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## Testosterone is related to GABA+ levels in the posterior-cingulate in unmedicated depressed women during reproductive life

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### Abstract

**Background:** The role of testosterone (T) in the pathophysiology of affective disorders and anxiety is broadly supported. Evidence suggests that T has anxiolytic and antidepressant properties. One proposed route for the central effects of T is its interaction with the gamma-aminobutyric acid (GABA) system. We explored the relationship between T levels and GABA+ levels in anterior-cingulate (ACC) and the posterior-cingulate (PCC) regions in depressed women, using magnetic resonance spectroscopy (<sup>1</sup>H-MRS).

**Methods:** Twenty-one depressed patients with regularly cycling who were not taking hormonal or psychotropic drugs were recruited. We assessed severity of depression using the Hamilton Depression Rating Scale (HDRS). Blood samples were taken for quantification of free (FT) and

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Author contribution

Study concept and design: MFR, SA, RAEE.

Acquisition of data: MLT, ECC, NBS.

Analysis and interpretation of data: MFR, SA, RAEE.

Drafting of the manuscript: MFR, SA, RAEE.

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Disclosures

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total testosterone (TT) on the day of the magnetic resonance (MR) scan. We evaluated GABA+ levels in the PCC and ACC, using the Hadamard Encoding and Reconstruction of MEGA-Edited Spectroscopy (HERMES) sequence. Pearson correlations were used to evaluate the association between FT, TT, GABA+ concentrations, and HDRS scores.

**Results:** TT and FT levels were positively correlated with GABA+ levels in the PCC. No correlation was observed between T levels and GABA+ levels in the ACC. The HDRS total scores correlated negatively with FT levels.

**Limitations:** Limitations include the cross-sectional evaluation and the lack of a comparative healthy group.

**Conclusions:** Our findings suggest that the potential anxiolytic and antidepressant properties of T are related to increased GABA+ levels in the PCC. This observation may contribute to increased understanding of the role of T in depressive and anxiety symptoms in women.

### Keywords

Testosterone; Major depressive disorder; Women; GABA; MR spectroscopy

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## 1. Introduction

Major depressive disorder (MDD) is a chronic illness that is approximately twice as common in women as in men. The complexity of this disorder notwithstanding, biological factors are suggested as triggers of a depressive episode in women (Albert, 2015). Similarly, anxiety is highly prevalent in women (Remes et al., 2016), and frequently MDD and anxiety are comorbid, suggesting common pathophysiological elements (Wichers et al., 2009). Gender differences in the presentation of MDD and anxiety disorders have been attributed to hormonal fluctuations, as is observed in studies that explore depressive and anxiety symptoms during periods of hormonal fluctuations, such as puberty, postpartum and perimenopause (Dahl and Gunnar, 2009; Douma et al., 2005; Halbreich, 2010; Flores-Ramos et al., 2014, 2017). A dysregulation in the stress system, which differs between sexes (Fernandez-Guasti et al., 2012; Oyola and Handa, 2017), is a proposed pathway to understand the hormonal influence on mood and anxiety, since it has been observed that differences in the cortisol response to specific stressors depend on sex and circulating hormones (Hollanders et al., 2017; Stephens et al., 2016). The hypothalamic-pituitary-adrenal (HPA) axis plays an important role in the regulation of stress (and its dysregulation in affective disorders) (Herman et al., 2005), which in turn is regulated by neurosteroids, such as testosterone and its metabolites, via GABAergic receptors (Gunn et al., 2015). Animal studies show that cerebrocortical neurosteroid concentrations increase after 1 h of restraint stress exposition (Mele et al., 2004); similarly foot shock stress elicits an increase of neuroactive steroids, which is blocked by abecarnil (a GABA<sub>A</sub> receptor agonist), suggesting a relationship between brain neuroactive steroids and GABA<sub>A</sub> receptor function (Barbaccia et al., 1996). Modulation of GABA receptors by neurosteroids has been proposed as a “fine tuning” HPA axis function, or in other words, a mechanism to return to homeostasis after an acute stressor (Gunn et al., 2015). In humans, some imaging studies support the role of neuro-steroids in stress regulation; for instance, a resting-state study

showed a modulation of amygdala connectivity by neurosteroids (allopregnano-lone and DHEA) (Sripada et al., 2014).

Interaction between the T and GABA systems has been supported by preclinical studies that demonstrate that anxiolytic effects of T are mediated by GABA-A receptors (Fernandez-Guasti and Martinez-Mota, 2005), with a minimal participation of estradiol aromatization (Gutiérrez-García et al., 2009). However, clinical studies of T participation in affective/anxiety symptoms have reported contradictory results that may depend on sex. In men, lower levels of bioavailable and total testosterone were observed in depressed patients compared to healthy males (McIntyre et al., 2006); furthermore beneficial effects of T supplementation have been demonstrated, improving mood and alleviating anxiety (Pope et al., 2000, 2010). In the case of women, data about the relation of T and depression are scarce and paradoxical results have been obtained. One study, based on a community sample, showed lower salivary testosterone levels in females with a current major depressive disorder and other anxiety disorders compared to healthy controls (Giltay et al., 2012).

Reproductive life stage could influence the relationship between testosterone and mood in the menopausal transition, an increased risk of depressive symptoms was related to higher levels of T in Caucasian, but not in African-American women (Milman et al., 2015). Similarly, a follow-up study observed that an increase in T levels was associated to an increased risk of depressive symptoms (Bromberger et al., 2010). Paradoxically, low-dose T supplementation in women seems to improve mood and reduce anxiety in women with treatment-resistant-depression, when used as a complementary therapy (Miller et al., 2009). Testosterone effects on mood also seem to be dose-dependent, with high T levels associated to depression (Rohr, 2002).

Magnetic resonance spectroscopy (MRS), which can measure levels of metabolites in the central nervous system (CNS) (Harris et al., 2017), has been applied in a number of studies of depression, hormones and GABAergic systems. Studies have demonstrated that GABA concentrations are diminished in the occipital cortex (OCC) in depressed patients compared to healthy controls (Sanacora et al., 1999), as well as in the prefrontal cortex (PFC) areas (Hasler et al., 2007). Moreover, in patients that have received an adequate treatment, such as serotonin selective reuptake Inhibitors (SSRI) or electroconvulsive therapy (ECT), normalization of GABA concentrations in the OCC is observed (Sanacora et al., 2002, 2003). Changes in GABA levels across the menstrual cycle have been reported, as well as an influence of the use of hormonal contraceptives (De Bondt et al., 2015), suggesting MRS is sensitive to the impact of hormonal fluctuations on the GABAergic system. To our knowledge, MRS studies have not evaluated the relationship between testosterone levels, mood and GABA levels. Therefore, this work explores testosterone levels in depressed women and their relationship to ACC and PCC GABA levels, based on the hypothesis that a positive relationship will be observed between free and total testosterone and GABA levels in the explored brain regions.

## 2. Methods

### 2.1. Participants

Twenty-one female patients with major depressive disorder (MDD), treatment-free and with regular menstrual cycling, participated in this study. Participants were recruited in the outpatient clinic of the National Institute of Psychiatry Ramón de la Fuente Muñiz (INPRFM), in Mexico City. The protocol fulfilled the requirements of the institutional ethical committee and all the participants signed an informed consent.

The MINI International Neuropsychiatric Interview (MINI) (Sheehan et al., 1998) was applied to evaluate psychiatric disorders. The inclusion criteria were: women; MDD diagnosis; 18 to 45 years old; regular menstrual cycles; reproductive period according to the Stages of Reproductive Age Workshop (STRAW) (Harlow et al., 2012). Patients with psychiatric comorbidities, except anxiety disorders, were excluded; medical illnesses were also not allowed according to medical examination and laboratory tests. Women using any hormonal-based compound (contraceptives or hormone therapy replacement), anti-epileptic drugs, and/or psychotropic medication were excluded; patients with alcohol use in the last 2 weeks or drug use in the last 6 months were also excluded. Lastly, patients treated with electro-convulsive therapy and transcranial magnetic stimulation in the last year were not included in the present study.

### 2.2. Procedures

MDD was diagnosed by a trained psychiatrist, according to the Diagnostic and Statistical Manual of Mental Disorders, fourth edition (DSM-IV) criteria (APATFo, 1991). The MINI was applied to confirm the diagnosis and to evaluate comorbidities. A complete clinical evaluation was performed for all the participants, including a menstrual cycle history to select only patients on the -5 to -3 reproductive stages according to STRAW, which means reproductive age and includes women with regular menstrual cycle, and does not include women in the menopausal transition (Soules et al., 2001). We recorded the last day of menstrual period and classified it in follicular or luteal menstrual phases. Severity of depression symptoms was evaluated by the 17-item Hamilton depression rating scale (HDRS-17) (Hamilton, 1960).

### 2.3. Laboratory test

Blood samples were taken between 7:00 and 9:00 AM. Quantification of hormones (progesterone, estradiol, free and total testosterone) was performed by chemoluminescence. Additionally thyroid tests were conducted in order to exclude thyroid abnormalities in the participants.

### 2.4. Magnetic resonance imaging (MRI) study protocol

All the MRI experiments were performed with a 3.0 Tesla scanner (Phillips Medical Systems) equipped with a 32-channel phased-array head coil. Before the MRS scans, a T1-weighted anatomical MRI was acquired for localization purposes with repetition time (TR) = 7.0 ms, and echo time (TE) = 3.5 ms.

MR Spectroscopy data were acquired from voxels centered on the anterior cingulate cortex ( $3 \times 3 \times 3 \text{ cm}^3$ ) and the posterior cingulate cortex ( $3 \times 3 \times 3 \text{ cm}^3$ ), avoiding the lateral ventricle and skull, as is shown in the Fig. 1. Signals of GABA and glutathione (GSH) were detected using Hadamard Encoding and Reconstruction of Mega-Edited Spectroscopy (HERMES) (Saleh et al., 2016). Briefly, HERMES allows the simultaneous J-difference editing of GSH and GABA signals by applying editing pulses to GABA spins (at 1.9 ppm) and GSH spins (at 4.56 ppm) in a Hadamard-encoded editing scheme. Four sub-experiments are performed and the GABA and GSH-edited difference spectrums are calculated. The HERMES sequence was acquired with the following parameters: TR = 2000 ms; TE = 80 ms; 320 signal averages (80 averages for each sub-experiment; 2 kHz spectral width; 2k data-points). Data were processed using Gannet, the GABA Analysis Toolkit, which allows the evaluation of the quality of data, removing rater-dependent variance, preventing the significance-based selection of processing parameters and widening access to the non-expert user (Edden et al., 2014). Post-processing frequency and phase correction was applied according to Mikkelsen et al. (2017). GABA values were corrected to account for differential gray matter/white matter/cerebrospinal fluid composition of the voxel (Porges et al., 2017; Harris et al., 2015). GABA levels were quantified relative to the unsuppressed water signal from the same region, corrected for the CSF volume-fraction  $f_{\text{CSF}}$  by dividing by  $(1 - f_{\text{CSF}})$ .

Additionally, since the edited signal at 3 ppm includes contributions from GABA, macromolecules and homocarnosine (Rothman et al., 1993), we refer to the signal obtained as GABA+.

## 2.5. Statistical analysis

Descriptive statistical analysis was carried out, and a two-tailed *t*-test was applied in order to evaluate differences in GABA+ levels between menstrual cycle phases. Associations between HDRS scores and GABA+ concentrations as well as FT and TT levels were assessed using Pearson correlation coefficient. Level of significance was set at  $p < 0.05$ . A Pearson correlation analysis was also carried out with other variables that could influence the results (progesterone, estradiol, number of previous episodes and age at first episode). Lastly, a partial correlation analysis was carried out to analyze the relationship between PCC\_GABA+ levels, ACC\_GABA+ levels, free testosterone and total testosterone, controlling by HDRS scores. False discovery rate (FDR) (Benjamini and Yekutieli, 2001) method was used to correct the correlations between the main variables according to our hypothesis (FT, TT, PCC\_GABA+ and ACC\_GABA+) before and after controlling by HDRS. All statistical analyses were carried using the Statistical Package for Social Sciences Software 20th version (SPSS-20).

## 3. Results

Mean age of the patients was 31.2 (sd 8.18); 12 patients were assessed to be in a follicular phase and 9 in the luteal phase. The clinical and socio-demographic characteristics of the patients are shown in the Table 1. GABA+ levels were normally distributed and are presented as means (sd), as well as testosterone levels. GABA+-edited spectra were

successfully acquired in all subjects, as shown in Fig. 2. No differences in the GABA+ concentrations in the PCC nor ACC regions were observed between menstrual phases ( $p = 0.83$ ;  $p = 0.55$ ).

Concentration of GABA+ in the PCC showed significant positive correlations with free ( $r = 0.53$ ,  $p = 0.039$ ) (Fig. 3B) and total ( $r = 0.64$ ,  $p = 0.009$ ) testosterone levels (Fig. 3A). Estradiol and progesterone levels were not correlated with brain metabolites in the regions of interest. Previous depressive episodes were observed in patients in a range from 0 to 12; we did not observe a relationship between the number of previous depressive episodes and GABA+ levels in the regions of interest. Age at first episode was also not related to GABA+ levels (Table 2a). According to our hypothesis we analyzed the correlation between TT, FT, PCC-GABA+ and ACC-GABA+, and after a correction with the FDR method the only correlation that remained significant was TT and PCC-GABA+. Lastly, HDRS total scores showed a negative relationship to FT ( $r = -0.50$ ,  $p = 0.02$ ) (Fig. 3C), but no relationship to TT. The partial correlation analysis between TT, FT, PCC GABA+ and ACC GABA+ controlling by HDRS showed that correlation between PCC GABA+ and FT, and correlation between PCC GABA+ and TT were significant ( $r = 0.67$ ,  $p = 0.032$ ;  $r = 0.813$ ,  $p = 0.004$ , respectively) (see Table 2b). After correction with the FDR method, only the correlation between PCC GABA+ and TT remained significant.

#### 4. Discussion

GABA is the main inhibitory neurotransmitter in the brain with an important role in maintaining excitation/inhibition balance, which is implicated in the pathophysiology of many neurological and psychiatric disorders. Reduced GABA neurotransmission has been observed in patients with MDD, underlining the importance of GABAergic systems in mood and anxiety symptoms. Biochemical measures of GABA in plasma and cerebrospinal fluid demonstrated decreased GABA levels in depressed patients (Luscher et al., 2011; Sanacora and Saricicek, 2007). Additionally, evidence coming from cortical inhibition, neuroimaging and neurophysiological studies converge to implicate GABAergic deficits in the pathophysiology of MDD (Croarkin et al., 2011). MRS studies have shown reduced GABA concentrations mainly in the occipital cortex (OCC, frequently studied for methodological reasons) and the ACC (Pehrson and Sanchez, 2015) as well normalization of GABA levels with appropriate treatment (Sanacora et al., 2002). The lack of a healthy control group in the current study design does not allow us to report group differences in GABA levels in our patients. We did not observe a relation between GABA+ concentrations and HDRS scores, in line with other studies using the same instrument to evaluate depressive symptoms (Wang et al., 2016) or using the Montgomery-Asberg Depression rating scale (Hasler et al., 2007). Other studies suggest that GABA changes are related to the diagnosis of depression, but not to the intensity of depressive symptoms. However, in the present sample of depressed women GABA+ levels showed no correlation to intensity of depressive symptoms measured by HDRS.

The main finding of our study was a correlation between GABA+ concentrations in the PCC and FT and TT serum levels. After correction for multiple comparisons and after controlling for HDRS, the relation between TT and GABA+ concentrations in the PCC remained



Lastly, we evaluated the number of previous depressive episodes in our patients, in order to correlate it with GABAergic concentrations, without finding any significant relation between those variables. A history of depression seems to be associated to low GABAergic concentrations in brain, which in turn, is associated with an altered neurosteroid profile (Girdler et al., 2012).

Limitations of the present study should be mentioned: first, we did not compare the brain metabolites concentration and testosterone levels with a healthy control group. The comparison between groups could add some information regarding the GABA/testosterone interaction in non-depressed women. Secondly, the cross-sectional evaluation in our patients represents a limitation because hormonal systems are dynamic and differences in the effect of testosterone on mood have been observed when cross-sectional or longitudinal evaluations are done (Kische et al., 2018). Third, a complete evaluation of hormone profile including precursors and metabolites of testosterone could add information about the interrelation between neurosteroids and GABA concentrations. Lastly, the sample size of the present study is small and our results should be taken cautiously, and apply only for women in reproductive age with depressive disorder.

Clinical implications could derive from our results, including the potential role of neurosteroids in alleviating depressive and anxiety symptoms in women, as well as the consideration of GABAergic path-ways in the depressive symptoms associated with hormonal profile in women (including mood changes associated to perimenopause and premenstrual dysphoric disorder). Lastly, considering that new rapid-acting antidepressants act on the GABAergic and glutamatergic systems, neurosteroid participation in the response to antidepressants should be explored.

Future studies about this topic should be conducted with a larger sample size, including control subjects and considering a longitudinal evaluation to better understand T effects on GABAergic systems; additionally, the measurement of T metabolites (such as dehydrotestosterone) could deliver insight on the effect of testosterone, its metabolites and GABAergic systems.

## 5. Conclusions

In women with major depression, Testosterone effects could be related to increased GABA+ levels in the PCC. This observation may contribute, at least in part, to an improved understanding of the role of T in depressive and anxiety symptoms in women.

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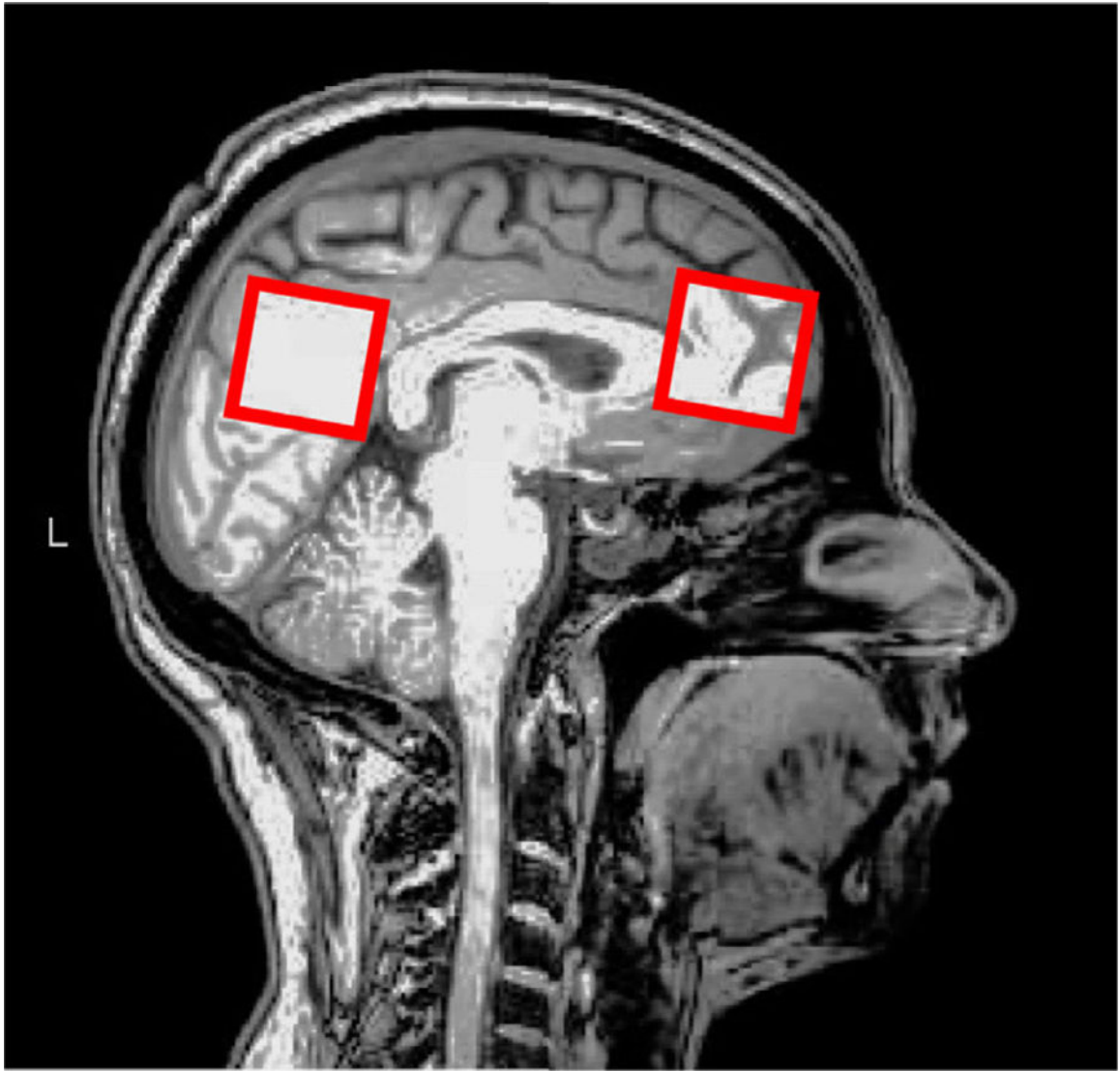
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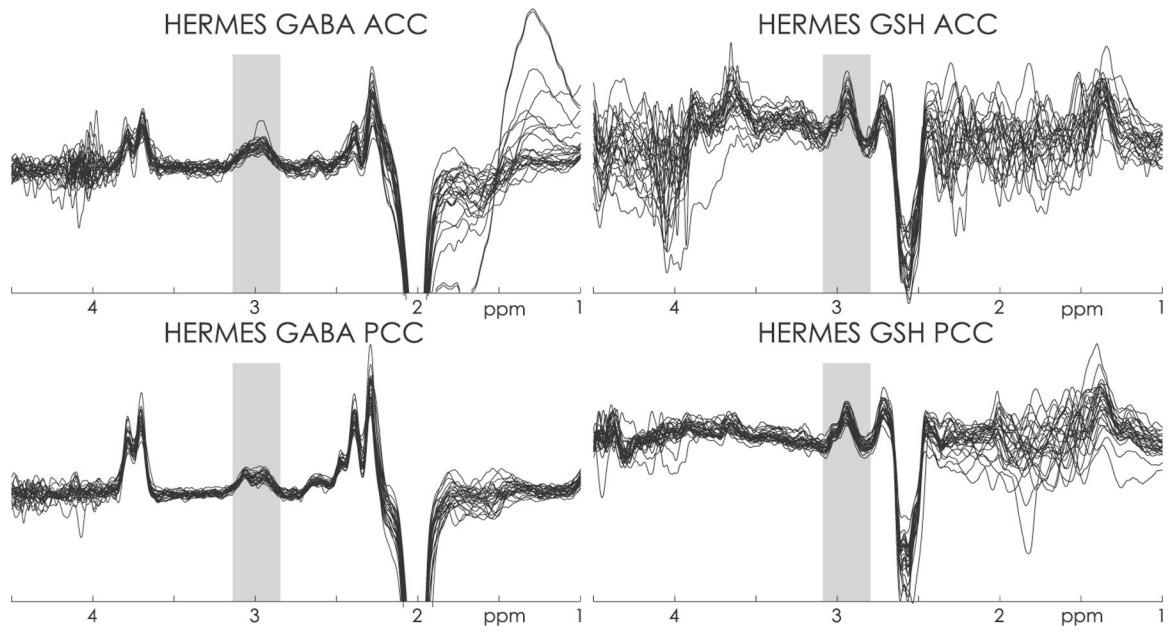
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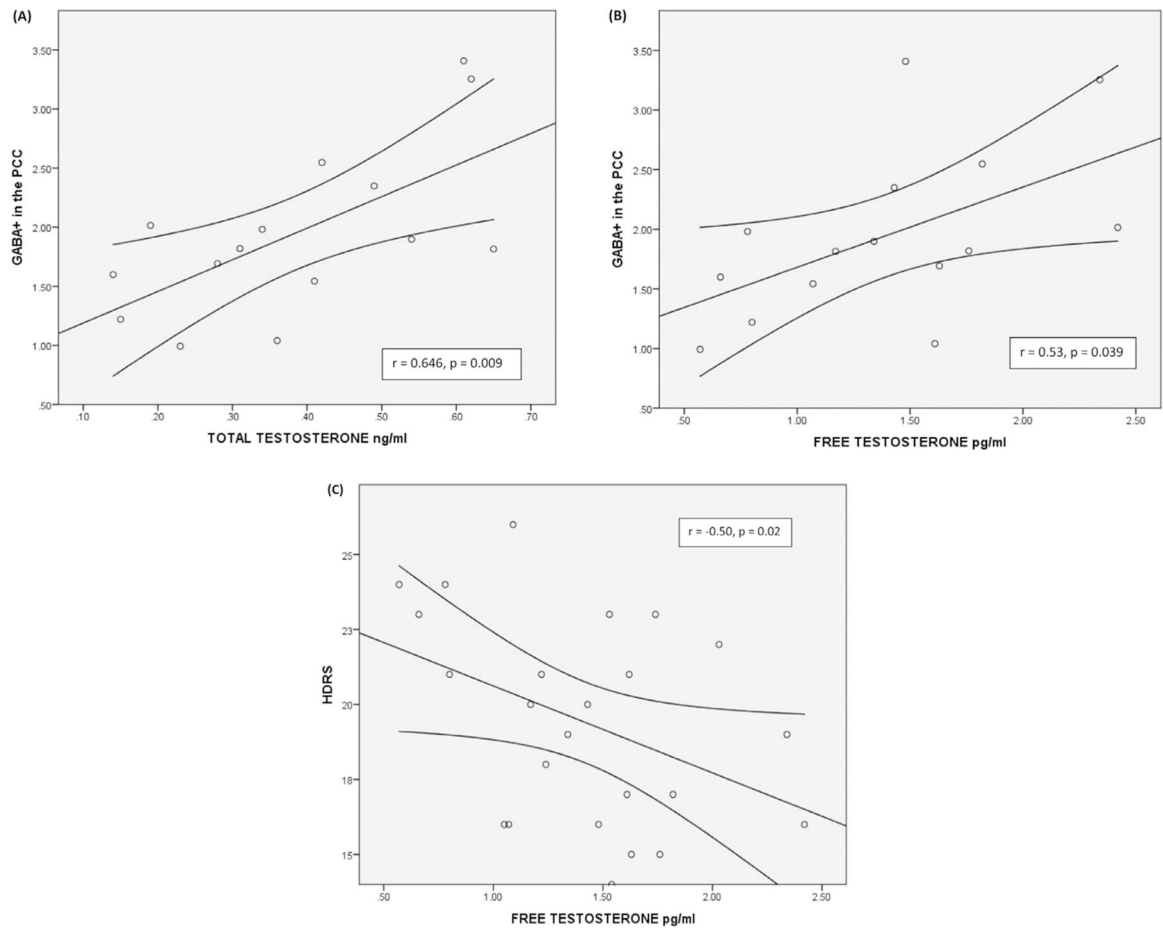
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**Fig. 1.**  
Location of MRS voxels in the anterior (A) and posterior (B) cingulate regions.



**Fig. 2.** GABA+ edited difference spectrum from one subject. The GABA+ signal being quantified is marked at 3 ppm.



**Fig. 3.** (A) GABA+ levels in PCC are correlated with total testosterone ( $r = 0.64$ ;  $p = 0.009$ ). (B) GABA+ levels in PCC are correlated with free testosterone ( $r = 0.53$ ;  $p = 0.039$ ). (C) HDRS total scores showed a negative relationship to FT ( $r = -0.50$ ,  $p = 0.02$ ).

**Table 1**

Clinical and socio-demographic characteristics of participants.

<b>Sociodemographic</b>	<b><i>n</i> = 21</b>
Age (years)	31.48 ± 9.41
Education (years)	13.43 ± 4.11
<b>Marital status, <i>n</i> (%)</b>	
Single	12 (57.1)
Married	8 (38.1)
Divorced	1 (4.8)
<b>Occupation, <i>n</i> (%)</b>	
Student	8 (38.1)
Employed	5 (23.8)
Not employed	8 (38.1)
<b>Economic status, <i>n</i> (%)</b>	
Low income	8 (38.1)
Medium income	8 (38.1)
High income	2 (9.5)
<b>Clinic</b>	
HDRS	19.05 ± 3.4
BMI	27.44 ± 7.4
Total testosterone (ng/ml)	0.34 ± 0.17
Free testosterone (pg/ml)	1.38 ± 0.49
ACC_GABA+ levels (mM)	1.72 ± 1.54
PCC_GABA+ levels (mM)	1.94 ± 0.70
<b>Menstrual phase, <i>n</i> (%)</b>	
Follicular	12 (57.1)

sd = standard deviation; HDRS = 17 items hamilton depression rating scale; BMI = body mass index; ACC = anterior cingulate cortex; PCC = posterior cingulate cortex; GABA = gamma amino-butyric acid.



Table 2a

Correlations between GABA levels, hormones and HDRS scores.

		ACC_GABA+	PCC_GABA+	PROG ng/ml	Estradiol pg/ml	TT ng/ml	FT pg/ml	HDRS scores	Age at FDE	No. PDE
ACC_GABA+	r	1	-.127	-.234	.401	.050	.353	-.451	-.161	.028
	p		.709	.463	.196	.877	.261	.142	.616	.932
PCC_GABA+	r		1	.273	-.442	.646**	.538	-.270	.066	-.384
	p			.324	.099	.009	.039	.330	.816	.157
PROG ng/ml	r			1	.246	.079	.233	-.538	.122	-.046
	p				.283	.734	.308	.012	.599	.843
Estradiol pg/ml	r				1	.140	-.108	-.331	-.280	.132
	p					.544	.642	.143	.220	.570
TT ng/ml	r					1	.208	-.286	-.202	-.382
	p						.365	.208	.381	.087
FT pg/ml	r						1	-.504	-.082	-.187
	p							.020	.724	.418
HDRS scores	r							1	-.208	.033
	p								.365	.886
Age at FDE	r								1	-.241
	p									.293
No. PDE	r									1
	p									

**Table 2b**

Partial correlation analysis controlling for HDRS scores.

Control variable	ACC-GABA+	PCC_GABA+	TT (ng/ml)	FT (pg/ml)
HDRS scores	ACC_GABA+ <i>r</i> 1.000	-0.167	-0.106	0.167
	<i>p</i> 0.645	0.645	0.770	0.644
	PCC_GABA+ <i>r</i> 1.000	1.000	0.677	0.813
	<i>p</i> 0.032	0.032	1.000	0.004
	TT (ng/ml) <i>r</i> 1.000	1.000	1.000	0.560
	<i>p</i> 0.092	0.092	0.092	0.092
	FT (pg/ml) <i>r</i> 1.000	1.000	1.000	1.000
	<i>p</i>			