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## DISEASE-SPECIFIC ALTERATIONS IN GLUTAMATERGIC NEUROTRANSMISSION ON INHIBITORY INTERNEURONS IN THE PREFRONTAL CORTEX IN SCHIZOPHRENIA

Tsung-Ung W. Woo, M.D., Ph.D.<sup>1,2,3</sup>, Amy M. Kim, B.A.<sup>1</sup>, and Emma Viscidi, B.S.<sup>1</sup>

<sup>1</sup>Laboratory for Structural and Molecular Neuroscience, McLean Hospital, Belmont, MA 02478

<sup>2</sup>Department of Psychiatry, Beth Israel Deaconess Medical Center, Boston, MA 02215

<sup>3</sup>Department of Psychiatry, Harvard Medical School, Boston, MA 02115

### Abstract

Glutamatergic modulation of inhibitory interneurons plays a crucial role in shaping the flow of information in the cerebral cortex. In a cohort of postmortem human brains from schizophrenia (n=20), bipolar disorder (n=20) and normal control (n=20) subjects, we colocalized the mRNA for the N-methyl-D-aspartate (NMDA) receptor NR2A subunit, labeled with [<sup>35</sup>S], and the mRNA for the  $\gamma$ -aminobutyric acid (GABA) synthesizing enzyme glutamic acid decarboxylase (GAD)<sub>67</sub>, labeled with digoxigenin. We found that the density of GAD<sub>67</sub><sup>+</sup> neurons in layers 2–5 of the prefrontal cortex was decreased by 27–36 % in both schizophrenia and bipolar disorder. In addition, the density of the GAD<sub>67</sub><sup>+</sup>/NR2A<sup>+</sup> neurons was decreased by 57 % and 49 % in layers 3 and 4, respectively, in schizophrenia, but it was unchanged in bipolar disorder. These findings raise the possibility that glutamatergic innervation of inhibitory interneurons via the NMDA receptor in the prefrontal cortex may be selectively altered in schizophrenia.

### 1. INTRODUCTION

Normal neurocognitive functioning is contingent upon the integrity of information processing in the cerebral cortex, with specific cortical areas contributing differentially to various aspects of cognition. For instance, the prefrontal cortex (PFC) plays an important role in the temporal organization of behavior, or executive functioning, via working memory [20,22]. Working memory is the ability to temporarily maintain “on-line” internal representations of information in the perceptual, cognitive, and emotive domains that are no longer immediately present, for a brief period of up to tens of seconds, in order to guide future behavior [5]. In other words, the integrity of working memory is critical for the sequential execution of motor or cognitive acts in a goal-directed manner; this capacity forms the basis of a variety of normal daily human activities, such as planning, reasoning, thinking and language. Patients with schizophrenia exhibit impairment in the performance of many of these activities [8]. In fact, working memory and executive functioning deficits are thought to represent core pathophysiologic or perhaps endophenotypic features of schizophrenia [22,41,49]. Although disturbances of the prefrontal

Send correspondence to: Dr. Tsung-Ung W. Woo, McLean Hospital, 115 Mill Street, Belmont, MA 02478, Tel: 617-855-2823, FAX: 617-855-3199, Email: wwoo@hms.harvard.edu.

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functional architecture have also been implicated in bipolar disorder [53], working memory, as measured by the delayed response paradigm, seems to be relatively intact in this illness [40]. Thus, the pathophysiologic nature of neural circuitry disturbances within the PFC may be dissimilar in the two conditions.

Inhibitory interneurons that utilize GABA ( $\gamma$ -aminobutyric acid) as a neurotransmitter play a crucial role in the maintenance of sustained neuronal activation during working memory by dynamically adjusting the conductances of the pyramidal neuronal network. Interestingly, the number of N-methyl-D-aspartate (NMDA) glutamate receptors on GABA neurons appears to be an important determinant of the stability of the network in sustaining working memory [34,54]. GABA neurons receive feedback excitatory modulation via local recurrent excitatory projections from the pyramidal neurons they innervate and, at the same time, they are also targets of feedforward excitatory modulation from axonal projections furnished by other pyramidal neurons, located both within the PFC and in other cortical areas [6,36–38]. Furthermore, projections from the thalamus provide additional excitatory drive to GABA cells [63]. The integrity of working memory and associated functions, such as executive control, depends on the complex interplay of feedback and feedforward mechanisms of modulation of cortical inhibitory activities via activation of glutamate receptors on GABA neurons [11,12, 46], which helps to regulate the temporal flow of information between spatially distributed populations of pyramidal neurons in a contextually meaningful manner.

Converging lines of evidence strongly suggest that, in the PFC, functional disturbances of GABA neurons represent a prominent pathophysiologic feature of schizophrenia [1,2,7,13, 33,47] and bipolar disorder [24,56,57]. In addition, alterations of glutamatergic modulation of GABA cells could further compromise PFC functions. In fact, in a recent study, we found that in the anterior cingulate cortex the density of GABA cells that expressed the NMDA NR2A subunit was decreased in schizophrenia and bipolar disorder [65]. It is currently unknown whether similar changes also occur in other brain regions, such as the PFC. In this study, we used double in situ hybridization to co-localize the mRNA for the NMDA NR2A subunit and that for the GABA synthesizing enzyme glutamic acid decarboxylase (GAD)<sub>67</sub> in the PFC in a cohort of 60 human subjects (Figure 1). We found that the pattern of changes in the density of GABA neurons was remarkably similar between schizophrenia and bipolar disorder, but the density of those GABA cells that expressed NR2A was altered in a diagnosis- and laminar-specific manner.

## 2. RESULTS

### 2.1. GAD<sub>67</sub> mRNA-Expressing Neurons

Overall, our analysis revealed a significant diagnosis effect ( $F_{2, 57}=11.71$ ,  $p<0.0001$ ; Figure 2). Post-hoc analyses further revealed that, in the schizophrenia subjects, the density of GAD<sub>67</sub> mRNA-expressing neurons in layers 2 ( $t=3.37$ ,  $p=0.002$ ), 3 ( $t=3.22$ ,  $p=0.003$ ), 4 ( $t=2.66$ ,  $p=0.01$ ) and 5 ( $t=2.91$ ,  $p=0.006$ ) was significantly decreased by 30.2%, 33.2%, 32.2% and 35.9% respectively, compared to the normal control subjects (Figure 2).

In the subjects with bipolar disorder, the pattern and magnitude of changes in cell density was virtually identical to what was observed in the schizophrenia subjects (Figure 2). Thus, the density of GAD<sub>67</sub> mRNA-expressing neurons was significantly decreased by 33.6% and 33.2% in layers 3 ( $t=2.99$ ,  $p=0.005$ ) and 4 ( $t=2.63$ ,  $p=0.01$ ), respectively. The density of these neurons also appears to be decreased by 26.7% and 32.0% in layers 2 ( $t=2.13$ ,  $p=0.04$ ) and 5 ( $t=1.94$ ,  $p=0.05$ ), respectively, but these differences did not achieve statistical significance after correction for repeated measures. Because a small number ( $N=4$ ) of the subjects with bipolar disorder had no history of psychosis, we also analyzed our data with the exclusion of these cases but this did not alter our results.

## 2.2. NR2A mRNA-Expressing GABA Neurons

A significant effect of diagnosis was detected by our ANOVA model ( $F_{2, 57}=4.38$ ,  $p=0.017$ ; Figure 2). During layer-by-layer post hoc analyses, we found that the density of the double-labeled neurons was decreased by 56.8% and 49.4% in layers 3 ( $t=2.36$ ,  $p=0.023$ ) and 4 ( $t=4.10$ ,  $p=0.0002$ ), respectively, in the schizophrenia subjects, although the reduction in layer 3 did not reach statistical significance after the stringent Bonferroni correction. Cell density also appeared to be decreased by 27% in layer 2 in the subjects with schizophrenia, but this difference was not statistically significant. In the subjects with bipolar disorder, we did not observe any statistically significant differences in neuronal density in any of the cortical layers. Exclusion of the 4 subjects with no history of psychosis also did not affect our results.

## 2.3. Cortical Thickness Measurements

The average cortical thickness ( $\pm$ S.D.) did not differ ( $F_{2, 57}=0.46$ ,  $p=0.63$ ) between the schizophrenia ( $1,755.0\pm 386.5$  mm), bipolar ( $1,863.6\pm 395.1$  mm) and normal control ( $1,766.8\pm 395.5$  mm) groups, suggesting that our observations of reduction in the density of both the single- and double-labeled neurons cannot be explained by differential tissue shrinkage in the 3 subject groups.

## 2.4. Expression Level of NR2A mRNA in GAD<sub>67</sub> mRNA-Expressing Neurons

The distributions of the frequency histograms of NR2A mRNA expression level per GAD<sub>67</sub>+ cell are shown in Figure 3. The Kruskal-Wallis test revealed that the effect of diagnosis on grain density was not statistically significant ( $H=5.05$ ,  $p=0.08$ ). Layer-by-layer analyses also failed to reveal any diagnosis effect (data not shown). Finally, the average size of the silver grain clusters also did not differ between the 3 groups.

## 2.5. Potential Confounding Variables

Within individual subject groups and when subject groups were combined, we found no evidence for any statistically significant correlation between any of the continuous variables and neuronal density measures or NR2A mRNA expression level. A small number of subjects with schizophrenia ( $n=4$ ) or bipolar disorder ( $n=5$ ) were not on any antipsychotic medications for a period of between 2 weeks and up to years before the time of death. In these subjects, their neuronal densities did not appear to be appreciably higher than neuronal densities in the subjects who were on these medications. Therefore, the observed changes in densities do not appear to be the result of antipsychotic treatment. Similarly, we found no evidence for any correlation between the dosage of divalproex and any of the density measures or NR2A mRNA expression level, suggesting that treatment with this mood stabilizer did not confound our findings. Due to sample size limitations, potential effects of lithium or other mood stabilizers could not be meaningfully evaluated and for the very same reason these medications would not have confounded our results. Finally, no statistically significant differences in neuronal densities or NR2A mRNA expression level were observed between the two sexes or the two hemispheres, either when individual subject groups were analyzed separately or when cases from the control group were combined with those from the schizophrenia or bipolar group.

## 3. DISCUSSION

Several conclusions can be drawn from the present study. First, the density of GAD<sub>67</sub> mRNA-containing neurons appears to be decreased in layers 2–5 of the PFC in schizophrenia. This observation replicates previous findings from several other laboratories [2,24,58] and adds to the consensus that disturbances of GABA neurotransmission represent a key pathophysiologic feature of schizophrenia [1,7,13,33]. Second, the density of GAD<sub>67</sub> mRNA-containing neurons in the PFC also appears to be decreased in bipolar disorder. Furthermore, the laminar pattern

of reduction is remarkably similar to that in schizophrenia. Third, the density of GABA neurons that express the NMDA NR2A subunit appears to be decreased in the middle cortical layers in schizophrenia, whereas the density of these neurons in bipolar disorder seems to be unaltered. Thus, in the PFC, glutamatergic innervation of GABA interneurons via NMDA channels may be deficient in schizophrenia, but not in bipolar disorder.

### 3.1. Methodological Considerations

Consistent with previous studies [2,24,58], we found that in 27–36% of the GABA cells, the expression of the mRNA for GAD<sub>67</sub> appears to be reduced to a level that is experimentally undetectable in subjects with schizophrenia. Furthermore, NR2A mRNA expression in the GABA cells with unaltered GAD<sub>67</sub> expression is significantly reduced. A very important question that cannot be addressed in the present study, however, is whether NR2A expression is also reduced in the GABA cells with undetectable GAD<sub>67</sub>. Because the expression of the 65 kD-isoform of GAD (GAD<sub>65</sub>) is not decreased in schizophrenia [24], this question can be indirectly addressed in future studies by comparing the expression of NR2A in GAD<sub>67</sub>- versus GAD<sub>65</sub>-expressing neurons.

In the quantification of GAD<sub>67</sub><sup>+</sup>/NR2A<sup>+</sup> cells, we included only those with the silver grain density that was 2X above background. In fact, the exclusive majority (>93%) of the double-labeled cells that were counted met this criterion. However, it is possible that the 7% of the cells that were excluded may have contained a small population of GABA cells with very low NR2A expression level. Nevertheless, the possible exclusion of this small contingent of cells should not have affected our conclusions, given the magnitude of change in the density of the double-labeled cells was observed to be in the 50–60% range. Conversely, we cannot rule out the possibility that some of the silver grain clusters included in our analysis may represent non-specific labeling. In this case, we would have underestimated the magnitude of reduction in cell density, but our conclusions should not have been affected.

In any postmortem human brain studies, medications pose a serious potential confound. For instance, in this study, the exclusive majority of the subjects were on psychiatric medications at the time of death, which include antipsychotic drugs in both the schizophrenia and bipolar subjects and mood stabilizers in the bipolar subjects. As described above, our statistical analyses do not support the notion that the observed reduction in the densities of both the GAD<sub>67</sub><sup>+</sup> and GAD<sub>67</sub><sup>+</sup>/NR2A<sup>+</sup> cells is the result of treatment with antipsychotics or mood stabilizers. Furthermore, it has been found that, in monkeys, long-term treatment with haloperidol does not appear to have any effects on the level of GAD<sub>67</sub> mRNA expression [58]. Also, the fact that the reduction in the density of GAD<sub>67</sub><sup>+</sup>/NR2A<sup>+</sup> cells occurred only in the schizophrenia group even though the majority (N=13) of the subjects with bipolar disorder were also receiving antipsychotic medications seems to argue against the idea that this observation represents the result of antipsychotic treatment. Lastly, many (N=11) of the subjects with bipolar disorder were on a combination of antipsychotics and mood stabilizers. When comparing the density of both the single- and double-labeled cells between these subjects and the rest of the subjects, the two groups did not appear to show any differences, suggesting that the combined use of antipsychotics and mood stabilizers does not appear to have affected our results.

Finally, cigarette smoking is exceedingly prevalent in patients with schizophrenia [9,15]. Unfortunately, the demographic information that is available did not allow us to systematically address this important potential confound. Nevertheless, previous studies in animals suggest that nicotine may actually increase the expression of NR2A in different brain regions, including the cerebral cortex [17,28,59]. If these findings apply to the human brain, we might have actually underestimated the magnitude of NR2A expression reduction in subjects with schizophrenia.

### 3.2. Disturbances of Inhibitory Interneurons in Schizophrenia and Bipolar Disorder

Our observations are in line with previous studies showing that the density of GAD<sub>67</sub> mRNA-expressing neurons appears to be decreased in the PFC [2,24,58] and in the anterior cingulate cortex [65] in schizophrenia. Together, these findings provide clear support for the growing understanding that GABAergic modulation of pyramidal cell activities in the cerebral cortex seems to be significantly altered in this disorder [7,13,33]. There has also been increasing evidence implicating GABA neurotransmission disturbances in bipolar disorder. For instance, using quantitative real-time reverse transcriptase polymerase chain reaction, it was found that the mRNA for GAD<sub>67</sub> in homogenized PFC tissues was significantly decreased by up to 50% in the PFC in subjects with bipolar disorder [24]. In the present study, we found that the density of GAD<sub>67</sub> mRNA-expressing neurons was decreased in layers 2–5; this pattern was virtually identical to what was observed in schizophrenia and is also consistent with the observations of 2 previous studies [24,57]. Therefore, it appears that disturbances of GABA interneurons in the PFC may represent a common pathophysiological element in both schizophrenia and bipolar disorder, although the neurobiological mechanisms that underlie these disturbances remain poorly understood. Because GABA cells are morphologically, neurochemically and functionally heterogeneous, with subpopulations of these neurons involved in different aspects of information processing in the cerebral cortex [16,25,29,35,61], a critical question that will need to be addressed in future studies is whether and if so how subpopulations of GABA interneurons may be differentially affected in the two conditions.

### 3.3. Disturbances of Glutamatergic Neurotransmission on Inhibitory Interneurons in Schizophrenia

Our observation suggests that glutamatergic innervation of GABA interneurons via NMDA receptors may be decreased in layers 3–4 of the PFC. This observation may at first glance appear to be at odds with previous findings [3] suggesting that NR2A expression in the PFC seems to be unaltered in schizophrenia. However, because GABA cells comprise only about 20 % of all neurons in the PFC, our observation of an approximately 50% decrease in NR2A transcript expressed by GABA cells suggests that NR2A transcript signal is decreased by only about 10% among all of the neurons. If we also take into consideration the amount of neuropil in the PFC, this signal-to-noise ratio will be further diluted and is unlikely to be detectable by film autoradiography [3].

The observed decrease in the density of NR2A+/GAD<sub>67</sub>+ cells in layers 3 and 4 of the PFC without an apparent decrease in the level of NR2A mRNA expression per GAD<sub>67</sub>-expressing cells suggests that NR2A mRNA expression in a subpopulation of GABA cells is reduced to an undetectable level. The origin of the glutamatergic terminals that innervate the GABA cells that exhibit decreased NR2A mRNA expression is at present unknown. However, it is known that deep layer 3 and layer 4 together represent the thalamocortical recipient zone in the PFC. In fact, there has been evidence suggesting that thalamocortical inputs to the PFC may be deficient in schizophrenia [4,21,32,39,42,43]. Furthermore, GABA cells in these layers, especially those that contain the calcium binding protein parvalbumin (PV), are known to be targeted by thalamocortical inputs [48,63]. Taken together, our findings are consistent with the idea that thalamocortical innervation of GABA neurons in the PFC may be aberrant in schizophrenia.

Pyramidal neurons in layer 3 of the PFC furnish the intrinsic excitatory circuit [44,64] and approximately 50% of the axon terminals arising from this circuit target GABA neurons [36]. Thus, these terminals may also be a source of the aberrant glutamatergic innervation of the GABA cells that exhibit reduced NR2A expression. It is postulated that this circuit may mediate sustained neuronal activation during working memory through reentrant oscillation of spatially-distributed functionally-segregated pyramidal cell ensembles [23,60]. Because



GABA interneurons play a crucial role in regulating the flow of information within and between these pyramidal cell ensembles, altered glutamatergic regulation of GABA neurons may compromise the precise spatiotemporal orchestration that is necessary for reentrant oscillatory dynamics [27,31], resulting in working memory and associated deficits in schizophrenia. In addition, thalamocortical projections may further contribute to reentrant oscillations of the intrinsic excitatory circuit [52]. As such, deficient thalamocortical modulation of GABA neurotransmission could further compromise the functional integrity of this circuit. Interestingly, the generation of neuronal oscillations, especially those in the gamma frequency band, has recently been shown to be disturbed in schizophrenia [10,50,51,55]. Furthermore, it is known that the fast-spiking GABA cells that contain PV are particularly important in mediating gamma band oscillations [14,19,62]. In addition, compared to pyramidal cells and other GABA cells, PV-containing GABA cells contain up to 5-fold the amount of NR2A at both the protein and transcript levels [30]. Taking together, these observations raise an interesting possibility that the PV-containing GABA cells may be among those GABA cells that exhibit drastically reduced NR2A mRNA expression.

### 3.4. Regional Similarities and Differences in Inhibitory Neural Circuitry Disturbances in Schizophrenia and Bipolar Disorder

The findings that the density of GAD<sub>67</sub> mRNA-expressing neurons is reduced in schizophrenia and bipolar disorder in both the PFC [2,24,58] and the anterior cingulate cortex [65] suggest that, in both of these conditions, GABA neurotransmission is compromised across cortical regions. However, while glutamatergic drive to GABA neurons via the NMDA receptor appears to be altered in both schizophrenia and bipolar disorder in the anterior cingulate cortex [65], in the PFC, this alteration may be specific to schizophrenia. This finding is consistent with the idea that NMDA receptors on GABA cells play a key role in regulating sustained neuronal activation [34,54], which forms the physiological substrate of working memory, and that working memory may be preferentially impaired in schizophrenia [40]. In future studies, it will be essential to more precisely delineate the similarities and differences in neural circuitry disturbances in these two cortical regions between schizophrenia and bipolar disorder in terms of cellular and molecular impairments within microcircuits.

## 4. EXPERIMENTAL PROCEDURE

### 4.1. Subjects

Sixty postmortem human brains from 20 subjects with schizophrenia, 20 subjects with bipolar disorder and 20 normal control subjects were obtained from the Harvard Brain Tissue Resource Center at McLean Hospital. As shown in Table 1, the 3 subject groups were closely matched for age and postmortem interval (PMI). In addition, the mean freezer storage time (days±S.D.) of brains was not significantly different ( $F_{2, 57}=1.32$ ,  $p=0.28$ ) between the normal control (1,822±997), schizophrenia (1,755±878) and bipolar (1,376±931) groups. The mean tissue pH (±S.D.) also did not differ ( $F_{2, 57}=0.77$ ,  $p=0.47$ ) between the three groups of subjects (normal control: 6.58±0.25; schizophrenia: 6.53±0.28; bipolar: 6.47±0.22).

Psychiatric diagnoses were established using a retrospective review of medical records and an extensive family questionnaire that included the medical, psychiatric, and social history of the subjects. For the diagnosis of schizophrenia, the criteria of Feighner et al [18] were used and the diagnoses of schizoaffective and bipolar disorder were made according to DSM-III-R criteria. The majority (N=16) of the subjects with bipolar disorder had a history of psychosis, even though they may not have been on antipsychotic medications at the time of death. The medication status of all of the cases can be found in Table 1. None of the subjects were suffering from any substance abuse/dependence disorder at the time of death, as confirmed by review of toxicology reports, medical records and family questionnaires.

## 4.2. Tissue Preparation

Human brains were blocked and blocks containing Brodmann's area 9 of the dorsolateral PFC were frozen over liquid nitrogen vapor before being stored at  $-70^{\circ}\text{C}$ . In preparation for in situ hybridization, tissue blocks were sectioned at a thickness of  $10\ \mu\text{m}$  on a cryostat. Two sections per subject and therefore six sections per matched triplet were used for the experiment. The six sections from each triplet were mounted on three slides as follows: 1) normal control +schizophrenia, 2) bipolar+normal control, and 3) schizophrenia+bipolar. This method of mounting sections controls for potential variability of hybridization signals due to, among other factors, variability in the thicknesses of emulsion between slides and possibly also on the same slides. All slides were processed at the same time in one single experiment.

## 4.3. Dual *in situ* Hybridization

### 4.3.1. Riboprobe Preparation

**4.3.1.1. Radiolabeled cRNA Probe for NR2A mRNA:** The cRNA probes for the NR2A subunit (kindly provided by Dr. Christine Konradi, Vanderbilt University) were transcribed in vitro from linearized cDNA subclones encoding the rat NMDA NR2A subunit, as previously described [65]. The specificity of the probe had been verified by northern blot analysis. The probe was derived from a cDNA spanning nucleotides 1185–2154 (Genbank Accession #M91561) within the coding region of the subunit. A corresponding sense probe was used as a control. Radiolabeled cRNA probe was prepared by first drying down [ $^{35}\text{S}$ ]UTP ( $500\ \mu\text{Ci}/\text{ml}$  of probe, New England Nuclear, Boston, MA) in a DNA Speed-Vac (Savant, Farmingdale, NY).  $100\ \text{ng}/\mu\text{l}$  of the cDNA template,  $0.1\ \text{M}$  dithiothreitol (DTT),  $3\ \text{U}/\mu\text{l}$  RNasin,  $5\ \text{mM}$  NTPs,  $0.8\ \text{U}/\mu\text{l}$  of T3 or T7 RNA polymerases (for antisense and sense probe, respectively), and  $5\text{X}$  transcription buffer were then added. The transcription mixture was subsequently incubated at  $37^{\circ}\text{C}$  for 1 hour. The cDNA template was digested by incubating the mixture with R1Q DNase at  $37^{\circ}\text{C}$  for 15 minutes. Unincorporated NTPs were removed by running the mixture through a Stratagene Nuc-Trap (La Jolla, CA) push column. The eluate was collected and probe concentration was determined by scintillation counting. The probe was then stored at  $-20^{\circ}\text{C}$  until use.

**4.3.1.2. DIG-Labeled GAD<sub>67</sub> mRNA Probe:** DIG-UTP-labeled cRNA probes were transcribed using  $100\ \text{ng}$  of full-length, linearized human cDNA clones inserted in a bluescript vector (kindly provided by Drs. Allan Tobin and Niranjula Tillakaratne, UCLA) in the presence of  $0.1\ \text{M}$  DTT,  $3\ \text{U}/\mu\text{l}$  Rnasin,  $0.8\ \text{U}/\mu\text{l}$  of T3 and T7 RNA polymerases,  $10\ \text{mM}$  of ATP, CTP, and GTP,  $6.5\ \text{mM}$  of UTP, and  $3.5\ \text{mM}$  of DIG-labeled UTP (Boehringer Mannheim, Indianapolis, IN). The mixture was incubated at  $37^{\circ}\text{C}$  for 1 hour. cDNA template was digested with RQ1 DNase. This same cRNA probe has been used in 2 previous postmortem studies from our laboratory [26,65].

**4.3.2. Hybridization—**To ensure adequate tissue penetration, the GAD<sub>67</sub> probe was hydrolyzed to  $0.8\ \text{kb}$  with an equal volume of sodium bicarbonate/carbonate buffer ( $\text{pH } 10.2$ ;  $40\ \text{mM}$   $\text{NaHCO}_3$ ,  $60\ \text{mM}$   $\text{Na}_2\text{CO}_3$ ) at  $60^{\circ}\text{C}$  for 6–10 minutes. Probes were then reconstituted in a hybridization buffer consisting of 50% formamide, 0.1% yeast tRNA, 10% dextran sulfate, 1X Dehardt's solution,  $0.5\ \text{M}$  EDTA, 0.02% SDS, 4X SSC,  $10\ \text{mM}$  DTT, and 0.1% ssDNA, at a final concentration of  $0.4\ \text{ng probe}/\mu\text{l}$  hybridization buffer. Before hybridization, mounted tissue sections were air dried and warmed to room temperature. They were then post-fixed in 4% paraformaldehyde for 10 minutes and incubated in  $0.1\ \text{M}$  TEA for 5 minutes at room temperature before being dehydrated in a graded series of ethanol. Probes were then added to slides for hybridization in a prewarmed, humidified dish. Sections were covered with coverslips and incubated at  $55^{\circ}\text{C}$  for 12 hours. At the end of hybridization, coverslips were soaked off in 4X SSC in the presence of  $100\ \mu\text{l}$  of 2-mercaptoethanol. Tissue was then incubated in  $0.5$

M NaCl/0.05 M PB for 10 minutes, 0.5 M NaCl with 0.025 mg/ml RNaseA at 37 °C for 30 minutes, followed by a high stringency wash with a solution containing 50% formamide, 0.5 M NaCl/0.05 M PB, and 100 µl 2-mercaptoethanol at 63 °C for 30 minutes. Sections were finally washed overnight in 0.5X SSC with 20 mM 2-mercaptoethanol at room temperature.

**4.3.3. Visualization of DIG Labeling**—After incubation in blocking buffer (100 mM Tris-HCL, 150 mM NaCl pH 7.5, 3% normal goat serum (NGS), 0.3% Triton X100), sections were incubated overnight at 4 °C in buffer containing 1:200 dilution of alkaline phosphatase-conjugated sheep anti-DIG antibody (Roche Diagnostics, Indianapolis, IN). Sections were then incubated in Vector Red™, which was prepared using the Vector Red™ alkaline phosphatase substrate kit (Vector Laboratories, Burlingame, CA), at room temperature for 40 minutes in complete dark.

**4.3.4. Emulsion Autoradiography**—It was determined that sufficient autoradiographic signal had developed after the slides were apposed to X-ray film (Kodak Biomax MS) for 10 days. The slides were then dipped in emulsion (Kodak NTB-2), air dried, and stored at 4 °C in darkness for 5 weeks. After development in the dark with Kodak D-19 developer, slides were counterstained with cresyl violet and coverslipped.

#### 4.4. Quantification of GAD<sub>67</sub> and NR2A mRNA Expression

All microscopic analyses were conducted by one investigator (EV) under strictly blind condition. [<sup>35</sup>S]-labeling of NR2A mRNA appeared as clusters of silver grains after processing for emulsion autoradiography. DIG labeling was visualized as a red-brown reaction product under a brightfield microscope equipped with polarizing filters to enhance the optical density of the reaction product (Figure 1). Neurons that were single-labeled with DIG and those that were double-labeled with DIG and [<sup>35</sup>S] were identified using a Leica Laborlux microscope equipped with a solid state CCD video camera and Bioquant Nova Image Analysis System (R&M Biometrics, Memphis, TN). Using an 100X oil immersion objective lens at a final magnification of 1,000X, the distribution of both single- and double-labeled neurons within a 250 µm-wide column extending from the pial surface to the border between layer 6 and the subcortical white matter were obtained for each section. A total of 120 sections (i.e. 2 sections per subject) were quantified. For the DIG + [<sup>35</sup>S] double-labeled cells, only those that had a grain density greater than 2x of background grain density (see below) were included. Neighboring sections were stained with cresyl violet for accurate determination of laminar boundaries. All sampling columns were placed within Brodmann's area 9, which was identified based on known cytoarchitectural criteria [45]. Densities of single- and double-labeled neurons for each cortical layer were then obtained by dividing cell counts by laminar areas. For each subject, density measures from the 2 sections were pooled to give rise to an average measure. Intra-rater reliability, as assessed by counting and recounting profiles within the same column, was established to be consistently between 93–95 % before the actual data collection process was begun and throughout the course of data collection. Quantification was performed on a regular and continuous basis in order to minimize any deviation in the data collection routines. The entire data quantification process was completed in four months.

To quantify the expression level of mRNA for the NR2A subunit in individual GABA cells, the area occupied by each silver grain cluster was outlined using a cursor displayed on the computer monitor. The cluster area was measured by highlighting the grains with a thresholding subroutine. The light intensity was then adjusted to ensure that the size of the grains was neither under- nor over-represented and both the threshold and the light intensity were held constant throughout the entire data collection process. The area covered by autoradiographic grains within the cluster area was automatically computed based on the threshold value and was represented as a pixel count for NR2A transcript expression level. The pixel count was



expressed as a function of cluster area. By subtracting the background grain density (i.e. pixel count of the area covered by autoradiographic grains per unit area in  $\mu\text{m}^2$  in the white matter), the corrected NR2A expression level was obtained. The average NR2A expression level in GABA interneurons (i.e. cells positive for GAD<sub>67</sub> mRNA) for each cortical layer for each case was then computed. For each diagnosis group, a histogram of the distribution of grain density was plotted.

#### 4.5. Statistical Analyses

The densities of GAD<sub>67</sub> mRNA+ and GAD<sub>67</sub> mRNA+/NR2A mRNA+ neurons were compared between the 3 subject groups across layers 2 through 6, using a repeated-measures analysis of variance (ANOVA), with diagnosis as the between-groups factor, layer as the within-group factor, and repeated measures on layer. For post-hoc testing of differences in group means, two-tailed unpaired t tests were used. Layer 1 was excluded in our analyses because of the very small number of GAD<sub>67</sub> mRNA-containing neurons that co-expressed NR2A mRNA in this lamina. The Bonferroni procedure was employed to correct for type 1 error as a result of multiple comparisons (i.e. layers 2, 3, 4, 5, and 6). Therefore, the alpha level for significance for all t tests was  $p=0.01$  (i.e.  $0.05/5$ ). In addition, in order to evaluate the effects of potential confounding factors, for continuous variables such as age, PMI, pH, freezer storage time, and exposure to antipsychotic medications or mood stabilizers, simple Pearson correlations were obtained for the individual groups and when cases from the control group were combined with those from the schizophrenia and the bipolar group, respectively. In addition, we used analysis of covariance (ANCOVA) to understand how these confounding variables might have affected our results. Because none of the conclusions derived from our findings were affected by the ANCOVAs, only results from repeated-measures ANOVAs are reported. Potential effects of nominal variables such as hemispheric laterality and sex on our findings were evaluated by using two-tailed unpaired t-tests to compare the density and expression level measures from the two hemispheres and those from the two sexes, both within individual groups and when cases from the control group were combined with those from the schizophrenia and bipolar group, respectively. To detect any differences in the expression of the mRNA for NR2A in GAD<sub>67</sub> mRNA-containing neurons, the non-parametric Kruskal-Wallis test was used.

#### ACKNOWLEDGEMENT

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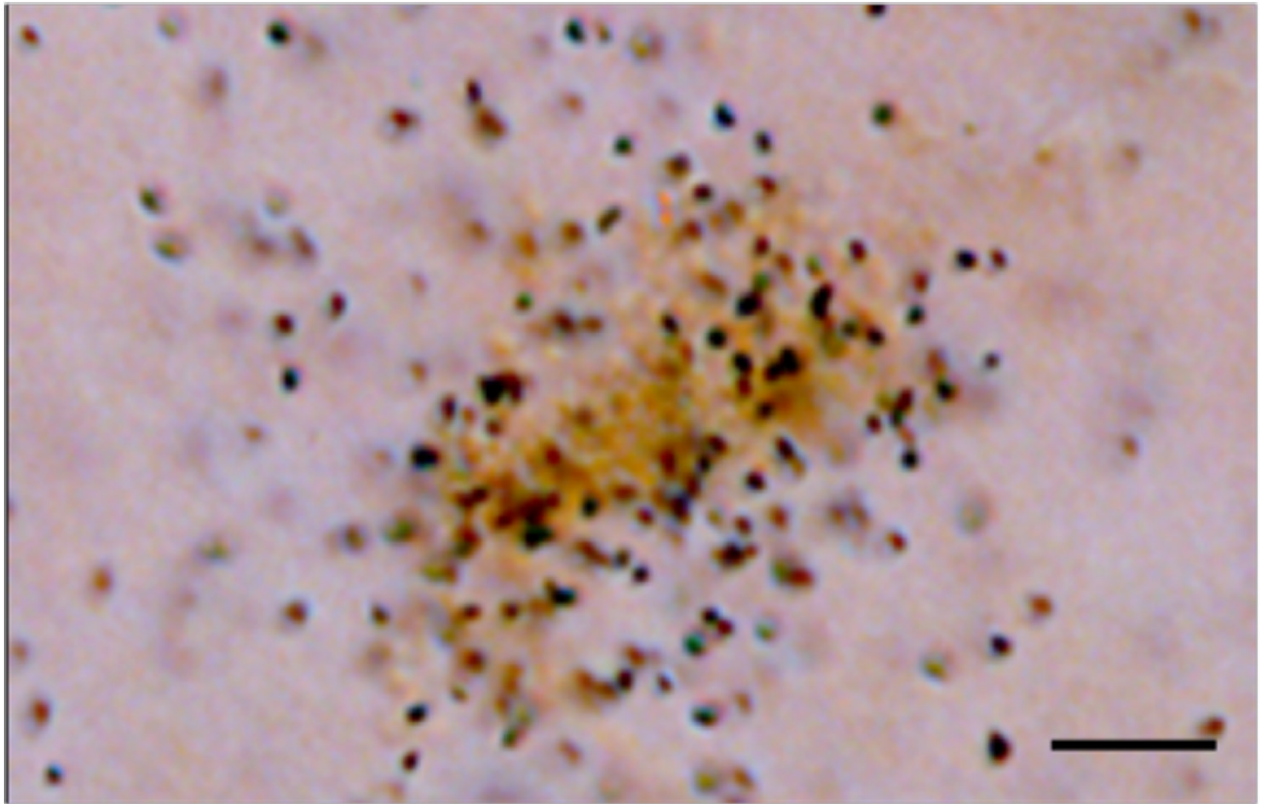
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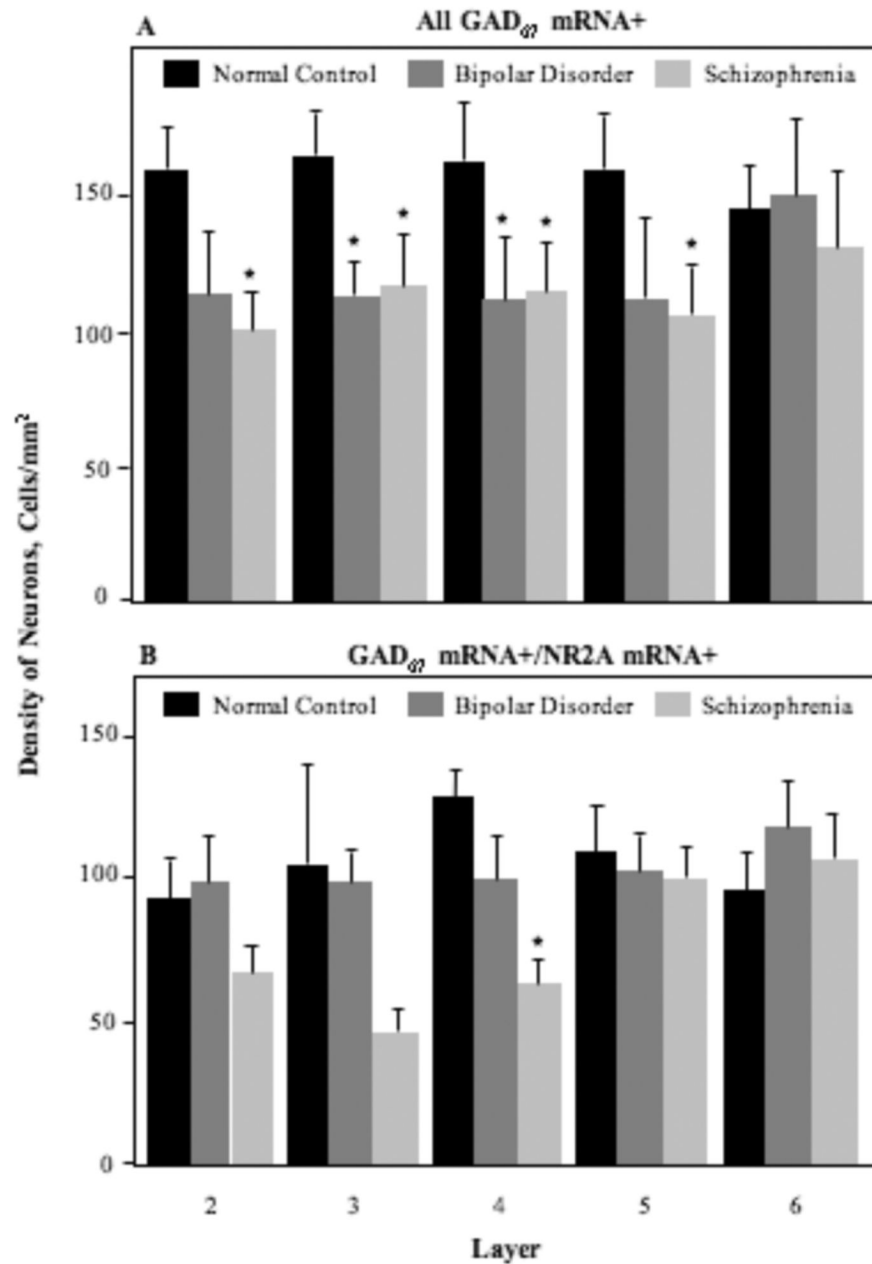
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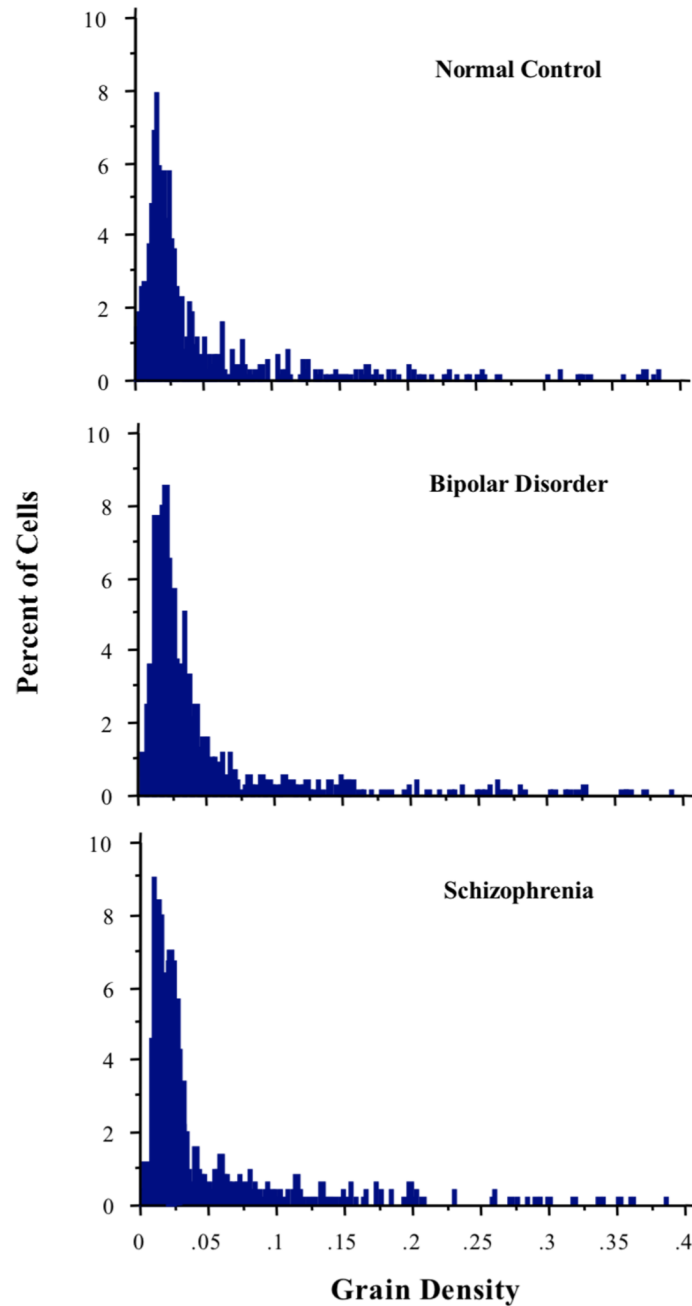


**Figure 1.**  
Photomicrograph of a double-labeled neuron with silver grains superimposed onto the digoxigenin reaction product. Scale bar=5  $\mu$ m.





**Figure 2.** (A) Mean density of all neurons that express GAD<sub>67</sub> mRNA. (B) Mean density of GAD<sub>67</sub> mRNA-expressing neurons that also express NR2A mRNA. Asterisks indicate statistically significant difference at p=0.01. Error bars represent SEM.



**Figure 3.** Grain density distribution is not significantly different between the 3 diagnosis groups, suggesting that the amount of NR2A mRNA being expressed by GABA interneurons with experimentally detectable transcript level is not altered in schizophrenia or bipolar disorder.

Table 1

Cases Used in the Present Study

Bipolar disorder										
Case	Age	PMI	Sex	Hemisphere	Race	CE	pH	Psychotropics received at time of death	Cause of death	
1	83	17.5	M	R	W	0	6.60	Divalproex	Cardiopulmonary failure	
2	74	22.9	F	R	W	100	U	Divalproex, lorazepam, olanzapine	Cardiopulmonary failure	
3	26	22.8	F	L	W	0	6.35	Lithium, lorazepam	Cardiopulmonary failure	
4	52	17.2	F	R	W	200	U	Olanzapine	Hepatic failure	
5	62	13.4	F	R	W	0	U	Divalproex, bupropion, sertraline	Breast cancer	
6	74	7.2	M	L	W	100	6.70	Gabapentin, olanzapine	Pneumonia	
7	85	27.5	F	L	W	534	6.28	Olanzapine, divalproex	Cardiopulmonary failure	
8	51	31	M	L	W	0	7.02	Clonazepam, gabapentin	Suicide by overdose	
9	69	29.5	M	L	W	0	U	Lithium	Pneumonia	
10	64	11	F	R	W	800	6.26	Trifluoperazine	Respiratory failure	
11	42	15.8	F	L	W	960	6.60	Lithium, divalproex, perphenazine	Medication overdose, rule out suicide	
12	29	10.7	F	L	W	200	6.70	Divalproex, lithium, clonazepam, olanzapine	Cardiac arrest	
13	36	9	M	L	W	U	U	Unknown	Suicide	
14	73	20.8	F	R	W	50	6.32	Carbamazepine, risperidone	Sepsis	
15	74	14.3	M	R	W	400	6.27	Divalproex, lithium, olanzapine, hydroxyzine prn, lorazepam prn, zolpidem prn	Pneumonia	
16	62	18.7	F	R	W	100	6.40	Divalproex, sertraline, risperidone, benzotropine, donepezil	Renal failure	
17	82	5.0	M	L	W	U	6.37	Unknown	Cardiopulmonary failure	
18	40	30.8	M	R	W	200	6.60	Gabapentin, ziprasidone, citalopram, risperidone, topiramate	Cardiac arrest	
19	38	22	M	R	W	200	6.24	Divalproex, paroxetine, olanzapine	Carbon monoxide poisoning	
20	47	16.3	F	R	W	50	U	Divalproex, topiramate, tiagabine, perphenazine, clonazepam,	Cardiopulmonary arrest	
Mean	58.1±18.7	18.2±7.7	M:F=10:10	R:L=11:9		212.3±282.8	6.48±0.23			
Schizophrenia										
21	85	15.7	F	R	W	150	U	Risperidone, lorazepam	Sepsis	
22	48	33.8	F	L	W	450	6.63	Risperidone, divalproex	Cardiac failure	

Bipolar disorder										
Case	Age	PMI	Sex	Hemisphere	Race	CE	pH	Psychotropics received at time of death	Cause of death	
23	44	19	M	L	W	266	6.20	Clozapine	Pneumonia	
24	89	13.5	F	L	W	20	U	Trifluoperazine	Pneumonia	
25	78	13.4	F	L	W	750	6.81	Haloperidol, lithium, cogentin	Sinus node disease	
26	61	19.9	M	R	W	300	6.68	Clozapine	Sepsis	
27	61	11	F	R	W	150	U	Paroxetine, clonazepam, clozapine	Myocardiac infarction	
28	84	25.8	F	R	W	0	6.14	None	Cardiac arrest	
29	26	16	M	R	W	357	6.75	Fluphenazine decanoate	Suicide by hanging	
30*	55	18	F	R	W	0	6.52	None	Oral cancer	
31*	47	19.2	M	R	W	0	6.57	Clonazepam, hydroxyzine prn	Lung cancer	
32	73	24	F	R	W	600	6.08	Risperidone, fluoxetine, chlorazepate, midazolam prn	Lung cancer	
33	49	19	M	L	W	500	6.60	Haloperidol decanoate, lorazepam	Suicide by hanging	
34	63	22.3	M	R	W	500	6.55	Clozapine, haloperidol, lorazepam, trazadone	Cardiac arrest	
45	72	21.7	F	R	W	400	6.65	Risperidone, paroxetine	Ovarian cancer	
36	66	22.1	M	R	W	1000	6.43	Haloperidol	Emphysema	
37	83	23.2	F	R	W	2000	6.91	Haloperidol decanoate, fluphenazine decanoate	Gastrointestinal bleed	
38	46	18.5	F	L	W	200	6.31	Olanzapine, divalproex	Sepsis	
39	42	27.1	M	R	W	0	6.64	None	Leukemia	
40	31	14	M	R	W	600	6.46	Risperidone, olanzapine, bupropion	Unknown (found dead in home)	
Mean	60.2±18.5	19.8±5.4	M:F=10:10	R:L=14:6		412.2±465.7	6.53±0.23			
Normal control										
41	49	24.6	M	L	W		6.76	None	Myocardiac infarction	
42	37	18.8	M	R	W		6.68	None	Electrocuton	
43	54	24.2	M	L	W		6.53	None	Cardiopulmonary arrest	
44	78	14.1	F	R	W		6.22	None	Myocardial infarction	
45	53	20.2	M	R	W		U	None	Cardiopulmonary arrest	
46	65	24.3	F	R	W		6.40	None	Lung cancer	
47	89	7.42	M	R	W		6.39	None	Cancer	

Bipolar disorder										
Case	Age	PMI	Sex	Hemisphere	Race	CE	pH	Psychotropics received at time of death	Cause of death	
48	69	15.3	M	R	W		6.88	None	Respiratory failure	
49	74	12.5	F	L	W		6.33	None	Cardiopulmonary arrest	
50	66	7.4	F	R	W		6.03	None	Cancer	
51	42	18.3	M	L	W		6.78	None	Myocardial infarction	
52	78	23.9	F	R	W		6.67	None	Breast cancer	
53	40	16.6	M	L	W		6.24	None	Myocardiac infarction	
54	67	22.3	M	L	W		6.42	None	Cardiopulmonary arrest	
55	70	22.5	F	L	W		6.26	None	Liver cancer	
56	66	18.7	M	R	W		6.76	None	Myocardial infarction	
57	79	20.9	M	L	W		6.74	None	Cancer	
58	38	28.8	M	L	W		6.53	None	Myocardiac infarction	
59	30	14.8	M	R	W		U	None	Blunt force trauma	
60	70	15	F	R	W		6.59	None	Cardiac arrest	
Mean	60.7±16.7	18.5±5.7	M:F=13:7	R:L=11:9			6.51±0.24			

CE: Chlorpromazine equivalent (in mg); PMI: Postmortem interval; U: Unavailable