The physiological roles of vesicular GABA transporter during embryonic development: a study using knockout mice

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Abstract

Background: The vesicular GABA transporter (VGAT) loads GABA and glycine from the neuronal cytoplasm into synaptic vesicles. To address functional importance of VGAT during embryonic development, we generated global VGAT knockout mice and analyzed them.

Results: VGAT knockouts at embryonic day (E) 18.5 exhibited substantial increases in overall GABA and glycine, but not glutamate, contents in the forebrain. Electrophysiological recordings from E17.5-18.5 spinal cord motoneurons demonstrated that VGAT knockouts presented no spontaneous inhibitory postsynaptic currents mediated by GABA and glycine. Histological examination of E18.5 knockout fetuses revealed reductions in the trapezius muscle, hepatic congestion and little alveolar spaces in the lung, indicating that the development of skeletal muscle, liver and lung in these mice was severely affected.

Conclusion: VGAT is fundamental for the GABA- and/or glycine-mediated transmission that supports embryonic development. VGAT knockout mice will be useful for further investigating the roles of VGAT in normal physiology and pathophysiologic processes.

Background

GABAergic and glycinergic neurotransmissions play critical roles in the central nervous system (CNS), because they regulate network activity and are essential for a number of brain functions, such as cognition, perception, movement and respiration. In the adult mammalian CNS, GABA and glycine are the main inhibitory neurotransmitters, but in fetal life and early postnatal development, both neurotransmitters act as either excitatory or inhibitory, depending on the intracellular chloride concentration.

GABA is synthesized from glutamic acid by glutamate decarboxylase (GAD) [1] and is accumulated into synaptic vesicles by the vesicular GABA transporter (VGAT) [2,3]. Two isozymes of GAD, GAD65 and GAD67, are primarily expressed in GABAergic neurons [4,5]. GAD65 knockout mice exhibit spontaneous seizures, elevated anxiety and altered sensitivity to pain [6,7]. GAD67 knockout mice die of cleft palate at birth [8]. VGAT is present in both GABAergic and glycinergic neurons and is also called the vesicular inhibitory amino acid transporter (VIAAT) [3,9]. In addition to its presence at GABAergic and glycinergic synapses, the role of VGAT/VIAAT in GABA and glycine release is supported by electrophysiological evidence from primary cultured hippocampal or spinal cord neurons of VGAT knockout mice [10] and VGAT-transfected secretory cells [11]. VGAT knockout mice die perinatally and show a hunched posture, cleft palate and omphalocele [10].

Divergent roles for the VGAT proteins are implicated in the nervous system. However, the contribution of
VGAT to tissues or cells outside of the CNS remains largely unclear. For example, neither muscle, lung nor liver phenotypes have been reported for these knockout mice. We independently generated VGAT knockout mice. To further investigate the roles of VGAT during development, we performed histopathological analyses in VGAT knockout muscle, lung and liver at an embryonic stage. These mice showed a reduction in the trapezius muscles, smaller saccules in the lung, and congestion in the liver. In addition, in VGAT knockout spinal cord motoneurons (MNs), spontaneous inhibitory postsynaptic currents (IPSCs) were absent. These experiments indicate that VGAT has an important role in the GABA- and/or glycine-mediated transmission that supports life. Preliminary results have been published in an abstract form [12].

Results

Generation of VGAT<sup>+/−</sup> mice

The targeting strategy used for the generation of VGAT knockout mice is shown in Figure 1A. Exons 2 and 3 encode the putative ten-transmembrane domain and C-terminus of the VGAT protein [3,13], and accordingly, the deletion of these regions was expected to destroy the function of the VGAT protein. Correctly targeted ES cell clones isolated were microinjected into blastocysts to generate chimeric mice. These mice were then crossed with C57BL/6 mice to generate heterozygous mice. We independently generated VGAT knockout mice by crossing VGAT<sup>flloxneo/+</sup> mice with CAG-Cre mice, in which Cre recombinase is expressed ubiquitously [14]. Genotyping was performed by Southern blot analysis (Figure 1B) and PCR (Figure 1C), and the DNA sequences around the loxP site in the knockout allele were also confirmed (data not shown). To obtain homozygous VGAT knockout (VGAT<sup>−/−</sup>) mice, we intercrossed the VGAT<sup>+/−</sup> mice. Western blot analysis revealed no VGAT protein expression in embryonic day (E) 18.5 VGAT<sup>−/−</sup> brain, whereas VGAT protein expression in VGAT<sup>+/−</sup> mouse brains was reduced to about half of the wild-type level (Figure 1D). All E18.5 VGAT<sup>−/−</sup> fetuses displayed cleft palate (Figure 1E) and omphalocele (Figure 1F), phenotypes that are consistent with those described by Wojcik et al. [10].

No VGAT<sup>+/−</sup> mice survived beyond birth (Table 1). When delivered by cesarean section on E18.5, both VGAT<sup>+/−</sup> (7 of 7) and VGAT<sup>−/−</sup> (11 of 12) fetuses began respiration, but none of the VGAT<sup>−/−</sup> fetuses (n = 7) began to breathe. Therefore, it is probable that VGAT<sup>−/−</sup> mice died at birth due to respiratory failure.

Elevations in GABA and glycine contents in VGAT<sup>−/−</sup> forebrains

In the absence of vesicular storage, neurotransmitter levels can be altered, and this alteration depends on the absence of the vesicular transporter. For example, monoamines are drastically reduced in vesicular monoamine transporter 2 knockout brains [15], but acetylcholine (ACh) is increased in vesicular acetylcholine transporter (VChT) knockout brains compared to control wild-type brains [16]. Therefore, we measured the amount of the neurotransmitters, GABA and glutamate in E18.5 VGAT<sup>−/−</sup> forebrain by HPLC. As shown in Figure 2A, VGAT<sup>−/−</sup> fetuses showed significant increases in both GABA and glycine, but not glutamate, compared to VGAT<sup>+/−</sup> fetuses. It is possible that the increase in GABA content in VGAT<sup>−/−</sup> fetuses was due to the elevated expression levels of GABA-synthesizing enzymes. To test for this possibility, we analyzed the expression levels of GAD65 and GAD67 in the embryonic brains. Our Western blot analysis showed that the expression levels of both GAD65 and GAD67 in VGAT<sup>−/−</sup> and VGAT<sup>+/−</sup> brains were similar (Figure 2B). These results indicate that the increase in GABA content was not derived from elevated amounts of GABA-synthesizing enzymes in VGAT<sup>−/−</sup> embryos.

Absence of functional inhibitory synaptic transmission in the VGAT<sup>−/−</sup> spinal cord

To examine the physiological nature of synaptic inputs to spinal MNs, we performed whole-cell patch-clamp recordings using isolated spinal cord preparations taken from VGAT<sup>−/−</sup> and control mouse embryos. In these preparations, the neuronal connections within the spinal cord are kept relatively intact [17]. In control lumbar MNs, spontaneous outward currents were observed when the membrane potential was depolarized at -40 mV above the chloride ion reversal potential (approximately -78 mV in the present experimental condition). These currents were blocked by bath application of the glycnergic antagonist strychnine and the GABAergic antagonist picrotoxin, indicating that the currents were IPSCs (n ≥ 8, Figure 3A). In contrast, we did not detect such spontaneous IPSCs in VGAT<sup>−/−</sup> MNs (n ≥ 12, Figure 3B). When the membrane potential was held at -70 mV, spontaneous inward currents were observed both in control and VGAT<sup>+/−</sup> MNs (Figure 3A and 3B). These inward currents were abolished by the concomitant bath application of the ionotropic
Figure 1 Generation of VGAT⁻/⁻ mice. (A) Schematic representation of the wild-type VGAT allele, the targeting vector, the VGAT-floxneo allele, and the VGAT knockout allele. Exons are represented by numbered white boxes. LoxP sites (open diamonds) and a PGK-Neo cassette (neo; gray box) flanked by the frt sites (open ellipses) were introduced into the wild-type VGAT locus by homologous recombination to produce the floxneo allele. The probes used for Southern blot analysis are indicated as white (5' probe) and red (3' probe) boxes. The expected sizes of the KpnI- and Ncol-digested genomic DNA fragments hybridized with the 5' and 3' probes, respectively, are indicated as lines under the schemes. Relevant restriction sites are indicated as follows: K, KpnI; N, Ncol. PCR primers are indicated as arrows. (B) (Left) Southern blot analysis of KpnI-digested genomic DNA isolated from VGAT+/+(+/+) and VGATfloxneo/+ (Floxneo/+) mice using the 5' probe indicated in A. The wild-type allele corresponds to the 15.2 kb band, whereas the floxneo allele corresponds to the 12.0 kb band. (Right) Southern blot analysis of Ncol-digested genomic DNA isolated from VGAT+/+(+/+), VGAT+/- (+/-), VGAT⁻/⁻ (-/-), and VGATfloxneo/+ (Floxneo/+) mice using the 3' probe indicated in A. The wild-type allele, the knockout allele, and the floxneo allele correspond to the 10.7 kb, 7.4 kb, and 6.3 kb bands, respectively. (C) Genotyping of offspring from intercrosses of VGAT+/- mice by PCR. Three primers were used (see Methods). Primers P2 and P3 produce a 183 bp fragment that represents the wild-type allele, whereas primers P1 and P3 produce a 430 bp fragment that represents the knockout allele. (D) Western blot analysis of E18.5 whole brain homogenates from VGAT⁺/⁺ (+/+), VGAT⁻/⁺ (+/-), and VGAT⁻/⁻ (-/-) using the anti-VGAT antibody directed against an N-terminal epitope. (E) Ventral views of the upper jaw of E18.5 VGAT⁺/⁺ (+/+), and VGAT⁻/⁻ (-/-) mice. In contrast to the completely fused palate of a VGAT⁺/⁺ mouse, secondary palatal shelves of a VGAT⁻/⁻ mouse did not contact each other (arrows), and its nasal cavity (asterisk) could be seen. (F) Lateral views of E18.5 VGAT⁺/⁺ (+/+), and VGAT⁻/⁻ (-/-) mice. An arrow indicates omphalocele in a VGAT⁻/⁻ mouse. In addition, the VGAT⁻/⁻ mouse showed an extremely hunched position in contrast to the VGAT⁺/⁺ mouse.
glutamate receptor blockers, a non-NMDA receptor antagonist 6-cyano-7-nitroquinoxaline-2,3-dione (CNQX) and an NMDA receptor antagonist D-2-amino-5-phosphonovaleric acid (AP5), indicating that MNs received excitatory synaptic transmission in the VGAT-/- spinal cord.

Alterations in body weight, response to stimuli, trapezius muscle, liver and lung of VGAT-/- mice

Wojcik et al. [10] reported that VGAT-/- mice display the phenotypes such as cleft palate, omphalocele, hunched posture, immobility and stiffness. However, we proposed that there would be some other alterations in VGAT-/- mice because VGAT is an essential molecule for GABAergic and glycineric transmission. To address the question whether VGAT is essential for fetal growth, we initially measured the body weight of VGAT-/- fetuses compared to VGAT+/+ and VGAT+/- littersmates. The body weight of the E18.5 VGAT-/- fetuses was significantly lower than that of VGAT+/+ and VGAT+/- fetuses (VGAT+/+: 1.18 ± 0.11 grams, n = 17; VGAT+/-: 1.20 ± 0.08 grams, n = 45; VGAT-/-: 1.05 ± 0.11 grams, n = 32 [mean ± SD], P < 0.001, one-way ANOVA, post hoc Fish-

Table 1 Genotypes of offspring from intercrosses of VGAT +/- mice and phenotypes of VGAT-/- mice

| Age       | Genotype | Phenotype                  | No. of +/- found dead | No. of +/- with omphalocele | No. of +/- with cleft palate |
|-----------|----------|----------------------------|-----------------------|-----------------------------|-----------------------------|
| E18.5     | +/-      |                            | 69 (24.5%)            | 2/77*                       | 77/77*                      |
|           | +/-      |                            | 136 (48.2%)           | 77 (27.3%)                  | 29/29*                      |
| Newborn   | +/-      |                            | 22 (19.6%)            | 14/14*                      | Not determined              |
|           | +/-      |                            | 76 (67.9%)            | 14 (12.5%)                  | Not determined              |

*aaffected/examined.

Comparison of cleft palate and omphalocele between VGAT-/- mice and GAD67-/- mice

Cleft palate is exhibited by VGAT-/- and GAD67-/- mice [8,10,23], demonstrating that GABAergic transmissions are involved in palatogenesis. Because VGAT and
Figure 2 Neurotransmitter contents and expression levels of GAD65 and GAD67. (A) Neurotransmitter contents of E18.5 mouse forebrain. VGAT<sup>−/−</sup> mice showed significantly higher levels of GABA and glycine than VGAT<sup>+/+</sup> mice, but not glutamate. Values represent means ± SD (*P < 0.05; Student’s t-test, n = 5-13 per group). (B) Western blot analysis. The expression level of GAD65 and GAD67 in whole brain homogenates was not significantly different between VGAT<sup>+/+</sup> (+/+) and VGAT<sup>−/−</sup> (-/-) mice. Equal amounts of protein were loaded and probed with an antibody that recognizes both GAD65 and GAD67. For the statistical comparison, the same blot was probed with anti-β-actin antibody as an internal control and measurements for GAD65 and GAD67 bands were normalized to the β-actin bands (GAD65: VGAT<sup>+/+</sup> 100 ± 23%; VGAT<sup>−/−</sup> 111 ± 20%, n = 4, P = 0.71) (GAD67: VGAT<sup>+/+</sup> 100 ± 18%; VGAT<sup>−/−</sup> 93 ± 17%, n = 4, P = 0.80).
GAD67 exhibit different molecular functions, we investigated whether the severity of cleft palate was different between VGAT-/- and GAD67-/- mice. Figure 5A shows hematoxylin-eosin staining of coronal sections from the oral region. In the cleft palate of VGAT-/- mice, the palatal shelves remained vertical along the sides of the tongue (3 of 3). However, in GAD67-/- mice, the palatal shelves were elevated up to a horizontal position (3 of 3). In one of the GAD67-/- mice, the palatal shelves even fused with each other completely (data not shown). These observations suggest that palatogenesis progresses further in GAD67-/- mice than in VGAT-/- mice. Our observations also suggest that cleft palate in VGAT-/- mice is more severe than in GAD67-/- mice.

The observation of omphalocele in VGAT-/- mice prompted us to investigate whether GAD67-/- mice displayed omphalocele. We found omphalocele in GAD67-/- and VGAT-/- mice (Figure 5B), indicating that GABA signaling is involved in its onset. The incidence rate of omphalocele in GAD67-/- mice was 43% (17 of 40), whereas the incidence in VGAT-/- mice was 100% (77 of 77; see also Table 1). Thus, the penetrance of omphalocele in GAD67-/- mice was lower than in VGAT-/- mice. The size of omphalocele appeared to be larger in VGAT-/- mice than in GAD67-/- mice. Taken together, these data suggest omphalocele in GAD67-/- mice is less severe than in VGAT-/- mice, similar to what was observed with the cleft palate.

**Discussion**

The present work addresses the contribution of VGAT to embryonic development. We generated VGAT mice and found that VGAT is fundamental for GABA and/or glycine release in the spinal cord. Moreover, in the absence of VGAT, there are profound effects on muscle, liver and lung during embryonic development. These observations implicate VGAT in the functional roles of its expression from the cellular to the whole-body level.

**Generation of VGAT knockout mice**

Wojcik et al. [10] generated VGAT knockout mice, in which a mutation was inserted into exon 1, and these mice exhibit cleft palate and omphalocele. Here, we generated floxed VGAT knockout mice, in which exons 2 and 3 of the VGAT gene were flanked by loxP sites. Crossing the floxed VGAT mice to CAG-Cre mice reproduced the phenotypes of cleft palate and omphalocele. Exons 1 and 2/3 encode the cytoplasmic domain and the transmembrane domain, respectively [3,13]. Our results demonstrate that exons 2 and 3 are dispensable for the function of VGAT.

VGAT-floxneo/floxneo mice were born at the expected frequency, were viable, did not have a cleft palate or omphalocele, and were overtly indistinguishable from their wild-type littermates. Western blot showed that the level of VGAT protein expression in VGAT-floxneo/floxneo brain was not different from the wild-type brain (Additional file 1: Supplementary Figure S1). These results suggest that a loxP sequence and an frt-flanked phosphoglycerate kinase promoter-driven neomycin-resistance gene (PGK-Neo) inserted into intron 1 and the 3'-flanking region of the VGAT gene, respectively, do not affect the expression level of VGAT protein. Therefore, the VGAT-floxneo
Figure 4  Histological analysis of VGAT<sup>−/−</sup> mice. Histological analysis (hematoxylin and eosin staining) of trapezius muscle (A, B), sagittal sections (C, D), liver (E, F) and lung (G, H) from E18.5 VGAT<sup>−/−</sup> mice (B, D, F, H) and control mice (A, C, E, G). (A, B) The trapezius muscle (bounded partly by white dashed lines) of VGAT<sup>−/−</sup> mouse (B) was thinner than the control mouse (A). Scale bar: 200 μm. (C, D) The VGAT<sup>−/−</sup> ribs (arrow in D) in the lower part were depressed, and their position was inside compared to the control ribs (arrow in C). H, heart; L, liver. (E, F) Red blood cell congestion was characteristic of VGAT<sup>−/−</sup> liver, but not control liver. Scale bar: 200 μm. (G, H) The VGAT<sup>−/−</sup> lung contained much less alveolar space than the control lung. Scale bar: 500 μm.
allele allowed for Cre-mediated conditional inactivation of the VGAT gene, and mice carrying these alleles will be useful in examining VGAT function at different developmental stages and in distinct cell types.

Increase of overall GABA and glycine contents in VGAT−/− forebrains

We showed that both GABA and glycine contents were increased in the VGAT−/− forebrain, but not an excitatory neurotransmitter glutamate, which is transported into synaptic vesicles by vesicular glutamate transporters. An increase in GABA content in VGAT−/− mice is similar to that in ACh contents in VACHT knockout mice [16], but it is opposite of the decrease in monoamine contents in vesicular monoamine transporter 2 knockout mice. In C. elegans, the mutational inactivation of VGAT also leads to an increase in GABA immunoreactivity in GABAergic neurons [24]. In VACHT knockout mice the amount of the ACh-synthesizing enzyme choline acetyltransferase (ChAT) is increased at the mRNA and protein levels compared to their wildtype littermates, suggesting that the change in ChAT expression may be related to a compensatory mechanism due to the lack of ACh release [16]. Conversely, the amounts of the GABA-synthesizing enzymes GAD65 and GAD67 were not different in between the brains of VGAT−/− mice and their control littermates (Figure 2B). Therefore, it is possible that the increased GABA in VGAT−/− brain was due to a reduction in their degradation. GABA and glycine are released from presynaptic
neurons into the synaptic cleft and are retrieved in neu-
rons and glial cells by plasma membrane transporters
[25,26]. GABA and glycine taken up in glial cells are
further metabolized, but the GABA and glycine taken
up in neurons are directly recycled into synaptic vesicles
[27,28]. Because degradation systems for both GABA
and glycine are mainly localized to glial cells [27,29],
the transport into glial cells from the synaptic cleft is im-
portant for their degradation. Because the synaptic release
of GABA and glycine was absent in VGAT−/− mice, the
deletion of VGAT may result in little or no transport of
GABA and glycine into glial cells. GABA and glycine
then accumulate in the GABAergic and glycinergic neu-
rons, respectively, but they are not degraded in the glial
cells of VGAT−/− mice.

**Contribution of VGAT to motor function**
In the embryonic spinal cord of rodents, synaptic trans-
mission to MNs mediated by GABA and glycine is pro-
minent from the early fetal period [30,31]. Our results
from the electrophysiological recordings of spinal cord
MNs indicate that the inhibitory synaptic transmission
was clearly absent in the VGAT knockout MNs, but
that the excitatory synaptic transmission was present.
Our results also suggest the absence of other functional
mechanisms that transport GABA and/or glycine into
synaptic vesicles in these synapses. VGAT−/− fetuses at
E17.5-18.5 not only were completely immobile and stiff,
but also none of them responded to mechanical stimuli
by pinching the limb or the tail. Therefore, it is probable
that the lack of inhibitory transmission onto MNs in
VGAT−/− fetuses resulted in defects in the spontaneous
and stimulus-induced movements in vivo despite the
presence of excitatory synaptic transmission.

In addition to the defect in motor movement, trape-
zius muscle displayed atrophy in VGAT−/− mice. Embryonic
myogenesis progresses by the proliferation of
myoblasts and fusion of myotubes, but it requires sub-
stantial cell death [32]. Physical forces play a significant
role in the development and maintenance of skeletal
muscle [33]. In cultured myoblasts, chronic and cyclic
stretch results in an increase in cell death, including
apoptosis [34]. Therefore, a possible explanation for the
atrophy in VGAT−/− trapezius muscle is that stretching
of the trapezius muscle due to the hunched posture
caused an increase in apoptosis during development.

**Phenotypes of VGAT and GAD67 knockouts outside
of the brain**
Ventral body wall closure abnormalities, such as ompha-
locele, are common human birth defects, but their
molecular and cellular bases are poorly understood [35].
The mouse provides a model system to study the
genetic defects and environmental insults that can lead
to ventral body wall closure abnormalities [20]. In this
study, omphalocele was observed in VGAT−/− and
GAD67−/− mice, indicating that the lack of GABA signal-
ing was involved in its onset. Omphalocele has been
observed in K+-Cl--cotransporter 2 (KCC2) knockout
mice [36], and KCC2 is required for GABA- and/or gly-
cine-induced hyperpolarizing responses [37]. In the
KCC2 knockout mice, GABA and/or glycine signals
continue to act in an excitatory, but not an inhibitory,
manner. Therefore, the omphaloceles observed in both
VGAT−/− mice and GAD67−/− mice resulted from defects
in the inhibitory neurotransmission derived from the
hyperpolarizing response. A lack of inhibitory transmis-
sion in VGAT−/− mice may lead to motor deficits, such
as a hunched posture. It is likely that the hunched pos-
ture resulted in increases in both intrathoracic and
intraabdominal pressures and this increased pressure
caused omphalocele.

Concerning the mechanism of onset of cleft palate,
studies using knockout mice have revealed associations
between cleft palate and mutation of genes related to
GABA signaling, such as GAD67 and GABRB3 [8,23,38].
Because the lack of the GAD67 gene leads to a reduc-
tion in tongue movement [39], the sluggish tongue may
be an obstacle to development of the palatal shelves.

The cleft palate and omphalocele phenotypes were
more severe in VGAT−/− mice than in GAD67−/− mice.
Glycinergic transmission is present in embryonic spinal
cord and brainstem [40]. Hyperekplexia is a neuroge-
etic disorder caused mostly by mutations in the gene
encoding the α1 subunit of glycine receptor and is char-
acterized by an exaggerated startle response and neo-
tal hypertonia. In patients with hyperekplexia, the
recurrent abdominal muscle contraction from the exag-
ggerated startle response can increase the abdominal
pressure and lead to omphalocele and inguinal hernia
[41,42]. These reports suggest that a defect in glyciner-
gic transmission is involved in the onset of omphalocele.
A small amount of GABA is synthesized by another
GAD isoform, GAD65, at the embryonic stage [8,43].
The differences in the severity between VGAT−/− and
GAD67−/− mice must be due to the presence of both gly-
cinergic and GAD65-produced GABAergic transmission
in GAD67−/− fetuses, but not in VGAT−/− fetuses.

**Conclusion**
In the present study, we established a VGAT knockout
mouse, with which we demonstrated that VGAT is fun-
damental for GABAergic and/or glycinergic transmis-

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transmission in mice. GABAergic neurons are classified into several subtypes according to the expression of chemical markers, such as parvalbumin and somatostatin [44,45]. Therefore, our floxed VGAT mice will be useful for conditional knockout studies to further investigate the role of VGAT in GABAergic neuronal subtypes.

Methods

Animals

All animal procedures were conducted in accordance with the guiding principles of the NIH under the review and approval of the Animal Care and Experimentation Committee, Gunma University, Showa Campus (Maebashi, Japan). Every effort was made to minimize the number of animals used and their suffering.

Construction of the Targeting Vector

Genomic BAC clones containing the mouse VGAT (mVGAT) locus were purchased, and the regions covering the entire VGAT gene were subcloned [13]. A genomic fragment spanning exons 1-3 of the mVGAT gene was used for the targeting vector (Figure 1; targeting vector). The HindIII (in the 5′-flanking region) - KpnI (in the 3′-flanking region) fragment (7.5 kb) was subcloned into pBluescript II KS(-), and the 5′-loxp site was introduced into the XbaI site (in intron 1). The 5′-loxP site was flanked by a KpnI site artificially introduced for Southern blot analysis. The 7.5 kb fragment was used as the 5′ homologous region containing the 5′-flanking region, exons 1-3 and the 3′-flanking region. The rfr-flanked PGK-Neo cassette for positive selection of ES clones and the 3′-loxP site were inserted into the KpnI site (in the 3′-flanking region). The KpnI-BstEII fragment in the 3′-flanking region (3.5 kb) was added as the 3′ homologous region. An MC1-DT-ApA cassette for negative selection [46] was ligated to the 3′ end of the homologous region.

Creation of a VGAT knockout allele

The linearized targeting vector was introduced by electroporation into ES cells (CCE) of 129/SvEv mouse origin, and G418-resistant colonies were screened by Southern blot analysis using probes outside of the target site. The KpnI-BstEII fragment (7.5 kb) was used for the targeting vector (Figure 1; targeting vector). KpnI-digested genomic DNA prepared from ES cell colonies was hybridized with 5′-loxP probes. The 7.5 kb fragment was used as the 5′ homologous region containing the 5′-flanking region, exons 1-3 of the VGAT gene and primers P2 and P3 amplified a 183-bp fragment specific for the 5′-flanking region of the VGAT gene. The primer set of P2 and P3 amplified a 183-bp fragment specific for the wild-type allele, and the primer set of P1 and P3 yielded a 430-bp fragment specific for the knockout allele.

Genotypes of VGAT+/+, VGAT+/- and VGAT-/- mice were determined by PCR using the following oligonucleotides: primer P1 (5′-AGTCGATCCCCTGGCATTCCTCAACTC-3′) corresponding to intron 1 of the VGAT gene and primers P2 (5′-TCAGAGGCTTCTCTCTAGGGCTGCTG-3′) and P3 (5′-GACCTCCCCCATTTGATAGATGGC-3′) corresponding to the 3′-flanking region of the VGAT gene. The primer set of P2 and P3 amplified a 183-bp fragment specific for the wild-type allele, and the primer set of P1 and P3 yielded a 430-bp fragment specific for the knockout allele.

GAD67 knockout mice

We used homozygous GAD67-GFP (△neo) (GAD67-GFP/GFP) mice as GAD67 knockout (GAD67-/-) mice. The generation of the GAD67-GFP (△neo) mice and their genotyping by PCR were described previously [47,48]. In the GAD67-GFP (△neo) mice, a cDNA encoding enhanced green fluorescent protein (EGFP) followed by an SV40 polyadenylation signal was targeted to the locus encoding GAD67 by homologous recombination, and the GAD67 gene was disrupted.

Western blotting and measurement of neurotransmitter contents

For Western blotting, homogenates prepared from E18.5 mouse brain were separated by 7.5% SDS-polyacrylamide gel electrophoresis, transferred to nitrocellulose membrane (Whatman, Maidstone, UK), and probed with antibodies specific for VGAT (1:1000) [49], GAD65/67 (1:1000) [50], synaptophysin (1:1000) (Synaptic Systems, Gottingen, Germany), and β-actin (1:10000) (Abcam, Cambridge, UK). After the membranes were washed with Tris-HCl buffered saline containing 0.05% (w/v) Tween 20, the bound antibodies were visualized with horseradish peroxidase-conjugated goat anti-mouse IgG or anti-rabbit IgG (Jackson ImmunoResearch Laboratories, West Grove, PA) using the ECL Western blotting detection system (GE Healthcare, London, UK). Protein levels were quantified using Light Capture and its quantification software (ATTO, Tokyo, Japan). Expression levels were normalized to β-actin or synaptophysin levels, and the values are expressed as means ± SE. Statistical significance was assessed using Student’s t-test.

GABA, glycine, and glutamate contents in the E18.5 mouse forebrain were measured according to previously described method [21,47].

Electrophysiological recording in spinal cord

Embryos (E17.5-18.5) of control (VGAT+/+; n = 3, VGAT+/-; n = 5) and VGAT-/- (n = 12) mice were examined using whole cell patch-clamp recordings.
decapitated and eviscerated, and the spinal cord was removed by ventral laminectomy. The isolated spinal cord was placed in a recording chamber perfused with oxygenated Ringer’s solution (118.4 mM NaCl, 3 mM KCl, 2.52 mM CaCl₂, 1.25 mM MgSO₄, 25 mM NaHCO₃, 1.18 mM KH₂PO₄, and 11.1 mM D-glucose aerated with 5% CO₂ in O₂) at room temperature. Recordings from MNs in the isolated spinal cord were performed as described previously [17]. Briefly, visually guided whole-cell tight-seal recording of MNs was performed with patch electrodes pulled from thick walled borosilicate glass to a final resistance of 5-8 MΩ. The electrode tips were filled with (in mM) 138 K-glucuronate, 10 HEPES, 1 CaCl₂, 5 ATP-Mg, and 0.3 GTP-Li. Intracellular signals were amplified with a Multiclamp 700B amplifier (Molecular Devices, Union City, CA), digitized at 5 kHz with the Digidata 1440A data acquisition system (Molecular Devices) and saved on a hard disk for off-line analysis. Electrical stimulations of lumbar ventral roots (VRs) were performed using glass suction electrodes. MNs were identified visually as cells with large soma in the ventral horn and by observing the antidromic firing activated by the electrical stimulation of the adjacent VR. All drugs (CNQX, AP5, strychnine and PTX) were bath-applied to the preparation. Analysis was performed using pClamp 10 software (Molecular Devices).

Histology
VGAT⁻/⁻, VGAT⁺⁻/⁻ and VGAT⁺⁺/+ mice at E18.5 were investigated. Samples were fixed in 10% (vol/vol) formaldehyde, dehydrated with a graded series of ethanol solutions and embedded in paraffin. Three-micrometer sections were prepared, subjected to paraffin removal by immersion in xylene, rehydrated, and stained with hematoxylin-eosin. VGAT⁺⁻/⁻ and VGAT⁺⁺/+ mice were mixed together and are referred to as control mice.

Additional material

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Authors’ contributions
Conceived and designed the experiments: KS, TK, HN, VN, MY, SK, AF, MF, KN, KO, YY. Performed the experiments: KS, TK, HN, TF, RH, ST, SE, MU, KI, MF, KN, KO, YY. Analyzed the data: KS, TK, HN, TF, RH, X, AF, MF, KN, KO, YY. Contributed new reagents/analytical tools: ST, MM, JM. Wrote the paper: KS, TK, HN, AF, MF, KN, KO, YY. All authors read and approved the final manuscript.

Competing interests
The authors declare that they have no competing interests.

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Additional file 1: Supplementary Figure S1. VGAT expression levels in VGAT mutant mice. (A, B) VGAT expression level was normal in adult VGATfloxed/floxed mice. Western blot of whole brain homogenates from adult VGAT⁻/⁻ (+/+), VGATfloxed/floxed (floxed/floxed) and VGAT⁺/+ (floxed/floxed) mice is shown (A). VGAT expression level normalized to β-actin was not different between VGAT⁺⁻/+ (+/+), VGATfloxed/floxed (floxed/floxed) and VGAT⁺/+ (floxed/floxed) mice (B) (C, D) VGAT expression level was reduced by about half in adult VGAT⁻/⁻ mice. Western blot of whole brain homogenates of adult VGAT⁻/⁻ (+/+), VGATfloxed/floxed (floxed/floxed) and VGAT⁺/+ (floxed/floxed) mice is shown (C). VGAT expression level normalized to synaptophysin was significantly different between VGAT⁺⁻/+ (+/+), VGATfloxed/floxed (floxed/floxed) and VGAT⁺/+ (floxed/floxed) mice (D). Significance was tested by Student's t-test (P < 0.05).
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