransmission are not well explored in current studies due to their complex genomic structure. Especially, intron sequences are poorly covered, because these regions are highly polymorphic and rich in copy number repeats. This fact challenges the analysis with currently available methods, as well as the interpretation of the results [60, 61]. For example, in GRIA1, 1394 SNPs can be found in the National Center for Biotechnology Information (NCBI) data base (<http://www.ncbi.nlm.nih.gov/SNP>, November 2008). The majority of these SNPs are located in intron sequences. However, only a very small percentage is represented on the commercially available GeneChip arrays. Those intron sequences may play a major role in gene expression regulation [62, 63]. In addition, evidence is emerging that a major source of variability between two individual genomes is encoded in structural variations, especially deletions, insertions, inversions and duplications of genetic material. The ability to detect these genomic variants with current SNP microarray technology is limited. The cumulative evidence of the importance of structural variants in psychosis and the information gap in genome-wide association studies justify further studies.

CLINICAL STUDIES

Considerable effort has been made to develop specific glutamate receptor antagonists and modulators, based on

accumulating evidence for the importance of glutamatergic neurotransmission in psychotic disorders. Both glutamate agonists, as well as antagonists have been tried in animal models and in humans; however, the results have been inconsistent.

Due to potential neurotoxicity, direct agonists of the NMDA receptor have not been used in clinical trials. Instead, modulators of the glycine binding site of the receptor, such as glycine, D-cycloserine, and D-alanine, have been more promising, especially in combination with typical and atypical antipsychotics [64]. These results, however, have not always been confirmed, and therefore, the enthusiasm about NMDA receptor modulators has declined [65]. Some evidence exists that modulators of the NMDA receptor are more effective in combination with typical anti-psychotics compared to atypical anti-psychotics.

The modulators of the AMPA receptor, such as ampakines, which selectively enhance glutamatergic neurotransmission, have improved cognitive deficits in SZ patients treated with clozapine in some studies [11].

The newest targets in these approaches are modulators of the metabotropic glutamate receptors. Mono-therapy with a selective agonist for the mGluR2/3 has led to significant improvement in both positive, as well as negative symptoms without the common side effects of atypical anti-psychotics [66].

The importance of glutamate in mood disorders has lately drawn attention [67]. The NMDA receptor antagonist ketamine has been reported to have strong and fast acting antidepressant effects [68] and modulators of the AMPA receptor have been shown to be advantageous in the treatment of depression and anxiety [69]. In fact, an extensive review of metabotropic glutamate receptor modulators has indicated their benefit for the treatment of mood and anxiety disorders [70].

New and currently tested anti-psychotic drugs, such as JL 13, might attenuate glutamatergic transmission in the PFC and in the basal ganglia through their effect on the NMDA and AMPA receptors, as well as through interaction with 5-HT_{2A} and α 1-adrenoceptors [71-73].

Even though still controversial, these studies support the role of glutamate in psychosis and justify further research in this important area. Based on these recent findings, new drugs for SZ are under development [74]. However, studies in BPD patients have not been conducted to date.

CONCLUSIONS

Psychiatric disorders are often associated with substance abuse and psychosis. Several theories attempt to explain this correlation. The experience of hallucinations and delusions is, most likely, a common endpoint to a number of neurophysiologic abnormalities. The striking inability of patients with BPD and SZ, however, to maintain balance and the increased vulnerability to addiction may indicate that certain abnormalities in homeostatic mechanisms are at the core of these phenotypes. The fact that substance abuse and dependence are so prevalent in this patient population may not only

be an unfortunate coincidence, but it may point to core pathophysiologic abnormalities in a subgroup of patients. Exploration of these correlations could help to improve our understanding of these psychotic disorders, their genetic risk factors and gene/environment interactions. Most of all, it could open up opportunities for the development of new and improved treatment options, as well as early diagnosis and intervention.

ABBREVIATIONS

PCP	=	Phencyclidine
BPD	=	Bipolar disorder
SZ	=	Schizophrenia
MFB	=	Medial forebrain bundle
NAC	=	Nucleus accumbens
D2	=	Dopamine receptor subtype 2
VTA	=	Ventral tegmental area
PFC	=	Prefrontal cortex
GABA	=	Gamma-aminobutyric acid
LTP	=	Long term potentiations
NMDA	=	N-methyl-D-aspartate
fMRI	=	Functional magnetic resonance imaging
BOLD	=	Regional blood oxygenation level-dependent
AMPA	=	Alpha-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid
GRIA	=	AMPA receptor gene
GRIA1-4	=	AMPA receptor subunit genes 1-4
GluR1	=	Murine glutamate receptor subunit 1
ECT	=	Electroconvulsive therapy
mGluR	=	Metabotropic glutamate receptor
5-HT	=	5-Hydroxy-tryptamine
5-HT _{2A}	=	5-Hydroxy-tryptamine 2A receptor
mGluR2/3	=	Metabotropic glutamate 2/3 receptor
LSD	=	Lysergic acid diethylamide
KO	=	Knock-out
KI	=	Knock-in
MDD	=	Major depressive disorder
SNP	=	Single nucleotide polymorphism
CNV	=	Copy number variation
RNA	=	Ribonucleic acid

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CONFLICT OF INTEREST

The author declares to have no conflict of interest.

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