In Vivo Preclinical Molecular Imaging of Repeated Exposure to an N-methyl-D-aspartate Antagonist and a Glutaminase Inhibitor as Potential Glutamatergic Modulators S

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ABSTRACT

And Experimental Therapeutics

Glutamate is the principal excitatory neurotransmitter in the brain and is at the base of a wide variety of neuropathologies, including epilepsy, autism, Fragile X, and obsessive compulsive disorder. Glutamate has also become the target for novel drugs in treatment and in fundamental research settings. However, much remains unknown on the working mechanisms of these drugs and the effects of chronic administration on the glutamatergic system. This study investigated the chronic effects of two glutamate-modulating drugs with imaging techniques to further clarify their working mechanisms for future research opportunities. Animals were exposed to saline ml/kg), (5S,10R)-(+)-5-Methyl-10,11-dihydro-5H-dibenzo[a,d]cyclohepten-5,10-imine (MK-801) (0.3 mg/kg), or ebselen (10 mg/kg) for 7 consecutive days. At the sixth injection, animals underwent a positron emission tomography (PET)/ computed tomography (CT) with (3-(6-methyl-pyridin-2ylethynyl)-cyclohex-2-enone-O-11C-methyl-oxime) (ABP-688) to visualize the metabotropic G protein-coupled glutamate

receptor 5 (mGluR5). After the seventh injection, animals underwent a magnetic resonance spectroscopy (MRS) scan to visualize glutamate and glutamine content. Afterward, results were verified by mGluR5 immunohistochemistry (IHC). PET/CT analysis revealed that animals receiving chronic MK-801 or ebselen had a significant (P < 0.05) higher binding potential $(2.90 \pm 0.47 \text{ and } 2.87 \pm 0.46, \text{ respectively})$ when compared with saline (1.97 \pm 0.39) in the caudate putamen. This was confirmed by mGluR5 IHC, with 60.83% \pm 6.30% of the area being highlighted for ebselen and 57.14% \pm 9.23% for MK-801 versus $50.21\% \pm 5.71\%$ for the saline group. MRS displayed significant changes on the glutamine level when comparing chronic ebselen (2.20 \pm 0.40 μ mol/g) to control $(2.72 \pm 0.34 \, \mu \text{mol/g})$. Therefore, although no direct effects on glutamate were visualized, the changes in glutamine suggest changes in the total glutamate-glutamine pool. This highlights the potential of both drugs to modulate glutamatergic pathologies.

[N-methyl-D-aspartate (NMDA), α-amino-3-hydroxy-5-methyl-

4-isoxazolepropionic acid, and kainite]. There are also three

Introduction

Glutamate is the principal excitatory neurotransmitter in the brain. It is transmitted via three families of ionotropic receptors, which have intrinsic cation permeable channels

groups of metabotropic G protein-coupled glutamate receptors (mGluRs) that act on membrane ion channels and second messengers such as cAMP and diacylglycerol to modify neuronal and glial excitability. Additionally, there are three neuronal glutamate transporters in the brain and two glial glutamate transporters: the glial glutamate and aspartate transporter and the glial glutamate transporter (Meldrum, 2000). It has been shown that overstimulation of the glutamatergic

neurotransmission is neurotoxic (Porciúncula et al., 2001), and glutamate abnormalities have been well characterized in a number of neuropathologies, such as Fragile X, autism, schizophrenia, and epilepsy. Also, in other disorders, such as depression, anxiety, and more recently obsessive compulsive disorder (OCD), the emerging role of glutamate signaling has been garnering interest (Coyle, 2006; Pittenger et al., 2011;

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ABBREVIATIONS: ABP-688, (3-(6-methyl-pyridin-2-ylethynyl)-cyclohex-2-enone-O-11C-methyl-oxime); BPnd, binding potential; CP, caudate putamen; Cr, creatine; CRLB, Cramér-Rao lower bounds; CT, computed tomography; Glu, glutamate; GPC, glycerophosphorylcholine; IHC, immunohistochemistry; mGluR, metabotropic G protein-coupled glutamate receptor; MK-801, (5S,10R)-(+)-5-Methyl-10,11-dihydro-5H-dibenzo [a,d]cyclohepten-5,10-imine; MR, magnetic resonance; MRS, magnetic resonance spectroscopy; NMDA, N-methyl-p-aspartate; OCD, obsessive compulsive disorder; PCr, phosphocreatine; PET, positron emission tomography; QP, quinpirole; VOI, volume of interest.

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Miladinovic et al., 2015). In addition, our group has previously shown that there is a role for glutamate in the quinpirole (QP) rodent model of OCD. In this model, animals are exposed to injections of QP and an open field test on a biweekly basis for a total duration of 5 weeks. These animals showed excessive compulsive behavior and were characterized by an increased distribution of mGluR5, as visualized by positron emission tomography (PET) (Servaes et al., 2017). Another group showed that in Sapap3 knockout mice, another preclinical model for OCD (Welch et al., 2007), mGluR5 signaling was elevated and associated with constitutively active postsynaptic receptors (Ade et al., 2016). As it is widely accepted that OCD results from a dysfunctional cortico-striato-thalamic circuitry (Fig. 1) (Denys et al., 2010), it is possible that, because the brain is exposed to higher levels of glutamate in OCD patients (Chakrabarty et al., 2005), the normal feedback loop is disrupted. Whether this is a consequence or a cause of the symptoms remains to be elucidated. Also, in other glutamate-related pathologies, such as Fragile X syndrome, autism, and epilepsy, there appears to be a specific role for the mGluR5 receptor (Pop et al., 2014; Chana et al., 2015; Kelly et al., 2018).

Taken together, glutamate has become a target for novel drugs. Therefore, drugs that can interact with the glutamatergic transmission, directly affecting the glutamate levels or the associated receptors in a positive or negative direction, could be of great assistance to further understand these pathologies. By modifying the glutamatergic system in either direction in a preclinical setting, more insight could be acquired in the manifestation of the pathologies mentioned before, either improving or aggravating the symptoms.

A variety of drugs interact with the glutamatergic system. Ebselen [2-phenyl-1,2-benzisoselenazol-3(2H)-one] is a seleno-organic compound of low toxicity that is characterized by a unique pharmacological profile (Schewe, 1995). Ebselen interacts directly with the glutamatergic system in rats (Nogueira et al., 2002). Although the exact mechanism of action has not been described yet (Azad and Tomar, 2014), preliminary results indicate that ebselen protects neurons from ischemic damage (Seo et al., 2009). Furthermore, in vitro studies have shown that ebselen inhibited [³H]glutamate release and uptake (Porciúncula et al., 2001). Importantly, ebselen is also an effective glutaminase inhibitor (Thomas et al., 2013). With glutaminase catalyzing the hydrolysis of glutamine to glutamate, this is a further indication of the direct effects of ebselen on the glutamatergic signal.

Another major group of glutamate-affecting drugs target the NMDA ionotropic glutamate receptor. NMDA receptors play a role in a variety of glutamate-associated pathologies, such as OCD (Arnold et al., 2004; Albelda et al., 2010), epilepsy (Ghasemi and Schachter, 2011), and autism (Lee et al., 2015). By directly modulating the NMDA receptors with NMDA receptor antagonists such as ketamine and (+)-MK-801 (dizocilpine maleate), [(5S,10R)-(+)-5-Methyl-10,11-dihydro-5H-dibenzo[a,d]cyclohepten-5,10-imine], the interneuron GABAergic receptors can be inhibited, which typically results in strong psychostimulatory and psychotomimetic effects. Furthermore, this modification could prove to be useful for investigating the NMDA abnormalities on the level of the receptor in the diseases mentioned earlier.

Ebselen's glutamate-reducing properties, together with its glutaminase inhibition and protection of ischemic damage, highlight its application potential for OCD. Additionally,

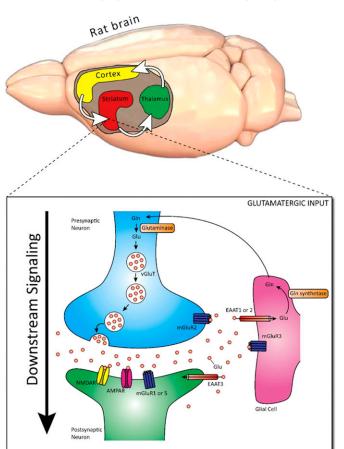


Fig. 1. The frontal cortex, responsible for functions such as error detection, working memory, and goal-directed behavior, sends signals through the striatum through glutamatergic afferents. From the striatum, this signal either inhibits or reduces the inhibition to the thalamus, depending on the levels of Dopamine present. From the thalamus, a signal is sent back to the frontal cortex, completing a feedback loop. It has been hypothesized that OCD results from a dysfunction in this circuit, in particular in the downstream signaling pathway. As the brain is exposed to higher levels of glutamate, this will likely cause overstimulation and disruption of the circuits and receptors in place. AMPAR, AMPA (α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid) receptor; NMDAR, NMDA receptor; VGluT, vesicular glutamate transporter.

although MK-801 will unlikely end up being a treatment option, the possibility for it to modulate glutamate levels could be of particular importance to further elucidate the underlying glutamatergic role in QP-induced OCD rats. Previously, MK-801 has been used in animal models for OCD to document the effect of activation of NMDA receptors on compulsions. Shannon and Love (2004) reported that MK-801 increased the rate of lever pressing in rats. In another animal model, MK-801 aggravated a transgene-dependent abnormal behavior likely due to the fact that it indirectly stimulates corticallimbic glutamate output (McGrath et al., 2000). Although a number of studies have looked at the acute effects of these drugs and their effect on glutamate release and the mGluR5 receptor, the result of chronic administration on neurotransmitter and receptor level remains elusive. Therefore, it is of critical importance to have an understanding of the effect of these drugs in a healthy animal to assess and influence their ability to modify the glutamatergic system. Ketamine, on the other hand, has previously been used in a treatment setting in OCD, with promising results (Bloch et al., 2012; Rodriguez et al., 2013).

However, previous research has shown that the underlying working mechanisms of ketamine may be unique to ketamine and thus not shared by all NMDA receptor antagonists (Hillhouse and Porter, 2014). Indeed, ketamine's mechanism of action remains mysterious, as next to its NMDA blockade, ketamine disturbs a wide range of other intracellular neuronal processes (Sleigh et al., 2014).

In this study, we evaluated the potential of both MK-801 and ebselen to modulate glutamate receptors and neurotransmitter levels in healthy naïve rats by means of magnetic resonance spectroscopy (MRS), PET, and immunohistochemistry (IHC). This allowed us to investigate the effect on 1) the neurotransmitter level by visualization of glutamate and glutamine with MRS and 2) on the neuroreceptor level by visualizing the receptor availability of mGluR5 and distribution by PET, as well as to verify this with mGluR5 IHC. For future preclinical research, such adds to elucidating the role of glutamate in neuropathologies such as OCD, in particular in the QP model discussed earlier, to establish whether modifying the glutamate levels has an effect on the symptoms.

Materials and Methods

The present study was carried out in accordance with the European Communities Council Directive of November 24th, 1986 (86/609/EEC) for the care of laboratory animals and after approval of the local ethical committee (University of Antwerp under number 2014-18). All efforts were made to minimize suffering and to reduce the number of animals. Naïve male Sprague-Dawley rats (n=32, 285-550 g during the experiment; Harlan, Netherlands) were housed in a temperature-and humidity-controlled vivarium in individually ventilated cages with a 12-hour light-dark cycle. All experiments were performed during daytime. Food and water were available ad libitum. All applicable institutional and national guidelines for the care and use of animals were followed.

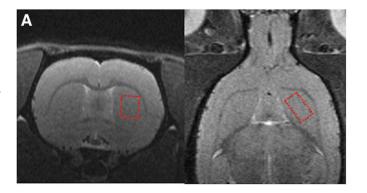
Experimental Setup. Rats were divided into three groups. Animals in the first group (n=12) received an intraperitoneal injection of MK-801 (0.3 mg/kg; dissolved in saline) every day at the same time (5–6 PM) for 7 consecutive days. This dosage was chosen based on previous work using MK-801 to induce glutamate toxicity (Kosten et al., 2016) and an acute study investigating a variety of dosages (Wyckhuys et al., 2013). Animals in the second group (n=12) were exposed to ebselen (10 mg/kg; dissolved in 0.5% carboxymethylcellulose) by oral gavage at the same time for 7 consecutive days also. The dosage of ebselen was based on previous studies (Dalla Corte et al., 2012; Yatmaz et al., 2013; Ozyigit et al., 2015; Oostwoud et al., 2016). Animals in the third group (n=8) received intraperitoneal injections of saline (1 ml/kg) and served as a control.

A day after the sixth injection of a solution, to ensure that the drug was cleared from the brain, animals in each group were anesthetized (induction of 5% isoflurane; maintained at 2% isoflurane mixed with medical oxygen) and subjected to a PET/computed tomography (CT) scan with (5S,10R)-(+)-5-Methyl-10,11-dihydro-5H-dibenzo[a,d]cyclohepten-5,10-imine (ABP-688), performed on two dedicated small-animal PET/CT scanners (docked PET/CT and Siemens Inveon multimodal PET/CT; Siemens Medical Solutions, Knoxville, TN) that had a reconstructed image resolution of approximately 1.2 mm at the center of the field of view. PET data were acquired for a total of 60 minutes in list-mode format (frames: 2×10 , 3×20 , 3×30 , 3×60 , 3×150 , and 9×300 seconds) with ABP-688 injected immediately after the start of the PET acquisition. After the PET scan, a CT image was acquired in three different bed positions. Small-animal µPET images were reconstructed by use of a two-dimensional ordered subset expectation maximization with four iterations and 16 subsets after Fourier

rebinning. The μ PET images were then reconstructed on a 128 \times 128 \times 159 grid with a voxel size of 0.776 \times 0.776 \times 0.776 mm. Normalization, dead time correction, random subtraction, CT-based attenuation, and single-scatter simulation corrections were applied (Watson, 2000).

The day after the seventh injection, animals underwent an MRS scan. All magnetic resonance (MR)—based measurements were performed on a seven Tesla Pharmascan (Avance III) with a horizontal bore (16 cm in diameter; Bruker Biospin, Ettlingen, Germany) equipped with a standard Bruker coil setup using a quadrature volume resonator for excitation and array rat head surface coil for signal detection. The system was interfaced to a Linux personal computer running TopSpin 3.1 and ParaVision 6.0.1 software (Bruker Biospin).

All rats were initially anesthetized using 5% isoflurane and maintained at 2% isoflurane in a mixture with medical oxygen. The volume of interest (VOI) for $^1\text{H-MRS}$ was localized using multislice Rapid Acquisition with Refocused Echoes (RARE) images using a TurboRARE pulse sequence (Hennig et al., 1986). A VOI (1.4 \times 2.2 \times 2.7 mm = 8.32 mm³) was placed in the caudate putamen (CP) of the rat brain, as shown in Fig. 2A, by using the Paxinos and Watson rat brain atlas (Paxinos, 2013) as a reference (between bregma ±-0.24 and ±1.92 mm). More details regarding this procedure can be found in the Supplemental Materials.



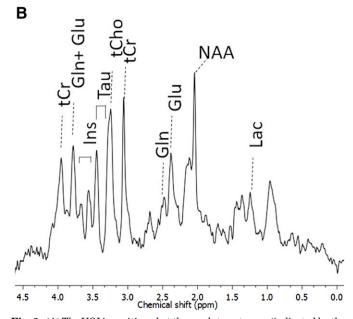


Fig. 2. (A) The VOI is positioned at the caudate putamen (indicated by the red boxes on T2 weighted axial and coronal images). (B) The major peaks are labeled on the representative spectra. Ins, inositol; Lac, lactate; NAA, *N*-acetylaspartate; PCho, Phospho-choline; Tau, taurine; total choline (tCho), Pcho + GPC; total creatine (tCr), PCr + Cr. The voxel size is 8.32 mm³.

Following the MRS scan animals, were sacrificed and their brains were removed and stored at $-80^{\circ}\mathrm{C}$. To verify the results of the PET scan, the brains of the animals in each group were subjected to a histologic verification with an mGluR5 staining. In total, eight animals in the ebselen group were compared with nine animals in the MK-801 group for the immunohistochemistry analysis, as one brain removal did not succeed.

PET Analysis. Dynamic images were processed using PMOD 3.3 (PMOD Technologies, Zurich, Switzerland). After cropping, the static image corresponding to the time-averaged frames of each dynamic acquisition was spatially transformed to an in-house-developed rat brain template for ABP in a brain normalization step (Verhaeghe et al., 2014). The obtained transformation was then applied to the dynamic image frames to equally transform those to the same ABP template. Extracerebral activity was removed by applying a brain mask available from the software. Time-activity curves of the investigated regions were extracted from the resulting dynamic images and used as the input for the simplified reference tissue model (Lammertsma and Hume, 1996), with the cerebellum as the reference. This kinetic modeling step subsequently produced the average binding potential (BPnd) of the VOI of each region. Parametric BPnd maps were generated by pixelwise kinetic modeling using the simplified reference tissue model 2. These BPnd output values were investigated by means of a VOI-based analysis (with a predefined rat brain VOI template available in the software), using a Mann-Whitney test, to compare the two conditions (MK-801 vs. Ebselen) crosssectionally. Additionally, a voxel-based analysis was performed using statistical parametric mapping on the aforementioned parametric BPnd maps. These data were analyzed using an unpaired two-sample t test with a cluster threshold of 125 voxels (1 mm³) and a significance level of 0.01. The resulting T maps were again overlaid on the aforementioned brain VOI template delineating the different brain regions displaying significant increases (hyper) in a red-yellow format and significant decreases (hypo) in a blue-green format.

MR Analysis. In vivo 1 H MR spectra were analyzed using an automated deconvolution program (LCModel) (Provencher, 2001) as described previously (Orije et al., 2015). The LCModel analysis was performed within the chemical shift range of 0.2–4.2 ppm. The following metabolites were included in the basis set of the LCModel (Fig. 2B): alanine, aspartate, creatine (Cr), phosphocreatine (PCr), γ -aminobutyric acid, glucose, glutamate (Glu), glutamine (Gln), glutathione, glycerophosphorylcholine (GPC), phosphorylcholine, inositol, lactate, N-acetylaspartate, N-acetylaspartylglutamate, scyllo-inositol, and taurine. In addition to this standard set of brain metabolites, LCModel includes nine stimulated macromolecule and lipid peaks (Pfeuffer et al., 1999) as shown in Fig. 2B. The results of macromolecules and lipids were not included in the study. The reliability of the metabolite quantification was assessed using the Cramér-Rao lower

bounds (CRLBs; estimated error of quantification) provided by LCModel (Table 1).

Poorly fitted metabolite peaks (CRLB of >50%) were not included in further analysis. Metabolites with CRLB ≤50% in at least 50% of the spectra were included in the neurochemical profile. Also, as neurometabolite levels fluctuate during the day, if the time of the MRS scan deviated from the originally foreseen time, the animal was not included in further analysis. This left us with a group size of 8 for the ebselen group, a group size of 10 for the MK-801 group, and a group size of 8 for the control group. A group correlation matrix for metabolites was created. If the correlation between two metabolites was consistently high (i.e., correlation coefficient higher than -0.5), the sum of the metabolites was reported (Provencher, 2001). We computed a correlation matrix for metabolites quantified for all animals to assess which metabolites could be reliably resolved from each other. If the negative correlation between two metabolites was very strong (i.e., correlation coefficient higher than -0.6), it was considered that these two metabolites cannot be individually quantified (Provencher, 2001). Therefore, their sum was reported. The amplitudes of Cr and PCr were strongly correlated: the corresponding matrix element equaled -0.9. The amplitudes of phosphorylcholine and GPC were strongly correlated: the corresponding matrix element equaled -0.9. For these metabolites, total creatine (Cr + PCr) and total choline (GPC) were reported. Due to the higher field, Glu and Gln could be quantified individually with good accuracy. Absolute concentrations (in micromoles per gram) of metabolites were determined relative to an unsuppressed water signal acquired from the same VOI (Minati et al., 2010). Further specifics on how these data were corrected can be found in the Supplemental Materials. For the purpose of this paper, we focused on the glutamate and glutamine levels and how these correlated with the other neurometabolites.

To estimate T2 maps of the CP, a rectangular region of interest $(\pm 0.05~{\rm cm}^2)$ was manually defined for the left and right CP separately using the Paxinos and Watson rat brain atlas (sixth edition) as a reference atlas. The data from left and right CP were averaged and recorded as the T2 relaxation time. The coronal images acquired with multi-slice multi-echo (MSME) sequence for T2 mapping were analyzed automatically with ParaVision 6.01 as described previously, using the image sequence analysis tool (Kara et al., 2015).

Immunohistochemistry. Coronal slices of the CP were made with a cryotome (Leica, Wetzlar, Germany) at a thickness of $20~\mu m$ on Superfrost Plus slides (Thermo Scientific, Merelbeke, Belgium). These slices were stored at -80° C until staining. The staining procedure can be found in the Supplemental Materials.

All pictures were taken with a Nikon Ti wide-field microscope (Nikon Instruments, Paris, France), digitalized by a Nikon Digital Sight DS-FI2 camera, using NIS Elements AR software, version 4.51.01 (Nikon Instruments).

TABLE 1 Metabolite concentrations (in micromoles per gram) and CRLB values

Metabolite	Concentration (mean \pm S.D.)			CRLB Control	CRLB Ebselen	CDI D MIZ 001
	Control $(n = 8)$	Ebselen $(n = 8)$	MK-801 (n = 10)	CKLB Control	CRLD Ebseien	CRLB MK-801
	μmol/g	μmol/g	μmol/g	%	%	%
tCr	6.1 ± 0.5	6.1 ± 0.3	6.0 ± 0.5	6	6	6
GABA	1.3 ± 0.3	1.5 ± 0.2	1.4 ± 0.4	29	24	26
Glc	1.3 ± 0.5	1.3 ± 0.3	1.4 ± 0.2	38	35	31
Gln	2.7 ± 0.3	2.2 ± 0.4	2.9 ± 0.5	18	22	18
Glu	8.4 ± 0.7	8.9 ± 1.1	8.8 ± 0.3	6	6	6
tCh	1.3 ± 0.1	1.4 ± 0.2	1.4 ± 0.1	9	8	8
GSH	1.1 ± 0.2	1.3 ± 0.5	1.4 ± 0.4	24	22	21
Ins	2.9 ± 0.5	2.7 ± 0.7	2.5 ± 0.6	14	16	17
Lac	1.3 ± 0.5	1.5 ± 0.7	1.4 ± 0.3	32	31	31
NAA	6.7 ± 0.4	6.4 ± 0.4	6.2 ± 0.5	6	6	6
Tau	9.0 ± 0.5	7.9 ± 0.6	8.3 ± 0.7	5	6	6

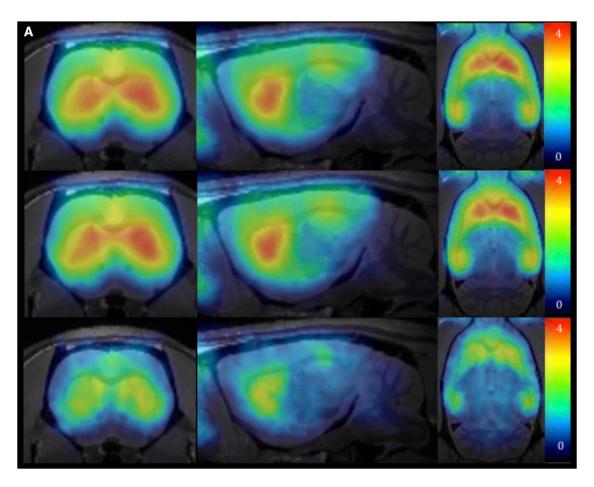
 $[\]operatorname{Glc}$, glucose; GSH , glutathione; Ins , inositol; Lac , lactate; NAA , N-acetylaspartate; Tau , taurine; tCh , total choline; tCr , total creatine.

Using ImageJ (version 1.47; National Institutes of Health, Bethesda, MD), mGluR5 receptor immunoreactivity was visualized. Analysis of the images consisted of the evaluation of the area fraction, which refers to the percentage of pixels in the image or selection that have been highlighted following a threshold, which was fixed at the same value for each image to eliminate any of the background.

Results and Statistical Analyses

Acute injection of MK-801 caused the animals of this group to display hyperactivity, consistent with what was previously described (Wu et al., 2005). No other unexpected

erratic behavior was displayed over the course of the experiment. During the course of the scans, no aberrant behavior was observed. The average weight of the animals receiving this drug at the start of the experiment was 326.42 \pm 11.88 g. At the end of the experiment, the average weight of these animals was significantly lower: 308.58 \pm 9.93 g. Animals receiving ebselen had an average weight of 331.33 \pm 11.02 g at the beginning of the experiment, which significantly decreased to 322.17 \pm 11.21 g at the end of the experiment. Animals of the control group had a weight of 353.96 \pm 9.94 g at the scanning time point, which was significantly higher than the weight at the start of the experiment (324.80 \pm 10 g).



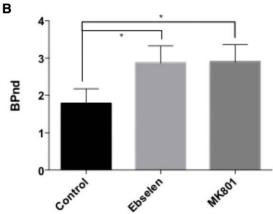


Fig. 3. (A) BPnd maps of ABP-688 for ebselen (top), MK-801 (middle), and saline (bottom) are displayed, with higher values displayed in red and lower values in a green-blue format. (B) BPnd values plotted showing the difference between the different groups.

PET Data. VOI-based analysis revealed a significant difference when comparing the BPnd in the CP between the control animals (1.97 \pm 0.39) and the animals receiving MK-801 (2.90 \pm 0.47, constituting an increase of 47.21% \pm 17.00%) or ebselen (2.87 \pm 0.46, constituting an increase of 45.69% \pm 16.37%) (Fig. 3).

This was confirmed with a voxel-based analysis (P=0.01, cluster size = 125 voxels) investigating all regions using statistical parametric mapping, displaying strong significant (P<0.01) increases after either drug in the frontal regions of the brain, in particular the majority of the CP, the cortex, hippocampus, and amygdala. Furthermore, there were significant (P<0.01) decreases in the majority of the cerebellum (Fig. 4).

Immunohistochemistry. Analysis revealed a significant (P < 0.05) difference in area percentage in the CP between control and ebselen, with an area fraction of $60.83\% \pm 6.30\%$ for ebselen and $50.21\% \pm 5.71\%$ for saline (Fig. 5). MK-801 had an area fraction of $57.14\% \pm 9.23\%$, which was not significantly different but with a tendency to increase.

MRS Data. No significant changes in glutamate level were found when comparing chronic ebselen and chronic MK-801 to control. However, a significant difference (P < 0.05) in the levels of glutamine between the ebselen and control animals was found. Animals treated with ebselen had a mean glutamine concentration of $2.2 \pm 0.4~\mu$ mol/g, which was $18.52\% \pm 5.42\%$ lower than animals in the control group who had a mean of $2.7 \pm 0.3~\mu$ mol/g. Animals of the MK-801 group had a mean of $2.9 \pm 0.5~\mu$ mol/g, which was $7.41\% \pm 2.10\%$ higher than the control animals. Also, when considering the ratio of glutamate versus glutamine, a significant difference (P < 0.01) was

present between the ebselen and control groups with a mean of 4.1 \pm 0.6 for ebselen, which was 32.26% \pm 8.88% higher than the ratio for the control animals (3.1 \pm 0.4; Fig. 6). This ratio was correlated (P < 0.05) with glutamine but not with glutamate.

A correlation between the MRS signal and the BPnd of the CP revealed no significant correlation with any of the neurometabolites (P > 0.05).

The mean T2 values of the CP of the rats receiving ebselen and MK-801 were not statistically significant compared to the rats receiving control (P < 0.05, two-tailed Student's test, n=8 for each group). The values were estimated as 40.76 ± 0.74 ms, 40.95 ± 0.80 ms, and 41.255 ± 0.68 ms for the groups exposed to MK-801, ebselen, and control, respectively.

Discussion

This study evaluated the potential of both ebselen and MK-801 to affect the glutamate receptors and neurotransmitter levels in healthy naïve rats, as possible modulators for the preclinical QP-induced OCD model. Previous work established that ebselen has a direct effect on the glutamatergic system by reducing the K⁺-evoked release of glutamate (Nogueira et al., 2002; Moretto et al., 2007), while low dosages of MK-801 selectively block NMDA receptors in the GABAergic interneurons, thus increasing glutamate release (Xi et al., 2009). A previous study performed by our laboratory confirmed glutamate increased when MK-801 was given acutely. Other NMDA antagonists, such as ketamine, when administered acutely, also increase glutamate (Wyckhuys et al., 2013).

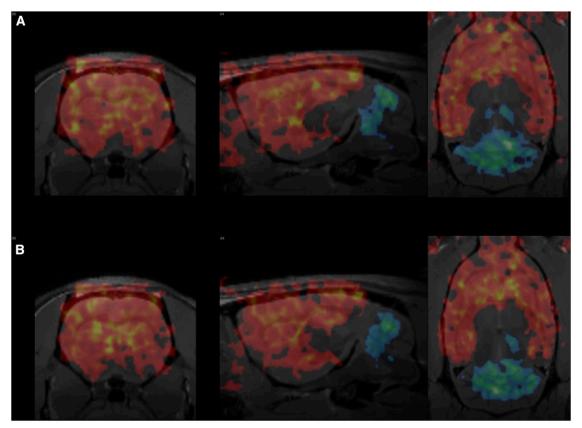
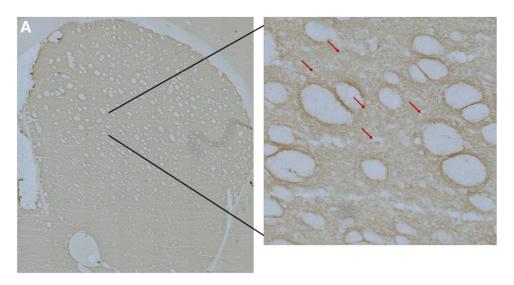
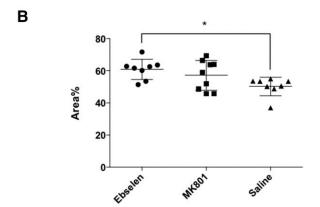


Fig. 4. Statistical parametric mapping maps of ABP-688 for both ebselen (A) and MK-801 (B) versus saline, with significantly higher t values (hypermetabolism) displayed in red and significantly lower t values (hypermetabolism) in a blue-green format.





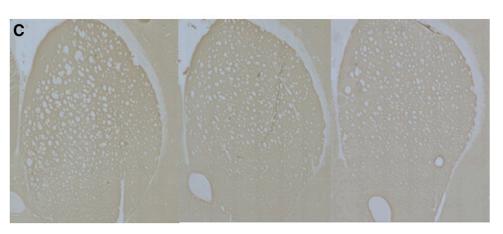


Fig. 5. (A) Overview image of the CP after immunohistochemistry (original magnification, $20\times$) with the mGluR5 antibody. Arrows point to cells with densely stained membranes rich in mGluR5. (B) Investigated parameter of the immunohistochemistry is presented here (area percentage) after chronic ebselen (n=8), chronic MK-801 (n=9), and saline (n=8). Individual points, median, and S.D. of each group are displayed. (C) Overview images of the three different conditions (left to right: ebselen, MK-801, saline). *P<0.05.

In contrast, MK-801 appears to decrease glutamate levels when administered chronically, which is potentially caused by chronic blockade of the NMDA receptor, making non-NMDA Glu receptors discompensatory and decreasing the terminal excitability of glutamatergic neurons, resulting in an attenuation of the Glu levels (Tsukada et al., 2005; Zuo et al., 2006). Alternatively, this decrease of Glu might also be caused by postsynaptic inhibition of other neuronal systems to NMDA receptor–mediated neurotransmission (Tsukada et al., 2005; Zuo et al., 2006). Another explanation can be found in the upregulation of Glu transporters (Tsukada et al., 2005; Zuo et al., 2006).

After chronic ebselen, the increase in mGluR5 area fraction measured by IHC was significant, indicating more mGluR5 distribution, possibly as a result of chronically less endogenous glutamate. For MK-801, a similar trend is noticeable, albeit not significant with the current power.

Also, the PET/CT results showed significant increases in BPnd after chronic administration of both ebselen and MK-801, also indicating more availability of the mGluR5 receptor. MK-801 is not the only NMDA antagonist that has been investigated by ABP-688. Previously, a clinical study used this tracer to investigate the acute effect of ketamine (0.23 mg/kg over 1 minute, then 0.58 mg/kg over 1 hour) with

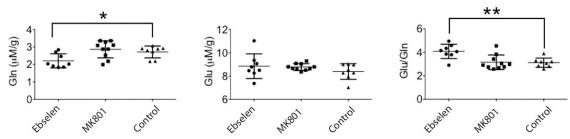


Fig. 6. The investigated MRS parameters (glutamate, glutamine, and the ratio between the two) extracted from the MRS spectrum are plotted here for both ebselen (n = 8) and MK-801 (n = 10). Individual points, median, and S.D. of the group are displayed. *P < 0.05; **P < 0.01.

the intention of increasing Glu, and reported significant decreases in BPnd. As ABP-688 binds at different sites on the receptor and hence does not compete with glutamate, at first sight one could hypothesize that glutamate fluctuations do not influence the BPnd. However, it should be noted that the affinity of the receptor for the tracer is in part determined by the endogenous glutamate. For example, DeLorenzo et al. (2015) used N-acetylcysteine, which is known to increase extracellular glutamate, and reported decreases in BPnd. Similarly, Zimmer et al. (2015) used ceftriaxone to decrease extracellular levels of glutamate and reported increases in the BPnd of ABP-688. Therefore, our results, displaying strong increases in BPnd after treatment with ebselen or MK-801 when compared with control, indicate lower levels of extracellular glutamate that positively affect the affinity of the tracer for mGluR5, consistent with previous data (Tsukada et al., 2005; Zuo et al., 2006). Thus, it is possible that the increase in mGluR5 distribution in the PET/CT results was overestimated due to the increase in tracer affinity caused by a decrease in extracellular glutamate. Alternatively, as NMDA and mGluR5 are interconnected in the brain, it is plausible for the changes induced by MK-801 to be present only on the level of receptor interaction and not on glutamate release.

In contrast to the results from the PET/CT and immunohistochemistry, indicating decreased levels of glutamate, the MRS scan showed that the chronic exposure to either drug did not affect glutamate levels in such a way that it differed from the control condition in the visualized time frame. When considering the PET/CT and immunohistochemistry results, this seems to contradict the hypothesis of a decrease in glutamate. However, it is important to note that MRS visualizes the total glutamate pool as it does not discriminate between intracellular and extracellular neurochemicals. The concentration of the neurochemicals is mostly driven by the intracellular concentration due to the negligible volume of the extracellular space. Thus, it could be that the release of glutamate in the extracellular space is, in fact, changed, directly affecting the glutamate receptors as visualized by PET/CT and immunohistochemistry. Interestingly, when considering this flux of Glu and Gln, there were significant decreases in Gln after chronic injection with ebselen. Because animals are repeatedly exposed to the drug, certain compensatory changes could occur. As glutamate is readily converted into glutamine through the glutamine synthetase pathway by astrocytes, it is possible that, due to decreased glutamate levels after chronic administration of ebselen, the available glutamine was converted into glutamate to preserve a normal level of glutamate, thereby depleting the Gln pool to compensate for this as the acute effect of ebselen wore off. Because the

amount of glutamate present in the synapse is maintained homeostatically at a low level at all times, with 85% of glutamate being converted to glutamine (Hertz, 2013) and ebselen acutely decreasing Glu levels, this could explain the distorted balance in the Glu/Gln ratio after administration of ebselen. Although this ratio had a tendency to increase after chronic administration of MK-801, this was not significant in this particular group.

In conclusion, both ebselen and MK-801 affect the mGluR5 receptor when administered chronically in the dosages described in this study. Although no direct effects on total glutamate were visualized with MRS, the changes in glutamine do hint toward changes in the total Glu-Gln pool.

Taken together, these two drugs are interesting modulators for further investigation in glutamatergic pathologies, such as obsessive compulsive disorder, Fragile X, epilepsy, and autism, and for further elucidation of their working mechanisms to optimize their use and potential.

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Authorship Contributions

Participated in research design: S. Servaes, Kara, Glorie, S. Staelens. Conducted experiments: S. Servaes, Kara.

Performed data analysis: S. Servaes, Kara.

Wrote or contributed to the writing of the manuscript: S. Servaes, Kara, Van Der Linden, S. Stroobants, S. Staelens.

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SUPPLEMENTARY TO: IN VIVO PRECLINICAL MOLECULAR IMAGING OF REPEATED EXPOSURE TO AN NMDA ANTAGONIST AND A GLUTAMINASE INHIBITOR AS POTENTIAL GLUTAMATERGIC MODULATORS.

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1. PET acquisition

Throughout the experiment, respiration rates were visually monitored. Body temperature of the animals was maintained by using a heated pad.

The PET data were reconstructed both into a single static image of the full 60 minutes and into 33 temporal bins of increasing duration (12×10 , 3×20 , 3×30 , 3×60 , 3×150 , and 9×300 seconds) using a two-dimensional ordered subset expectation maximization (2D-OSEM) with 4 iterations and 16 subsets after Fourier rebinning. The μ PET images were then reconstructed on a $128 \times 128 \times 159$ grid with a voxel size of 0.776 \times 0.776 \times 0.796 mm. Normalization, dead time correction, random subtraction, CT-based attenuation and single-scatter simulation corrections were applied (Watson, 2000).

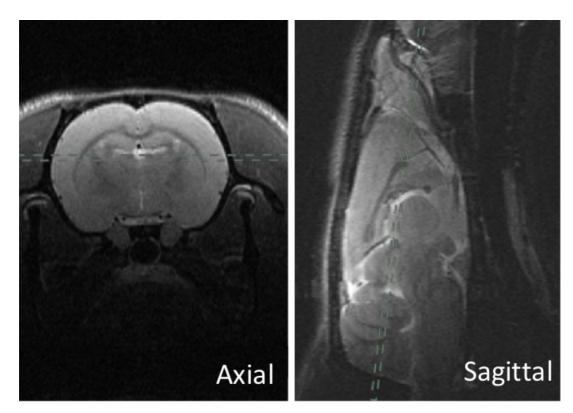
2. MR acquisition

Throughout the MR studies, respiration rate was constantly monitored by a pressure sensitive respiration sensor, placed under the abdomen. Body temperature was monitored with a rectal temperature probe and was maintained at 37.0 ± 0.5 °C using a warm air system with a feedback unit (SA Instruments, NY, USA). The respiration and body temperature control systems were connected to a computer with pcSam monitoring software (SA Instruments, NY, USA), and these parameters were monitored throughout the acquisition.

The TurboRARE images were acquired in three orthogonal directions (axial, coronal and sagittal) with following parameters: repetition time (TR) = 2500 ms, echo time (TE)

= 33.442 ms, matrix size (256 x 192), field of view = (20 x 20) mm2, 16 slices, slice thickness = 0.5 mm, RARE factor = 8.

Consistent voxel placement for 1H-MRS was achieved in each rat by using anatomical markers such as the external capsule, lateral ventricle, and anterior commissure in three orthogonal RARE slices. While positioning the voxel on the brain, T2-weighted RARE images at three orthogonal position were checked to avoid contribution of cerebral spinal fluid and white matter volume fractions. MAPSHIM was employed to automatically adjust first- and second-order shim coils for the VOI. Spectral line widths of water around 9.8 - 11.7 Hz were obtained. In vivo localised 1H MR spectra were acquired with localisation by Adiabatic Selective Refocusing (LASER) sequence in combination with outer volume suppression (OVS) and VAPOR water suppression (Tkáč et al., 1999; Garwood and DelaBarre, 2001). The following parameters were used: TR = 4000 ms, TE = 20.46 ms, number of averages = 512, water (4.7 ppm) as frequency on resonance, spectral width = 8012.82 Hz, number of acquired points = 2048 yielding a spectral resolution of 1.96 Hz/points. The total acquisition time amounted to ± 34 minutes. Each MRS scan was, during acquisition, dynamically frequency drift corrected. To this end, the water unsuppressed signal was used as an in vivo reference. For each animal, an unsuppressed water signal (TE = 20.46 ms, TR = 4000 ms, 16 averages, scanning time = \pm 1 min, water suppression turned off) was acquired immediately after acquiring the water-suppressed spectrum. During postprocessing, this unsuppressed water signal was used for Eddy-current correction by the LCModel, where the water-suppressed-MRS signal (in time domain) is point-wise divided by the phase part of the unsuppressed water signal (Mandal, 2012).



Supplementary Figure 1 The figures depict T2 weighted anatomical images (axial and sagittal). For T2 relaxometry, a single slice is positioned parallel to the cerebral cortex using the anatomical markers (i.e olfactory bulb, cerebral cortex and corpus callosum) allowing the slice to cover the caudate putamen.

3. MR analysis

Due to some presumption of LC model default set up, the concentration units acquired from LC model needed to be corrected to acquire absolute μ mol/g unit. LC model default set up assumed that water attenuation factor (ATTH20) 0.7 and metabolite attenuation factor was 1, and NMR visible water concentration of white matter as 35880 mM (tissue water reference based on results of (Ernst *et al.*, 1993)).

To remove the default correction of LC model, first we multipled the concentration output values of LC model with $\left[\left(\frac{1}{35880X0.7}\right)\right]$. After removal of default correction, this value was multipled with water attenuation factor which was calculated using the following formulate ATTH20= exp (-TE/T2). TE and tissue T2 from the current study were used for this estimation. Third, the absolute concentrations of metabolites were corrected for partial volume effects by taking the volume fractions of cerebral spinal fluid, white and gray matter into account in the quantification (Provencher, 1993, 2014). These volume fractions were used to correct the total water concentration for each VOI using the following equation: water concentration = (43300 x fraction gray matter + 35880 x fraction white matter + 55556 x fraction cerebrospinal fluid) / (1fraction cerebrospinal fluid) (Provencher, 1993). No visible white matter and/or cerebral spinal fluid contribution was detected after inspection of T2 weighted images which were used to localize the MRS voxel. Therefore, in this study the mean water tissue concentrations (from the gray matter) as a reference was assumed as 43300 mM. Since we employed long TR and short TE, correction for relaxation effects of metabolites had minimum effect on metabolite concentration (Provencher, 2014). Therefore, metabolic T1 and T2 relaxation correction was neglected. In addition, T1 relaxation time correction for water attenuation was neglected.

Tissue water T2 relaxation time was estimated using multi slice multi echo (MSME) sequence with following parameters: TR = 2500 ms, TE = 8 ms, number of averages = 1, echo spacing = 8ms, echo images = 20, image size = 256x 192, field of view = 35 mm x 35 mm, slice number = 1, slice thickness = 0.4 mm and acquisition time = 8 min.

Anatomical markers (i.e. olfactory bulb, cerebral cortex and corpus callosum) were used to position the single slice to cover the CP.

4. Immunohistochemistry

Before staining, slides were acclimatized at room temperature during 5 min before being fixed in 4% paraformaldehyde (PFA) during 10 min. They were washed 6 times in PBS (pH 7.4) for 5 min each after which they were immersed in water containing 3% H₂O₂ for 15 min at room temperature to quench endogenous peroxidases. After three washes of 5 min, each in PBS, sections were preincubated with 10% normal donkey serum (NDS) and 0.3% Triton X-100 in PBS at room temperature for 1 h. After 3 washes in PBS for 5 min, the slides were blocked with avidin during 15 min, rinsed briefly and subsequently blocked with biotin during 15 min, followed by incubation with polyclonal mGluR5 primary antibody (1/2500 in PBS with 1% NDS, 10% milk, 0.2% Triton X-100 and 1% BSA; AB5675 Millipore) overnight at room temperature. Subsequently, after three 5 min rinses in PBS, sections were incubated with ExtrAvidin peroxidase 1/1000 for 1 h. After three washes of 5 min each in PBS, the peroxidase reaction was carried out with DAB reagent (Dako) for 2 minutes. The reaction was stopped in tap water. The slides were dehydrated through subsequent immersion in 80%, 95% and two changes of 100% etOH for 5 min each, and finally immersion in two changes of Xylene before mounting with DPX mounting medium (Sigma Aldrich) (Kosten *et al.*, 2016).

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