



Modulation of Monoaminergic Systems by Antidepressants in the Frontal Cortex of Rats After Chronic Mild Stress Exposure

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Abstract

The standard pharmacological treatment of the major depressive disorder (MDD) is still grounded in a monoaminergic approach. Consequently, antidepressant treatments pursue to heighten serotonergic and noradrenergic neurotransmissions. Thus, the aim of this study was to assess the impact of exposure to a well-characterized animal model, the chronic mild stress (CMS) on serotonin (5-HT) and noradrenaline (NE) levels, and reuptake transporters and receptors in the frontal cortex (FC) of CMS-exposed rats. Moreover, considering the diverse pharmacological profiles of existing antidepressants and the large number of patients not responding to treatments, we have investigated whether generally utilized antidepressants can modulate their expression. Male Wistar rats were exposed to CMS and some of them treated with desipramine, escitalopram, or duloxetine. Possible changes in the described monoaminergic neurotransmission elements were evaluated. CMS induced differences in the expression of reuptake transporters and receptors involved in the monoaminergic neurotransmission pointing towards the weakening of their signaling. CMS antidepressant-treated rats showed an improvement of the monoaminergic tone, being desipramine and duloxetine more influential than escitalopram over noradrenergic elements and having the three antidepressant-tested effects on serotonergic transmission. In summary, there are molecular alterations on the monoaminergic neurotransmission in FC induced by CMS exposure. Besides, antidepressant treatments modulate the elements of these neurotransmission systems differentially.

Keywords Chronic mild stress · Depression · Frontal cortex · Antidepressants · Monoamine receptors · Monoamine transporters

Introduction

Major depressive disorder (MDD) affects over 322 million people worldwide being one of the leading causes of disability with enormous personal, medical, and economic costs [1].

However, the etiology and pathophysiology of this disease are still unknown, although different hypotheses (not exclusive but complementary) have been proposed, trying to shed light on its mechanisms [2].

Near one-third of patients with MDD are treatment-resistant to the available drugs, whose majority is still based on the classic monoaminergic hypothesis. Consequently, current treatments try to enhance serotonergic and noradrenergic neurotransmission through actions on degradation enzymes (monoamine oxidase inhibitors—MAOIs), reuptake transporters (tricyclic antidepressants—TCAs—and selective reuptake inhibitors), receptors (multi-receptor agonists or antagonists), or on these last two elements simultaneously (serotonin partial agonist reuptake inhibitors—SPARIs) [3].

Serotonin (5-HT) is one traditional therapeutic target of study in MDD. Nevertheless, there is still controversy on the detailed implications of this complex receptor system and how to overcome the technical difficulties of drug development due to the heterogeneous distribution and functionalities of the different 5-HT receptor types. To date, seven families

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and 14 5-HT receptors have been described with different expression and functional implications depending on their cerebral location, molecular features, second messengers, and affinity for their ligands among other factors (Barnes et al., IUPHAR database). Within them, the 5-HT_{1/2/7} families have arisen as the most connected with depressive-like behavior and antidepressant effect [4].

Norepinephrine (NE) is other neurotransmitter classically involved in the development of depressive-like behaviors. There are α - and β -adrenoceptors and up to nine subtypes with different implications and affinities for their ligands [5]. Given their widely spread location throughout the central nervous system (CNS), including the frontal cortex (FC), it has been shown that noradrenergic signaling participates in many different processes, some of them intimately bounded to depression (e.g., stress and immune responses, memory consolidation, plasticity, and sleep regulation) [6].

New drugs designed for the treatment of depression aim to modulate serotonergic and noradrenergic transmissions acting on highly specific targets. Studies about the actual decrease on the monoamine metabolites in biological fluids and the CNS in mental diseases convey the idea that antidepressants might improve the monoaminergic transmission modifying the expression of monoamine receptors and reuptake transporters [7]. This can contribute to the fine regulation needed for their efficacy due to the serious side effects arising when monoamine signaling is deficient or overstimulated [8].

Antidepressants (ADs) might modulate other biological systems worth to study; antioxidant [9], anti-inflammatory [10], epigenetic [11], neurotrophic [12], and mitochondrial [13] actions have been already described for ADs [14].

There is a profuse volume of published studies dealing with serotonin and norepinephrine neurotransmissions and their modulation by ADs. Nevertheless, these investigations focus on some particular elements of these signaling systems and, given the cross-regulation among all of them, these previous approaches would be partially explaining some of the conflicting results obtained.

The first aim of this study was to evaluate the impact of exposure to a well-characterized multidimensional animal model, the chronic mild stress (CMS) on 5-HT and NE levels, as well as the expression of their respective reuptake transporters and receptors in the FC of rats. Besides, available antidepressants have heterogeneous pharmacological profiles and there is an important number of patients not responding to classical treatments. Thus, new molecular insights in the activity of classic ADs over the monoaminergic neurotransmission are essential to improve ADs design and make it more accurate, powerful, and with a better tolerance profile. Hence, the second aim of this study was to examine whether commonly used antidepressants can modulate the expression of the monoaminergic elements explored in the FC.

Materials and Methods

Animals

Male outbred Wistar Hannover rats (HsdRccHan:Wist, from Envigo, Spain) initially weighing 200–225 g were housed one per cage. Animals were maintained under standard conditions of temperature and humidity and in a 12-h light/dark cycle (lights on at 08:00 h) with free access to food and water. Rats were handled daily for the change of cage and bedding for 14 days before the beginning of the stress protocol. All experimental procedures adhered to the guidelines of the Animal Welfare Committee of the Universidad Complutense (PROEX421/15) following European legislation (2010/63/EU) and they were carried out in the Animal Facility CAI-UCM. Animal studies are reported in compliance with the ARRIVE guidelines and all efforts were made to minimize animal suffering and to reduce the number of animals used.

Experimental Groups

Animals which were randomly assigned to the following groups ($n = 6–8$ in each group) were used: (1) a control (animals were handled for few seconds once at 10:00 h) group (CT); (2) a control group with intraperitoneal injection (i.p.) of sterile saline (used as vehicle) injected daily for 7 days (CT+Veh); (3) a chronic mild stress (CMS) group; and (4) a CMS injected (i.p.) with vehicle (CMS+Veh) group (days 14–21).

For experiments involving the i.p. injection of ADs (from day 14 to day 21), three additional experimental groups were employed: (5) a CMS group injected with desipramine (CMS+Desip); (6) a CMS group injected with escitalopram (CMS+Escit); and (7) a CMS group injected with duloxetine (CMS+Dulox). See the “[Pharmacological Tools](#)” section for ADs-specific details.

Vehicle-injected groups did not differ from the same experimental groups without injection in any of the parameters analyzed, and their values have been pooled in all the figures.

Chronic Mild Stress (CMS) Protocol

The CMS protocol used was a modification of the one proposed by Willner [15]. Our group has previously used this CMS protocol showing that it induces a depressive-like behavior when analyzed by means of the modified forced swim test, sucrose test, splash test, and elevated plus maze test, as well as a response to the administration of the ADs here employed [9, 16, 17]. The protocol consists of a series of different stressors that were changed daily (two stressors/day), given in an unpredictable basis, for a period of 21 days. The stressors included the following: (a) food deprivation, (b) water deprivation, (c) cage tilting, (d) soiled cage, (e) grouped housing after a period of water deprivation, (f), stroboscopic

illumination (150 flashes/min), and (g) intermittent illumination every 2 h.

Pharmacological Tools

The employed ADs (from Sigma-Aldrich, Spain) were the tricyclic antidepressant (TCA) Desipramine hydrochloride (D3900) (20 mg/kg, i.p.), the selective serotonin reuptake inhibitor (SSRI) Escitalopram oxalate (E4786) (15 mg/kg, i.p.), and the serotonin-norepinephrine reuptake inhibitor (SNRI) (S)-Duloxetine hydrochloride (SML0474) (15 mg/kg; i.p.).

In order to explore therapeutic effects of the drugs, we choose a protocol close to clinical reality: vehicle or ADs were administered during the last 7 days of the CMS protocol. The administered doses have shown antidepressant activity in rodents in previously published studies using similar protocols. The drugs employed are neither new nor experimental drugs; these are well-established ADs, widely used in preclinical studies and in the treatment of patients, and without effects on non-stressed animals. Accordingly, and in order to reduce the number and the suffering of animals following the 3Rs principles, we did not perform behavioral tests on these rats. The result that these tests would yield, that the CMS protocol induces a depressive-like behavior, and that treatment with canonical antidepressants (i.e., TCAs, SSRIs, and SNRIs) reverses the behavior induced by the stress protocol has been extensively published with these drugs and doses in this model [18, 19].

Tissue Samples

To avoid variations in corticosterone levels caused by the circadian rhythm, biological samples were always obtained at the same time of the day, namely between 3 and 4 p.m. Samples from CMS-exposed animals were taken immediately after day 21 of stress.

After terminal anesthesia using sodium pentobarbital (320 mg/kg i.p. Vetoquinol®, Spain; CN:570681.8), the brain was removed from the skull and frontal cortical areas were excised from the brain and frozen at $-80\text{ }^{\circ}\text{C}$ until assayed.

Homogenization of the Samples

A widely utilized method that provides a high-purity cytosolic fraction, practically without nuclear contamination, was employed [20]. Briefly, FC was homogenized in 300 mL buffer [10 mmol/L N-2-hydroxyethylpiperazine-N-2-ethanesulfonic acid (pH 7.9); 1 mmol/L EDTA, 1 mmol/L EGTA, 10 mmol/L KCl, 1 mmol/L dithiothreitol, 0.5 mmol/L phenylmethylsulfonyl fluoride, 0.1 mg m/

L aprotinin, 1 mg/mL leupeptin, 1 mg/mL Na-p-tosyll-lysinechloromethylketone, 5 mmol/L NaF, 1 mmol/L NaVO_4 , 0.5 mol/L sucrose, and 10 mmol/L Na_2MoO_4]. After 15 min, Nonidet P-40® (Roche, Mannheim, Germany) was added to reach a 0.5% concentration. The tubes were gently vortexed for 15 s, and nuclei were collected by centrifugation at 5000g for 5 min. Supernatants were considered as the cytosolic fraction. All steps of the fractionation were carried out at $4\text{ }^{\circ}\text{C}$.

Western Blot Analysis

To determine the expression levels of serotonin transporter (SERT) and norepinephrine transporter (NET), cytosolic extracts from FC samples were used (see previous section for details).

After adjusting protein levels (see the “Protein Assay” section) in the homogenates and mixing them with *Laemmli* sample buffer (Bio-Rad, Hercules, CA, USA), 20 mL (1 mg/mL) was loaded and the proteins were size-separated in 10% SDS-polyacrylamide gel electrophoresis (90 V).

After the gel electrophoresis, the membranes were blocked in 30 mL Tris-buffered saline containing 0.1% Tween 20 and 5% skim milk/BSA, then the membranes were incubated with specific primary antibodies against SERT (sc-1458, Santa Cruz Biotechnology, 1:500) and NET (sc-67,216, Santa Cruz Biotechnology, 1:750). After washing with a TBS-Tween solution, the membranes were incubated with the respective horseradish peroxidase-conjugated secondary antibodies for 90 min at room temperature and revealed by ECLTM-kit following the manufacturer’s instructions (Amersham Ibérica, RTN2236; Spain).

Blots were imaged using an Odyssey® Fc System (LI-COR Biosciences) and quantified by densitometry (NIH ImageJ® software). All densitometries are expressed in arbitrary units of optical density (O.D.). Several exposition times were analyzed to ensure linearity of the band intensities. Loading controls (blots shown in the respective figures) were β -actin (Sigma A5441).

Protein Assay

Protein levels were measured using Bradford method based on the principle of protein-dye binding.

PCR Analysis

Primer oligonucleotides for PCR were designed with the Primer3 tool [21]. Target specificity was checked by *in silico* PCR using the USCS GenomeBrowser [22] and

Blast (NCBI) for cDNA and gDNA; only primer pairs with no unintended targets were selected (Table 1).

PCR analyses were carried out homogenizing frontal cortex in 600 μ L of TRIZOL® reagent (Invitrogen, Life Technologies, CA, USA) in the TissueLyser LT (QUIAGEN®, Venlo, Netherlands). Frequency used was 50 oscillations per second for 5 min at 4 °C.

Total cytoplasmic RNA was prepared from samples following TRIZOL® datasheet; aliquots were converted to cDNA using random hexamer primers. Semi-quantitative changes in mRNA levels were estimated by real time-PCR (RT-PCR) using the following cycling conditions: 35 cycles of denaturation at 95 °C for 10 s, annealing at 58–63 °C for 15 s depending on the specific set of primers, and extension at 72 °C for 20 s. Reactions were carried out in the presence of SYBR green (Quantimix Easy Master Mix Biotools, B&M labs 10607-4154) in a 20- μ L reaction in a Rotor-Gene (Corbett Research, Mortlake, NSW, Australia). Relative mRNA concentrations were calculated from the take-off point of reactions using included software, and tubulin levels used to normalize data.

As some of target genes have only one exon, the intron-spanning primer design was impossible in these cases. For this reason, purity and impact of possible genomic DNA contamination were monitored using both RNA samples and ValidPrime™ protocol (TATAA Biocenter, Odinsgatan, Sweden) as control for genomic background.

Serotonin and Norepinephrine Brain Levels

Serotonin and norepinephrine levels in the FC tissue homogenate (see the “[Homogenization of the Samples](#)” section) were measured using commercially available ELISA kits (Labor Diagnostika Nord, Nordhorn, Germany) following the manufacturer’s instructions.

Chemicals and Statistical Analyses

Unless otherwise stated, the chemicals were from Sigma-Aldrich (Spain). Data are expressed as mean \pm SEM and a one-way ANOVA with a Tukey post hoc test was employed for comparisons between groups. Data were analyzed using the Brown-Forsythe test to assess Gaussian distribution. In those cases in which the data did not follow a Gaussian distribution, a nonparametric ANOVA with a Kruskal-Wallis test followed by a Dunn’s post hoc test was performed. A p value < 0.05 was considered statistically significant.

Results

Effects of CMS and Antidepressant Treatments on the Whole Serotonin (5-HT) and Norepinephrine (NE) Levels in the FC

No differences were detected in 5-HT levels among all the experimental groups (one-way ANOVA $F_{(4,33)} = 1.598$, $p = 0.1980$; Fig. 1a).

Whole NE levels did not change after CMS, but desipramine decreased NE amount compared with the CMS and CMS+Escitalopram groups (one-way ANOVA-Tukey $F_{(4,38)} = 5.198$, $p = 0.0019$; Fig. 1b).

Effects of CMS and Antidepressant Treatments on the Expression of Serotonin and Norepinephrine Transporters in the FC

CMS did not affect the serotonin transporter (SERT) protein expression in the FC. Desipramine and escitalopram treatments did not change the expression of the SERT but Duloxetine upregulated the SERT expression compared with the control and CMS+Desipramine groups (one-way ANOVA-Tukey $F_{(4,36)} = 5.349$, $p = 0.0017$; Fig. 2a).

The CMS exposure upregulated the expression of the norepinephrine transporter (NET) in the FC (Kruskal-Wallis test-Dunn, KW statistic = 14.57, $df = 4$, $p = 0.0057$; Fig. 2b). NET protein expression was not affected by treatments, but they seemed to prevent the increase induced by the CMS, being the CMS+Duloxetine group, the one that showed a clearer trend towards avoiding the CMS-induced NET increment.

Effects of CMS and Antidepressant Treatments on the Expression of the Serotonin Receptors 5-HT1A, 5-HT1B, 5-HT2A, 5-HT2B, 5-HT2C, and 5-HT7 in the FC

CMS increased mRNA expression levels of 5-HT1A and the treatments seemed to blunt the effects of CMS on 5-HT1A although without statistical differences (one-way ANOVA-Tukey $F_{(4,38)} = 2.85$, $p = 0.0368$; Fig. 3a). The CMS exposure had no effect on 5-HT1B mRNA but desipramine induced an upregulation in 5-HT1B mRNA levels compared with control (Kruskal-Wallis test-Dunn, KW statistic = 18.13, $df = 4$, $p = 0.0012$; Fig. 3b).

The postsynaptic receptor 5-HT2A did not change under CMS conditions but desipramine and duloxetine upregulated its levels compared with the control and CMS groups (one-way ANOVA-Tukey $F_{(4,39)} = 7.579$, $p = 0.0001$; Fig. 3c). 5-HT2B increased in the FC after CMS but none of the antidepressants affected the induction of the expression of 5-HT2B caused by CMS,

Table 1 Primer sequences used for PCR analyses

Protein	Forward	Reverse
5-HT1A	TGGGTACTCTCATTCTG	CAGCACTGATACCATGAG
5-HT1B	GGTGGACTATTCTGCTAAA	GGAGTAGACCGTGTAGAG
5-HT2A	ATGCTGCTGGGTTCTTGT	TTGAAGCGGCTGTGGTGAAT
5-HT2B	GGCCAATCAGTGCAATCCC	AAAGCAGCCAGTGACCCAAA
5-HT2C	CCAACGAACACCTTCTTCC	GCATTGTGCAGTTTCTTCTCC
5-HT7	TCTCTTCGGATGGGCTCAGA	TTCCTGGCGGCCTTGTAAT
α 1A	TGTCCTGCGAATCCAGTGTC	TGTCTCTGTGCTGTCCCTCT
α 1B	TCCGCTACCCTAGAAGTGCT	TAGCGGTCAATGGAGATGGC
α 1D	TCTTCGTCCTGTGCTGGTTC	AGATGAGCGGGTTCACACAG
α 2A	TGGCAACGTGCTGGTTATTATCGC	TCTCGCACCACACCTTACCAAAGT
α 2B	TCCTGATCACTGGCATTACCCCTT	ACATTTGCTGGCTCTCTAGCTCGT
α 2C	ATCAGGACTTCAGGCGCTCTTCA	AACTCTGGAGAAGCCCACTGGTT
β 1	AGACGCTCACCAACCTTTCATCA	ACATAGCACGTCTACCGAAGTCCA
β 2	GGTCATCACAGCCATTGCCAAGTT	GGCTGTGACGCACAACACATCAAT
β 3	GTCTTCTGTGCAGTACGGT	CCTGTTGAGCGGTGAGTCT

although duloxetine made its levels similar to control (one-way ANOVA-Tukey $F_{(4,39)} = 6.674$, $p = 0.0003$; Fig. 3d). No effects were detected in 5-HT2C mRNA (one-way ANOVA-Tukey $F_{(4,37)} = 1.163$, $p = 0.3427$; Fig. 3d).

There were no significant differences in 5-HT7 mRNA when all the groups were analyzed (one-way ANOVA-Tukey $F_{(4,38)} = 2.539$, $p = 0.0556$; Fig. 3f). However, there was a trend to increase 5-HT7 levels when comparing CMS with the control group.

Effects of CMS and Antidepressant Treatments on the Expression of the Adrenergic Receptors in the FC

The α 1A expression did not change in our experimental set (one-way ANOVA-Tukey $F_{(4,39)} = 2.003$, $p = 0.1131$; Fig. 4a). The CMS exposure trended to decrease α 1B mRNA and desipramine and duloxetine upregulated it (one-way ANOVA-Tukey $F_{(4,39)} = 10.87$, $p < 0.0001$; Fig. 4b). CMS and CMS+Desipramine levels of α 1D were increased compared with the control group (Kruskal-Wallis test-Dunn, KW statistic = 18.75, $df = 4$, $p = 0.0009$; Fig. 4c).

Desipramine and duloxetine upregulated α 2A mRNA (one-way ANOVA-Tukey $F_{(4,39)} = 21.67$, $p < 0.0001$; Fig. 4d). α 2A also trended to decrease under CMS conditions (two-tailed unpaired t test $t = 3.216$, $df = 25$, $p = 0.0036$). The experimental procedures applied did not alter α 2B mRNA expression in the FC (one-way ANOVA-Tukey $F_{(4,39)} = 1.426$, $p = 0.2436$; Fig. 4e). Only desipramine raised α 2B levels under CMS conditions (one-way ANOVA-Tukey $F_{(4,39)} = 10.78$, $p < 0.0001$; Fig. 4f).

β 1 mRNA expression increased with the duloxetine treatment (one-way ANOVA-Tukey $F_{(4,39)} = 4.713$, $p = 0.0034$; Fig. 4g). No changes detected in the FC for β 2 (one-way ANOVA-Tukey $F_{(4,39)} = 0.3681$, $p = 0.8299$; Fig. 4h) and β 3 (one-way ANOVA-Tukey $F_{(4,39)} = 2.198$, $p = 0.0871$; Fig. 4i), but a clear difference was revealed between CMS and control for β 3.

Escitalopram had no effects on noradrenergic receptors.

Discussion

The inhibition of the monoamine reuptake still epitomizes the first pharmacological approach for mood disorders treatment [23]. This study examines alterations caused by CMS on the monoaminergic neurotransmission in FC of rats. Specifically, it explores possible changes in 5-HT and NE reuptake transporters and the 5-HT and NE signaling receptors expression.

Our results did not expose changes in serotonin and norepinephrine levels after CMS in FC. The method used for the assay refers to the whole levels of neurotransmitters and it does not represent the levels acting on the synaptic cleft. Other studies describe a decline of monoamines in FC after CMS exposure employing longer experimental protocols [24, 25]. Our results may rather reflect an equilibrium among synthesis and degradation of neurotransmitters. Nevertheless, the discussion about the real existence of changes in monoamine levels in MDD continues, and there are conflicting results showing either a decrease [26] or no effects [27] on serotonin and norepinephrine plasma concentrations in patients with MDD. This situation points out to a feasible implication of other neurotransmission elements involved in the antidepressant action of classic pharmacological treatments.

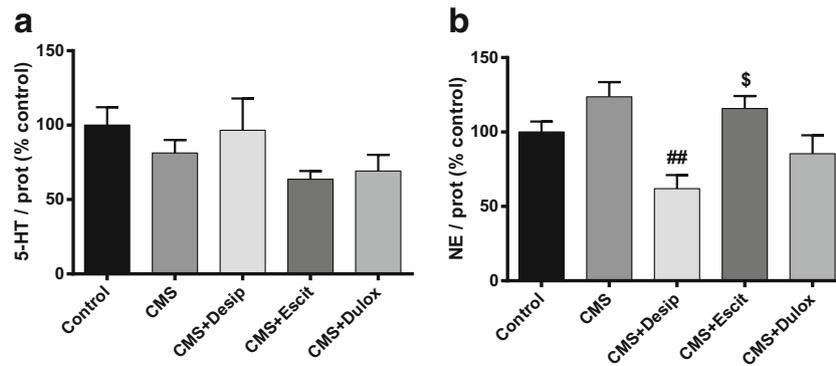


Fig. 1 Effects of CMS and AD treatments on the whole serotonin (5-HT) and norepinephrine (NE) levels in the FC. No changes detected in the 5-HT levels after the CMS or the antidepressant treatments (a). The CMS did not modify the NE amount but desipramine decreased it (b). The

vehicle-injected groups did not differ from the same experimental groups without injection, and their values have been pooled in the figure. Data are means \pm SEM of 6–8 rats per group; ^{##} $p < 0.01$ vs. CMS; ^{\$} $p < 0.01$ vs. CMS+Desip; One-way ANOVA following Tukey post hoc test

The desipramine-induced noradrenergic tone reduction detected could be part of a compensatory mechanism facing the high concentrations reached in the synaptic cleft after desipramine administration. Even more, it would agree with a hypothesis proposing the diminishing in the *locus coeruleus* activity as necessary for the efficacy of an antidepressant drug [28].

We have employed a stress-related protocol, which as inducer of the hypothalamic-pituitary-adrenal (HPA) axis stimulates the release of catecholamines, including norepinephrine (NE). The NET serves as the primary mechanism for the inactivation of noradrenergic signaling [29]. Thus, CMS-induced NET overexpression could be a mechanism aimed to decrease the trend of augmented NE levels compared with

control detected in CMS. Furthermore, antidepressant treatments prevented the CMS-induced NET overexpression agreeing with studies supporting the ability of some ADs to restore the baseline NET expression levels. A possible hypothesis is that neural adaptation resulting in enhanced monoaminergic neurotransmission may be mediating antidepressant actions [30]. In the same vein, the absence of effects of CMS on SERT expression could be explained by the fact that CMS did not modify 5-HT levels, although further research is warranted to support these hypotheses.

Inhibition or desensitization of 5-HT_{1A} receptors exhibits antidepressant effects in the raphe nuclei [31, 32]. The 5-HT_{1A} heteroreceptors in FC can exert an indirect suppression of the serotonin cell firing in the dorsal raphe nucleus [33] and

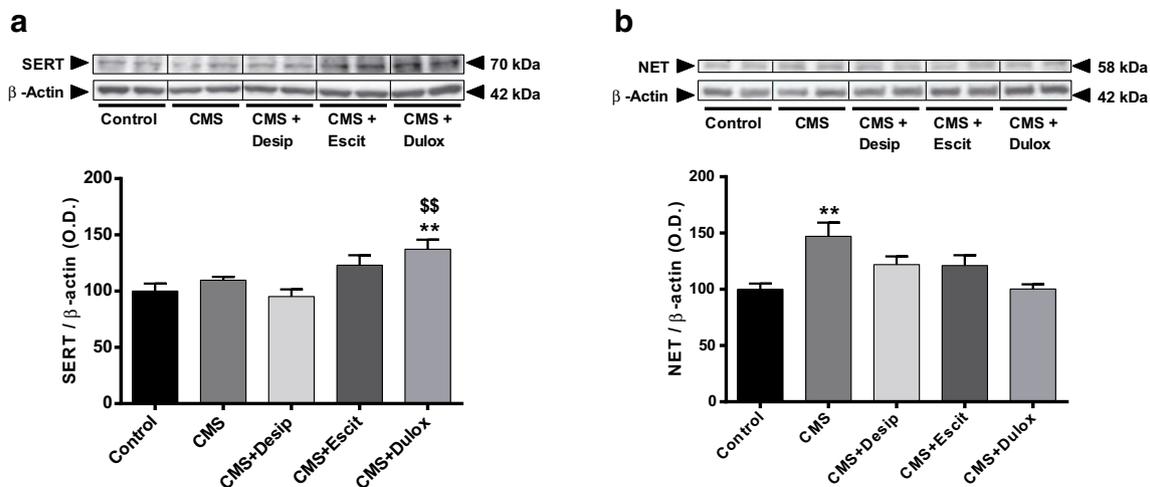
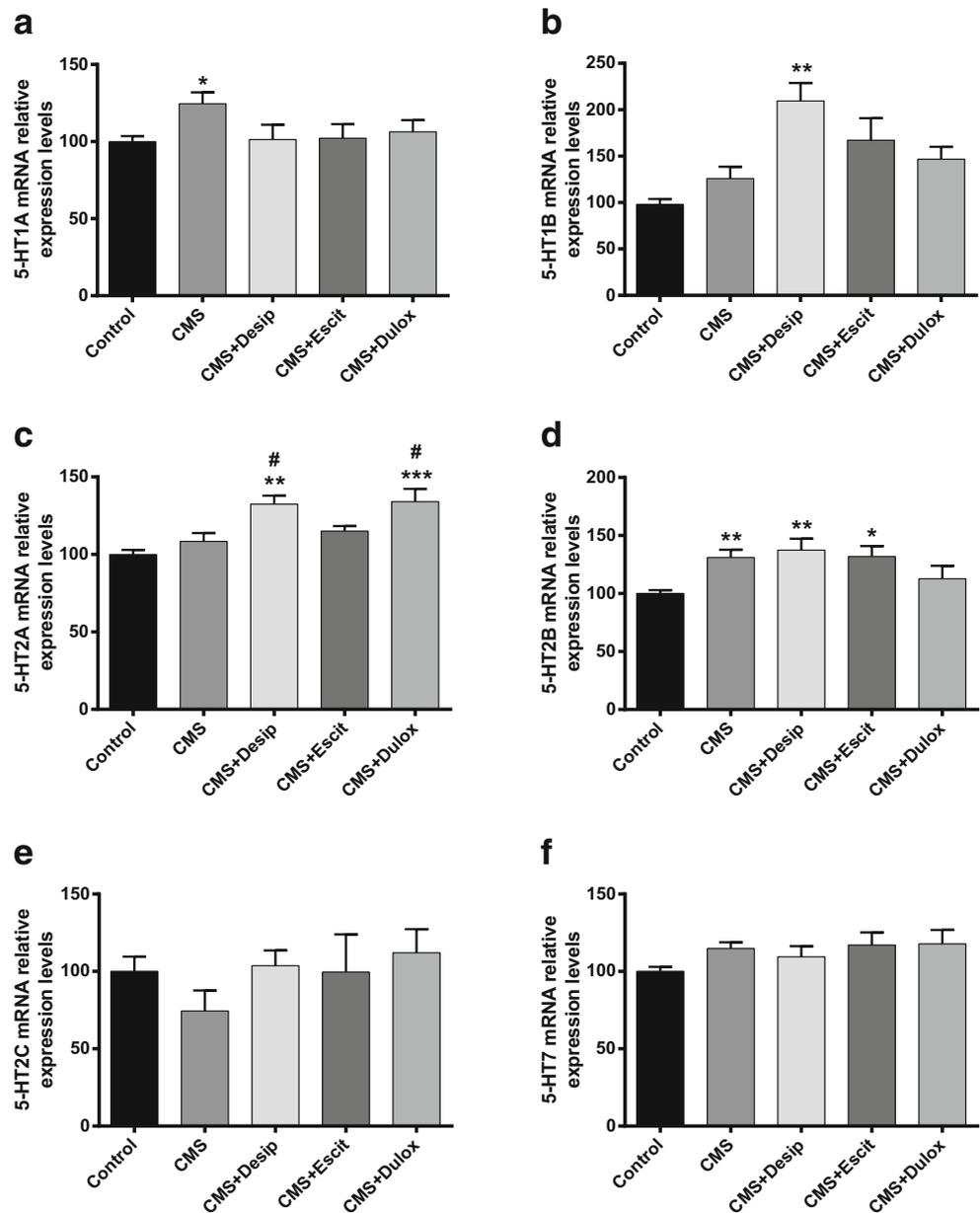


Fig. 2 Effects of CMS and AD treatments on the protein expression of 5-HT and NE transporters in the FC. CMS did not affect the protein expression of the serotonin transporter (SERT) (a) and upregulated the expression of the norepinephrine transporter (NET) (b). Desipramine and escitalopram did not change the expression of the SERT but duloxetine increased it (a). The NE transporter (NET) expression was not affected by ADs (b). The vehicle-injected groups did not differ from the same experimental groups without injection, and their values have been pooled in the

figure. The densitometric data of the band of interest were normalized by beta-actin. Blots were cropped (black lines) for improving the clarity and conciseness of the presentation. Data are means \pm SEM of 6–8 rats per group; ^{**} $p < 0.01$ vs. control; ^{\$\$} $p < 0.01$ vs. CMS+Desip; One-way ANOVA following Tukey post hoc test. The values of NET (b) did not follow a Gaussian distribution; thus, a nonparametric ANOVA with a Kruskal-Wallis test followed by a Dunn's post hoc test was performed

Fig. 3 Effects of CMS and AD treatments on the expression of the 5-HT receptors 5-HT1A, 5-HT1B, 5-HT2A, 5-HT2B, 5-HT2C, and 5-HT7 in the FC. CMS increased the mRNA expression levels of 5-HT1A (a), 5-HT2B (d), and 5-HT7 (f) ($p = 0.0053$; two-tailed t test) when compared with the control group. CMS had no effect on mRNA levels of 5-HT1B (b), 5-HT2A (c), and 5-HT2C (e). Desipramine treatment upregulated 5-HT1B mRNA levels compared with the control group (b). Desipramine and Duloxetine upregulated 5-HT2A mRNA levels compared with the control and CMS groups (c) but none of the ADs could affect the induction of the expression of 5-HT2B caused by the CMS protocol although duloxetine made its levels similar to control (d). No effects recorded for the ADs over 5-HT2C (e). Treatments did not have any effect on 5-HT7 mRNA levels (f). The vehicle-injected groups did not differ from the same experimental groups without injection, and their values have been pooled in the figure. Data are means \pm SEM of 6–8 rats per group; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. control; # $p < 0.05$, vs. CMS; One-way ANOVA following Tukey post hoc test



the selective activation of postsynaptic 5-HT1A receptors plays an important role in the antidepressant response [34]. Even more, chronic fluoxetine treatment decreases the 5-HT1A binding in prefrontal cortex and hippocampus of rats [35]. Our study shows that CMS upregulates 5-HT1A, and treatments block that upregulation, further supporting the idea of antidepressant effects mediated by 5-HT1A inhibition.

The absence of 5-HT1B expression changes after CMS seems coherent with other study demonstrating that 5-HT1B remains unaltered in the anterior cingulate cortex of patients with MDD [36]. In any case, the 5-HT1B auto and heteroreceptors populations are hard to study separately due to their overlapping diffuse localization in FC; thus, 5-HT1B receptor agonists and antagonists can exhibit antidepressant

properties. The use of experimental strategies that allow the study of them separately represents an important issue for future research to reach reliable findings about the 5-HT1B and its potential as a therapeutic tool [37].

Postsynaptic 5-HT2A and 5-HT2B receptors seem to be necessary for the action of ADs [38] and their upregulation empowers the neurotransmission. Thus, data observed in the CMS could form part of a compensatory mechanism triggered to maintain a minimum of serotonin working in the synaptic cleft. The effect of ADs upregulating this kind of 5-HT receptors seems to be consistent with other studies indicating that they are essential for the effects of treatments [38].

The potential of the 5-HT2C receptor is being explored in psychiatric diseases [39] and an inverse agonism/antagonism

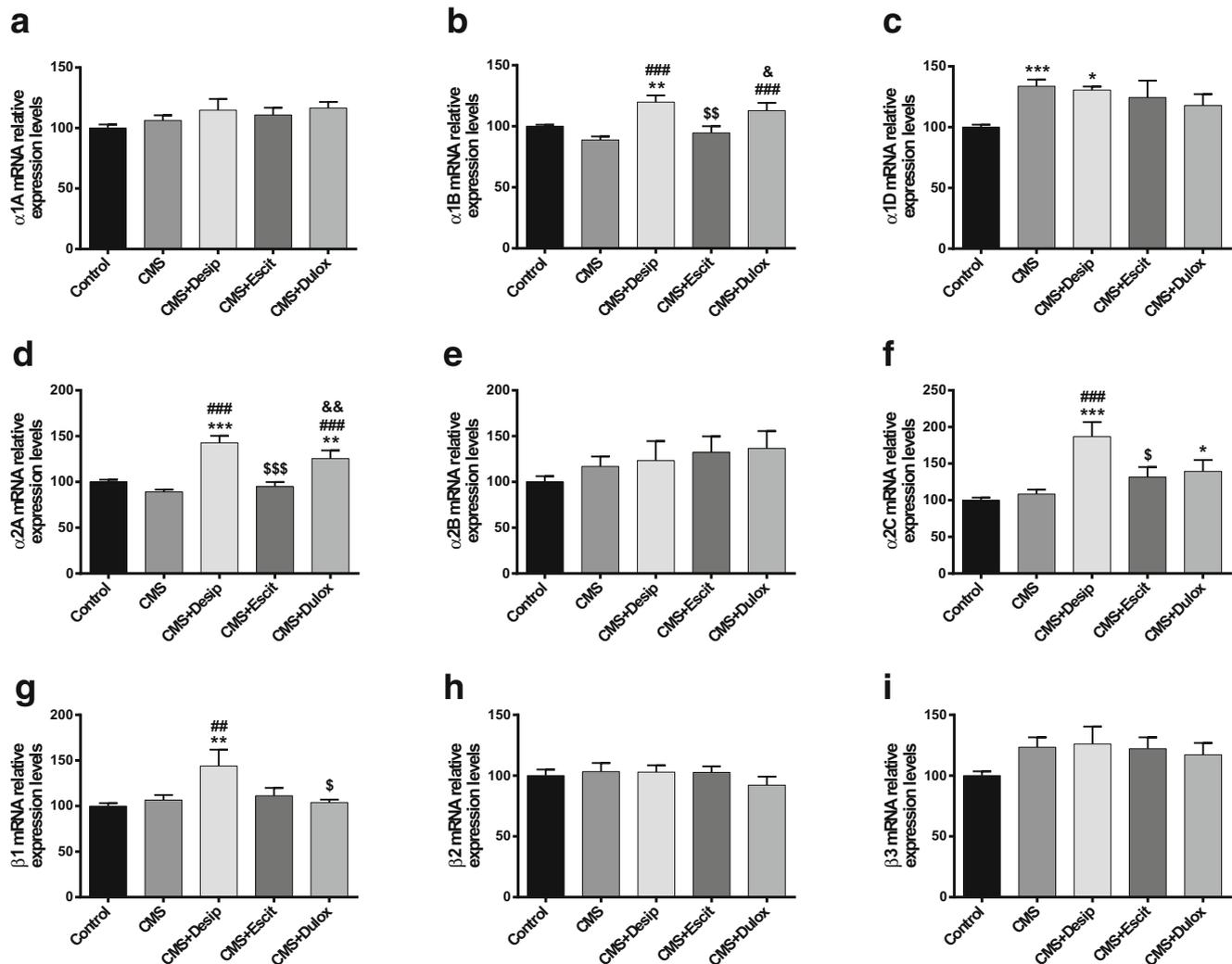


Fig. 4 Effects of CMS and AD treatments on the expression of NE receptors $\alpha 1A$, $\alpha 1B$, $\alpha 1D$, $\alpha 2A$, $\alpha 2B$, $\alpha 2C$, $\beta 1$, $\beta 2$, and $\beta 3$ in the FC. The CMS increased $\alpha 1D$ (c) and tended to diminish $\alpha 1B$ (b) and $\alpha 2A$ (d) (two-tailed unpaired *t* test, $p = 0.0012$ and $p = 0.0036$, respectively) and to upregulate $\beta 3$ (i) (two-tailed unpaired *t* test, $p = 0.0121$) mRNA expression. Desipramine and duloxetine increased $\alpha 1B$ (b) and $\alpha 2A$ (d) and desipramine also increased $\alpha 2C$ (f) and $\beta 1$ (g) levels compared with the CMS group. No effects on any of the noradrenergic receptors were detected after the escitalopram treatment. The vehicle-injected groups did

not differ from the same experimental groups without injection, and their values have been pooled in the figure. Data are means \pm SEM of 6–8 rats per group; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. control; ## $p < 0.01$, ### $p < 0.001$ vs. CMS; \$ $p < 0.05$, \$\$ $p < 0.01$ vs. CMS+Desip; & $p < 0.05$, && $p < 0.01$ vs. CMS+Escit. One-way ANOVA following Tukey post hoc test. The values of $\alpha 1D$ (c) did not follow a Gaussian distribution; thus, a nonparametric ANOVA with a Kruskal-Wallis test followed by a Dunn's post hoc test was performed

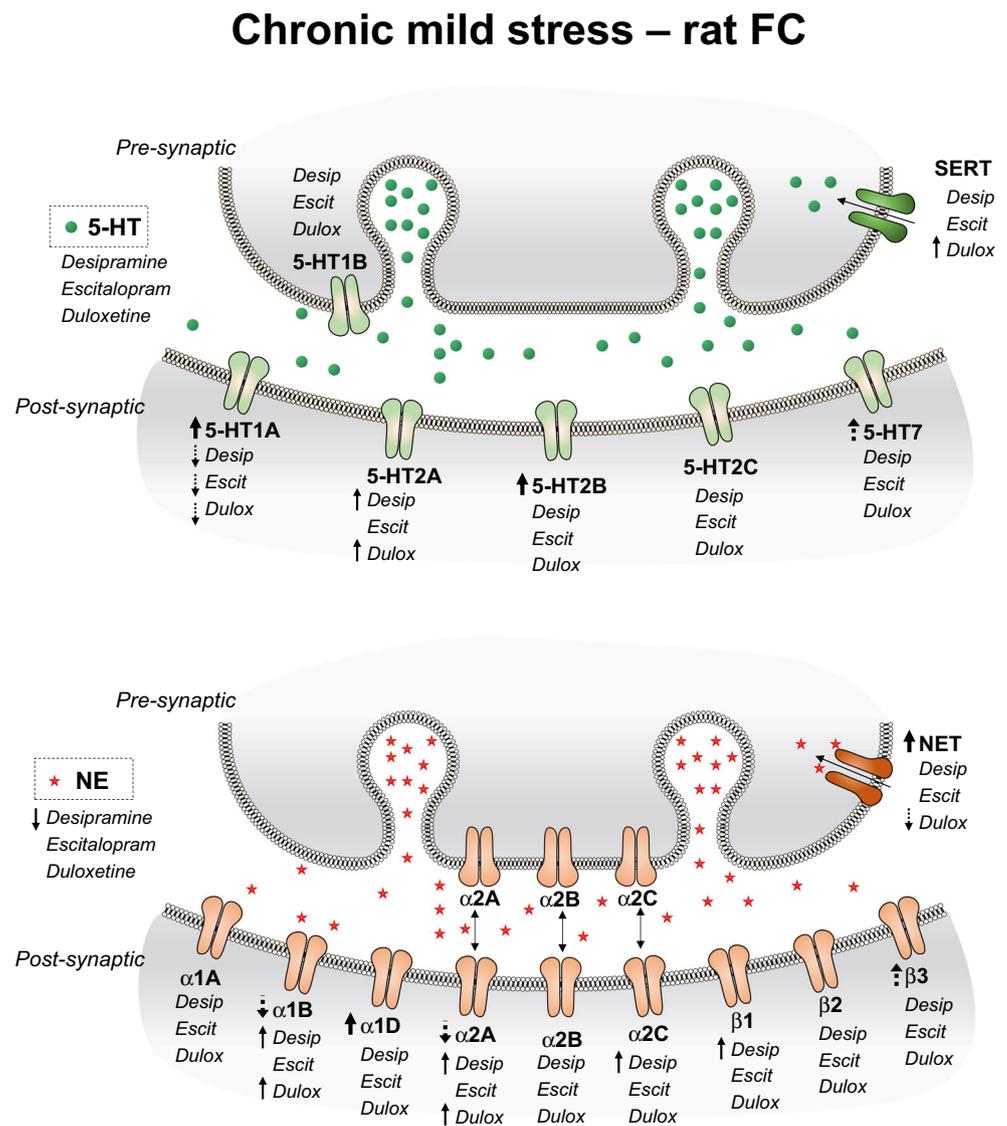
has shown antidepressant properties [40]. However, 5-HT_{2C} mRNA suffers from extensive RNA editing processes and these post-transcriptional modifications are critical for its function [41], making mRNA expression analysis maybe insufficient to explain 5-HT_{2C} implication in CMS. In addition, 5-HT_{2C} agonism, partial agonism, or antagonism are still undergoing testing to assess their antidepressant ability [42].

Latest observations in rodent models and patients suggest that 5-HT₇ blockade mediates antidepressant efficacy [43]. Thus, taking also into account that 5-HT₇ is upregulated after CMS, future research should focus on the potential as therapeutic targets of this family of receptors for the treatment of mood disorders.

Escitalopram did not affect noradrenergic receptors, which is coherent with its specificity for serotonin since escitalopram is a SSRI. Nevertheless, we found a possible effect for escitalopram in NET expression, indicating the significance of the crosstalk among monoamine neurotransmission systems and their relevance to depressive-like behaviors [44].

Alpha (α) receptors have been related to antidepressant responses [45]. CMS exposure decreased $\alpha 1B$ and increased $\alpha 1D$ adrenoceptors, whereas desipramine and duloxetine increased $\alpha 1B$ expression. These results are consistent with studies showing that activation of the $\alpha 1B$ -adrenoceptors associated with antagonism of $\alpha 1D$ -adrenoceptors might be one of the mechanisms involved in the antidepressant-like activity

Fig. 5 CMS effects on the monoaminergic neurotransmissions and their modulation by ADs. Schematic representation of 5-HT and NE synapses including the neurotransmitters, their receptors (5-HT1A, 5-HT1B, 5-HT2A, 5-HT2B, 5-HT2C, and 5-HT7 for serotonin; α 1A, α 1B, α 1D, α 2A, α 2B, α 2C, β 1, β 2, and β 3 for noradrenaline), and their reuptake transporters (SERT and NET) and their up or downregulation with Desipramine (Desip), Escitalopram (Escit), and Duloxetine (Dulox) under CMS conditions



of imipramine and other tricyclic antidepressants in rodents [46]. Nevertheless, the lack of deep research involving α 1 subtypes brings to light the need of further research to elucidate their contribution to the pathophysiology of depression.

The α 2-adrenoceptors can function as autoreceptors regulating NE release or heteroreceptors modulating other neurotransmitters, including serotonin [47]. Our results showed a decrease in α 2A after CMS, agreeing with reports showing that α 2A knockout mice develop depressive-like phenotypes [48]. Moreover, desipramine induced the expression of α 2A and α 2C, and duloxetine increased the expression of α 2A subtype. These results are consistent with the literature suggesting that α 2 receptors are essential for the antidepressant activity [48, 49] and α 2A specifically for the efficacy of desipramine through its direct interaction with the receptor [50]. These results are also consistent with a previous study wherein an α 2-adrenoceptor antagonist blocked the antidepressant

effect of acutely administered desipramine [51]. The absence of changes in α 2B subtype could be explained due to its low expression profile in FC [52]. Actually, it seems like regulation of receptors in depressive disorders is likely to vary considerably among different brain regions and even highly localized subregions [53].

The overlap between the auto and heteroreceptor population hinders the interpretation of mRNA α 2 expression results. On one hand, α 2A receptors are increased in FC of suicide depressive victims [53, 54]. Moreover, mirtazapine and mianserin are α 2 antagonists with widely proved antidepressant properties, although it cannot be ignored that they exert actions on other receptors [3]. Alternatively, the action upon postsynaptic α 2 is critical for antidepressant effects [51]; it could improve the working memory [55] and it promotes brain plasticity [56]. Bearing in mind the opposite actions described, discerning the pre or postsynaptic nature of the

changes on $\alpha 2$ receptors represents an essential issue for future research.

There are post mortem studies showing either increased, decreased, or unaffected levels of β -receptors in subjects with depression [57, 58]. Furthermore, it has been described that chronic signaling through $\beta 3$ -adrenergic receptors during social stress is an adaptive response that improves behavioral function and that $\beta 3$ signaling may promote emotional coping with stress [59]. Our research has found upregulated $\beta 3$ expression after CMS; thus, a plausible explanation could be based on the previously mentioned adaptive response to improve behavioral performance during stress. Moreover, antidepressant-treated groups show values close to the CMS increase, reflecting the positive effects of the $\beta 3$ stimulation.

Agonism of $\beta 1$ and $\beta 2$ receptors (e.g., by intraventricular administration of isoproterenol) produces behavioral changes similar to those observed after administration of proven antidepressant drugs [60]. Thus, desipramine induction of $\beta 1$ receptor could explain part of the beneficial effects of this antidepressant drug. This outcome seems contrary to the decrease in $\beta 1$ using an 8-week CMS reported in a previous study [61]. The length of the stress exposure and the multiple adjustments among the different subtypes of noradrenergic receptors could explain this downregulation. In any case, the increase described after imipramine treatment agrees with the desipramine results in our experiment. What is more, polymorphisms for $\beta 1$ receptor have been associated with antidepressant outcomes [62].

For a better interpretation, and a summary, of the alterations caused by CMS and the actions of antidepressants over the serotonergic and noradrenergic neurotransmission in FC, a diagram with their possible synaptic implications is included in Fig. 5.

Conclusions

We have identified imbalances in reuptake transporters and receptors involved in the monoaminergic neurotransmission pointing towards the weakening of their signaling. Our results also show that antidepressant treatments strive for the improvement of the monoaminergic tone, being desipramine and duloxetine more influential than escitalopram over noradrenergic elements.

To our knowledge, our study is the first one tackling a whole characterization of the serotonergic and noradrenergic receptors and reuptake transporters in the most used and validated depression model—the CMS. Moreover, the simultaneous analysis of three antidepressants with different mechanisms of action adds strength to this research allowing their comparison and providing new insights into their molecular effects.

In summary, there are molecular alterations on serotonergic and noradrenergic neurotransmissions in FC induced by CMS exposure. Yet more, antidepressant treatments modulate the elements of these neurotransmission systems differentially.

There is a need of more accurate and effective drugs to treat depression with lower side effects. Our research offers new insights in this direction providing some advices for possible multi-target drug designs.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

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