

## Article

# Methyleugenol Has an Antidepressant Effect in a Neuroendocrine Model: In Silico and In Vivo Evidence

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**Citation:** Oliveira, M.C.N.; Cavalcante, I.L.; de Araújo, A.N.; Ferreira dos Santos, A.M.; de Menezes, R.P.B.; Herrera-Acevedo, C.; Ferreira de Sousa, N.; de Souza Aquino, J.; Barbosa-Filho, J.M.; de Castro, R.D.; et al. Methyleugenol Has an Antidepressant Effect in a Neuroendocrine Model: In Silico and In Vivo Evidence. *Pharmaceuticals* **2023**, *16*, 1408. <https://doi.org/10.3390/ph16101408>

Academic Editors: Angel Josabado Alonso-Castro and Hye Won Lee

Received: 1 July 2023

Revised: 18 August 2023

Accepted: 21 August 2023

Published: 4 October 2023



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**Abstract:** Major depressive disorder is a severe mood disorder characterized by different emotions and feelings. This study investigated the antidepressant activity of the phenylpropanoid methyleugenol (ME) in adult female mice exposed to a stress model induced by dexamethasone. The animals were randomly divided into groups containing eight animals and were pre-administered with dexamethasone (64 µg/kg subcutaneously). After 165 and 180 min, they were treated with ME (25, 50 and 100 mg/kg intraperitoneally) or imipramine (10 mg/kg intraperitoneally) after 45 min and 30 min, respectively; they were then submitted to tests which were filmed. The videos were analyzed blindly. In the tail suspension test, ME (50 mg/kg) increased latency and reduced immobility time. In the splash test, ME (50 mg/kg) decreased grooming latency and increased grooming time. In the open field, there was no statistical difference for the ME groups regarding the number of crosses, and ME (50 mg/kg) increased the number of rearing and time spent in the center. Regarding in silico studies, ME interacted with dopaminergic D1 and α1 adrenergic pathway receptors and with tryptophan hydroxylase inhibitor. In the in vivo evaluation of the pathways of action, the antidepressant potential of ME (50 mg/kg) was reversed by SCH23390 (4 mg/kg intraperitoneally) dopaminergic D1 receptor, Prazosin (1 mg/kg intraperitoneally) α1 adrenergic receptor, and PCPA (4 mg/kg intraperitoneally) tryptophan hydroxylase inhibitor. Our findings indicate that ME did not alter with the locomotor activity of the animals and shows antidepressant activity in female mice with the participation of the D1, α1 and serotonergic systems.

**Keywords:** phenylpropanoid; dopaminergic; α1 adrenergic; serotonergic; in vivo; in silico

## 1. Introduction

Major depressive disorder (MDD) is a mood disturbance influenced by different emotions and feelings. Typically, MDD may feature feelings of sadness and helplessness, reduced or lost interest in daily activities (anhedonia), lack of motivation, depressed mood, irritability, and even suicidal ideation [1,2].

The World Health Organization (WHO) reported that in 2017 there were about 322 million people with depression worldwide (World Health Organization, 2017). The first year of the COVID-19 pandemic intensified the prevalence by 25% and the records show that,

by 2030, depression is projected to become the most prevalent disease globally, becoming the main cause of population loss among all health problems (WHO 2022). Depression is about twice as prevalent in women than in men, which may be related to increased sensitivity to stress, hormonal variations, and genetic or social factors [3,4]. It is known that stress causes changes that may favor the development of depression [5]; stress is an important physiological response so that when exposure is high, hyperactivity of the hypothalamic-pituitary-adrenal axis (HPA) occurs [6], which stimulates the paraventricular nucleus of the hypothalamus to release corticotropins, which in turn promotes the production of adrenocorticotrophic hormone that acts on the adrenal gland and allows the release of cortisol, which is the main stress response hormone and acts on numerous glucocorticoid receptors [7,8]. Hyperactivity of the HPA axis negatively affects the brain's noradrenergic, serotonergic, and dopaminergic systems, which play crucial roles in regulating mood, attention, and various cognitive functions [9,10]. It is known that one of the causes identified for the development of depression is the reduction of monoamines (noradrenaline, serotonin, and dopamine) in the central nervous system [11–13]. Thus, the antidepressants used in the clinic make it possible to reduce antidepressant symptoms through mechanisms that increase the concentration of monoaminergic neurotransmitters, through the action of blocking reuptake and/or monoaminergic receptors [11,12].

Treatment for MDD includes behavioral interventions/self-care strategies, psychotherapy, and pharmacotherapy. Antidepressant drugs are considered essential to treat moderate and severe depression. However, antidepressants do not have a favorable short-term outcome (a latency period is required for a satisfactory effect to occur), and adverse effects may include cognitive and motor changes and increased vulnerability of individuals to become tolerant and dependent [13]. About 30% of patients do not achieve remission [14]. Therefore, developing new effective antidepressants is important.

An alternative for researching new drugs involves products from medicinal plants, especially those producing essential oils, which constitute a large and diverse deposit of pharmacologically active compounds [15].

Plants such as *Ginkgo biloba* L. [16], *Alpinia zerumbet* (Pers.) BLBurt & RMSm. [17], and *Lafoesia pacari* A.St.-Hil. [18] showed antidepressant activity in animal models and are mainly composed of phenylpropanoids, such as trans-anethole [19] and methyleugenol [20,21]. A study by [22] found that the phenylpropanoid eugenol (present in clove oil) generated antidepressant activity similar to that of imipramine in the behavioral tests of forced swim and tail suspension in mice.

Methyleugenol (ME, 1,2-dimethoxy-4-prop-2-en-1-ylbenzene) (an analog of the phenolic compound eugenol) is a phenylpropanoid that can be extracted from essential oils of plants, such as *Melissa officinalis* L. and *Ocimum basilicum* L. Previous studies revealed that methyleugenol has antinociceptive and anti-inflammatory pharmacological properties [23,24]. A study verified antidepressant activity using [25] male rats. Thus, the present study is the first to evaluate the antidepressant potential of phenylpropanoid methyleugenol in vivo using female mice and the possible pharmacological targets of these findings through in silico computational prospecting.

## 2. Results

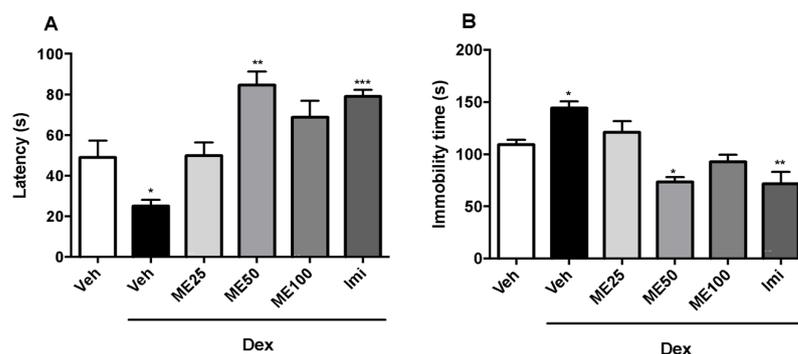
### 2.1. Behavioral Pharmacology Screening and LD50 Estimation

Under the conditions evaluated, the results showed that during the 14 days of observation, there were no deaths or lasting behavioral changes in the animals treated with a dose of 2000 mg/kg.

The phenylpropanoid ME by i.p. route showed low toxicity, with no evidence of changes in skin, hair, or eyes (a priori in class 5); thus, the LD50 was estimated to be found in values above 2000 mg kg<sup>-1</sup> and below 5000 mg kg<sup>-1</sup>. In this manner, a range of secure doses could be determined for subsequent in vivo execution.

## 2.2. Tail Suspension Test

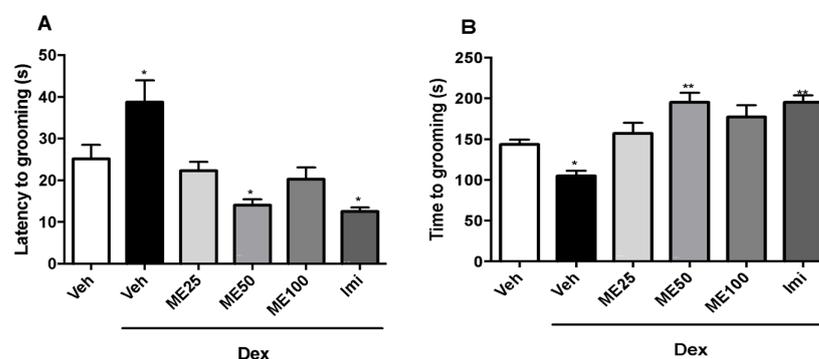
During the evaluation of latency to immobility (Figure 1A), results of one-way ANOVA revealed that acute treatment with ME (at a dose of 50 mg/kg [ $84.7 \pm 6.6$ ]) and the reference antidepressant (imipramine [Imi] [ $79 \pm 3.1$ ]) showed higher latency to immobility. The group of animals receiving dexamethasone (Dex) ( $25 \pm 3.1$ ) had their latency reduced when compared to the group receiving vehicle (Veh) ( $49.8 \pm 6.5$ ), ( $F(5, 42) = 12.51$ ;  $p < 0.0001$ ). Regarding immobility time, animals treated with ME 50 mg/kg ( $73.2 \pm 4.7$ ) and Imi ( $71.7 \pm 11.3$ ) reduced this parameter, while the dexamethasone (Dex) group ( $144.1 \pm 6.5$ ) increased this parameter compared to the Veh group ( $109.1 \pm 4.7$ ) [ $F(5, 42) = 12.7$ ;  $p < 0.0001$ ] (Figure 1B).



**Figure 1.** Effect of methyleugenol (ME 25, 50 and 100 mg/kg, i.p.), vehicle (Veh i.p.), dexamethasone (Dex 64  $\mu$ g/kg s.c.), and imipramine (Imi 10 mg/kg i.p.) on latency to immobility (A) and Time to immobility (B) in the tail suspension test. Values are expressed as mean  $\pm$  SEM ( $n = 8$ ). One-way ANOVA followed by Dunnett's Test. \*  $p < 0.05$ ; \*\*  $p < 0.01$ ; \*\*\*  $p < 0.001$  versus Vehicle (Veh).

## 2.3. Splash Test

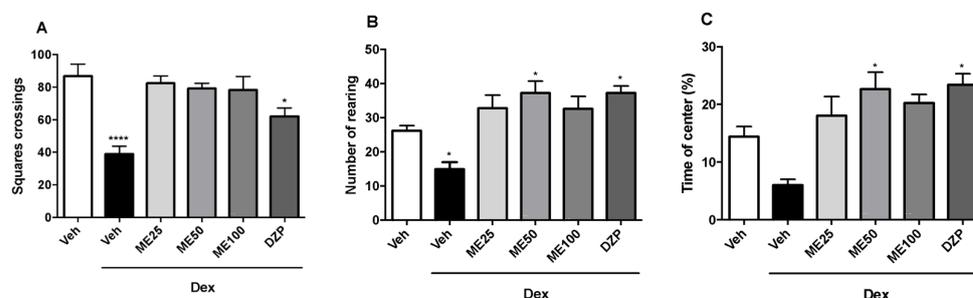
Methyleugenol in the intermediate dose (50 mg/kg) ( $22.2 \pm 2.1$ ) and imipramine (Imi) ( $12.5 \pm 0.9$ ) decreased the latency in seconds for grooming (Figure 2A) and the dexamethasone (Dex) group ( $38.7 \pm 5.1$ ) increased this parameter compared to the vehicle (Veh) group ( $25.1 \pm 3.3$ ), ( $F(5, 42) = 10.0$ ;  $p < 0.0001$ ). The grooming time (Figure 2B) increased in the groups treated with ME at a dose of 50 mg/kg ( $195 \pm 11.9$ ) and imipramine (Imi) ( $195.3 \pm 8.4$ ), while the dexamethasone (Dex) group ( $105.1 \pm 5.9$ ) had a reduction in this parameter compared to the vehicle (Veh) group ( $143.8 \pm 5.7$ ) (Figure 2B) [ $F(5, 42) = 10.7$ ;  $p < 0.0001$ ].



**Figure 2.** Effect of ME (25, 50 and 100 mg/kg i.p.), vehicle (Veh) (i.p.), dexamethasone (Dex) (64  $\mu$ g/kg s.c.) and imipramine (Imi) (10 mg/kg i.p.). (A) Latency to grooming and (B) Grooming time in the Splash Test. Values expressed as mean  $\pm$  SEM ( $n = 8$ ). One-way ANOVA followed by Dunnett's Test. \*  $p < 0.05$ , \*\*  $p < 0.01$  versus Vehicle (Veh).

#### 2.4. Open Field Test

Figure 3 demonstrates the outcomes derived from the open field assessment. In the evaluation of the number of crossings in the arena (Figure 3A), the group treated with dexamethasone (Dex) ( $38.8 \pm 4.7$ ) and diazepam (DZP) ( $61.8 \pm 5.2$ ) showed a reduction in the number of crossings and differed from the vehicle group (Veh) ( $86.6 \pm 7.4$ ) ( $F(5, 42) = 9.3$ ;  $p < 0.0001$ ).



**Figure 3.** Effect of ME (25, 50 and 100 mg/kg i.p.), vehicle (Veh) (i.p.), dexamethasone (Dex) ( $64 \mu\text{g}/\text{kg}$  s.c.) and imipramine (Imi) ( $10 \text{mg}/\text{kg}$  i.p.) on the number of crosses (A); Rearing (B); Time of center (C) in Open Field Test. Values expressed as mean  $\pm$  SEM ( $n = 8$ ). One-way ANOVA followed by Dunnett's Test. \*  $p < 0.05$ ; \*\*\*\*  $p < 0.0001$  versus Vehicle (Veh).

Regarding the rearing number (Figure 3B), animals treated with ME at the intermediate dose (50 mg/kg) ( $37.2 \pm 3.4$ ) and diazepam (DZP) ( $37.2 \pm 2.0$ ) performed this behavior more than the dexamethasone (Dex) group ( $14.8 \pm 2.1$ ) compared to the vehicle (Veh) group ( $26.1 \pm 1.6$ ) ( $F(5, 42) = 8.2$ ;  $p < 0.0001$ ).

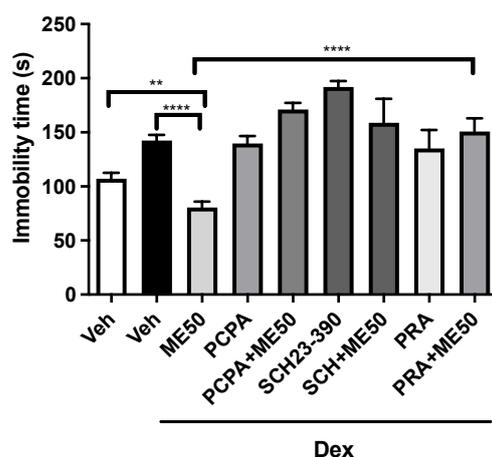
Regarding the percentage of time that the animals remained in the center of the arena (Figure 3C), treatments with ME 50 mg/kg ( $22.6 \pm 2.9$ ) and diazepam (DZP) ( $23.3 \pm 1.9$ ) increased this permanence in the center of the open field, while those treated with dexamethasone (Dex) ( $6.0 \pm 1.1$ ) reduced this percentage compared to the vehicle (Veh) group ( $14.3 \pm 1.8$ ) ( $F(5, 42) = 8.5$ ;  $p < 0.0001$ ).

#### 2.5. Assessment of the Mechanisms of Action in the Antidepressant Effects of Methyleugenol

When investigating the pharmacological pathways, we found that all groups, Veh ( $107.3 \pm 5.3$ ), DEX ( $141.3 \pm 6.5$ ), PCPA ( $139.8 \pm 6.7$ ), PCPA + ME ( $171.2 \pm 5.9$ ), SCH2330 ( $191.9 \pm 5.4$ ), SCH23390 + ME50 ( $158.8 \pm 4.5$ ), prazosin (PRA) ( $134.9 \pm 3.9$ ), and PRA + ME50 ( $150.8 \pm 4.4$ ), increased immobility time compared to the ME50 group ( $80.5 \pm 5.4$ ), which had this time reduced. A previous administration of PCPA (serotonin synthesis inhibitor), SCH-23390 (D1 antagonist), and prazosin ( $\alpha$ 1-adrenergic antagonist) prevented the antidepressant-like effect of both ME ( $F(8, 65) = 37.3$ ;  $p < 0.0001$ ) (Figure 4).

#### 2.6. In Silico Methodologies

To investigate whether the D1 dopaminergic and  $\alpha$ 1 adrenergic pathways are involved in the antidepressant impact of methyleugenol, we carried out a study using molecular docking. Through this study, it is possible to assess the probability of interaction between ME and the active site of the selected proteins. The interaction probability can be analyzed through the MolDock score, which calculates the interaction energy, where the higher the stability of the interaction, the lower the energy value. We also analyzed it through intermolecular interactions between ME and amino acid residues present in the binding site of proteins. We validated the docking analysis through redocking and RMSD calculation.



**Figure 4.** Assessment of the mechanisms of action involved in the antidepressant activity of methyleugenol (ME 50 mg/kg, i.p.) on immobility time in TSC. Vehicle (Veh i.p.), dexamethasone (Dex) (64  $\mu$ g/kg s.c.); PCPA (100 mg/kg, i.p.); SCH23390 (1 mg/kg s.c.); prazosin (PRA) (1 mg/kg i.p.). Values expressed as mean  $\pm$  SEM (n = 8). One-way ANOVA followed by Tukey's test. \*\*  $p < 0.01$ ; \*\*\*\*  $p < 0.0001$  versus ME50.

Table 1 shows the MolDock score energy values for ME of the three proteins, as well as of the molecules SCH 23390 inhibitor of the dopaminergic D1 pathway, prazosin inhibitor of the  $\alpha$ 1 adrenergic pathway, and PCPA inhibitor of the tryptophan hydroxylase pathway, and also the compounds crystallized in the binding site of each enzyme: fenoldopam (catechol agonist of the D1 dopaminergic pathway), (+)-cyclazosin (an inverse agonist of the  $\alpha$ 1 adrenergic pathway), and LP533401 (inhibitor of the tryptophan hydroxylase pathway). For a fair comparison between the energy values for every compound within the binding site of each enzyme, the measurements were also standardized according to the value of their molecular masses. Table 1 also provides the RMSD value for the crystallized ligand of each protein.

**Table 1.** The MolDock score and the RMSD of the docking molecular.

Molecules	D1 Dopaminergic		$\alpha$ 1 Adrenergic		Tryptophan Hydroxylase	
	Kcal/mol	Stand. <sup>1</sup>	Kcal/mol	Stand.	Kcal/mol	Stand.
Methyleugenol	-25.1705	-0.1412	-63.7474	-0.3576	-76.295	-0.428
Prazosin <sup>2</sup>	-	-	-120.102	-0.3132	-	-
(+)-Cyclazosin <sup>3</sup>	-	-	-135.19	-0.309	-	-
SCH 23390 <sup>4</sup>	-31.5101	-0.11	-	-	-	-
Fenoldopam <sup>5</sup>	-85.5469	-0.2798	-	-	-	-
PCPA <sup>6</sup>	-	-	-	-	-82.953	-0.4155
LP533401 <sup>7</sup>	-	-	-	-	-130.797	-0.2484
RMSD		0.19		0.3		0.6

<sup>1</sup> standardization of energies by the molar mass of each molecule; <sup>2</sup>  $\alpha$ 1 adrenergic antagonist control; <sup>3</sup>  $\alpha$ 1 adrenergic inverse agonist (PDB ligand); <sup>4</sup> D1 dopaminergic antagonist control; <sup>5</sup> D1 dopaminergic agonist (PDB ligand); <sup>6</sup> tryptophan hydroxylase inhibit control; <sup>7</sup> tryptophan hydroxylase inhibit (ligand PDB).

### 3. Discussion

In the present study, we investigated and confirmed the antidepressant psychopharmacological profile of the phenylpropanoid methyleugenol using in vivo behavioral and in silico computational methodologies.

The concept that products obtained from natural origin possess important pharmacological properties is widely spread by the scientific community and the population. However, it is important to be aware of the possibility that they present toxic effects that are often overlooked [26]. Pharmacological screening qualitatively assesses their actions on

the central nervous system (CNS) and autonomic nervous system (ANS) and estimating the acute toxicity of any substance is of utmost relevance to ensure the safety of its use. Methyleugenol at a dose of 2000 mg/kg caused behavioral changes suggestive of drugs with CNS activity but without causing toxic effects and lethality. This estimate of LD50 allowed us to evaluate the doses that were used in the behavioral tests, free of toxic effects.

The antidepressant effect of methyleugenol was investigated in a model of depression induced by dexamethasone applied i.p. [27]. Major depressive disorders have been linked to dysregulation of the hypothalamic-pituitary-adrenal (HPA) axis, which involves the secretion of glucocorticoids in excess [28], and persistent activation of its receptors by high levels of glucocorticoids can cause neural damage, with molecular and behavioral alterations, resulting in MDD and other neuropsychiatric disorders [29]. Patients treated with glucocorticoids have exhibited psychiatric and cognitive symptoms similar to depression. Already in animal models, chronic and acute glucocorticoid administration caused behavioral modifications similar to those found in humans with the disorder [30,31]. Thus, in this animal model of inducing depressive-like behavior, it is possible to mimic symptoms associated with the disease and develop pathophysiological aspects analogous to the human condition where treatment with antidepressant drugs can reverse the changes in the model [32,33].

Besides the choice of model induction that induces depressive-symptomatic behavior, the tests chosen to evaluate this behavior are also important. Indeed, the tail suspension test is one of the most widely used tests for screening antidepressant drugs [34] and is considered sensitive and specific for most classes of antidepressant drug [35]. Thus, the animal placed in an inescapable situation (trapped by the tail) develops a posture of immobility after escape attempts, and we perceived this as giving up, which characterizes a depressive state. The antidepressant substances analyzed can reverse such a condition by reducing the time of immobility [36,37]. Methyleugenol (at a dose of 50 mg/kg) increased latency in the tail suspension test, showing an effect analogous to that of imipramine and showing the antidepressant-similar activity of the substance. Although little evaluated, the verification of latency contributes to further understanding of the effects of drugs, as certain classes of antidepressants alter this parameter [38].

In 2014, Li and collaborators conducted a study using the Dex protocol, in which Dex was administered during the neonatal period (1 to 3 days) in doses ranging from 0.1 to 0.5 mg/kg body weight. Despite differences in the protocol when compared to the one used in this study, Dex increased the immobility time of the animals in the tail suspension test when compared to the control group, confirming the results obtained in this study. Still validating our data, [39] found the same result as the previous authors when administering a higher dose of Dex (1.5 mg/kg) in rats. Recently, an increase in immobility time was observed in male mice submitted to single-dose injections of Dex (64 µg/kg) in the forced swim test [31,40].

To corroborate our findings in the tail suspension test, we performed the splash test. Animals submitted to this test exhibit a shorter grooming time, which is interpreted as anecdotal behavior. In contrast, administering classical antidepressants increases the time spent on this behavior [41,42]. In turn, methyleugenol (intermediate dose 50 mg/kg) decreased the latency to grooming as well as increased the total grooming time, again showing an antidepressant-similar effect in female mice and similar to the standard imipramine. In a study similar to ours, acute treatment of a single dose of dexamethasone increased immobility time in the tail suspension test, inducing depressive behavior in animals, which was counteracted with venlafaxine, a conventional antidepressant [43].

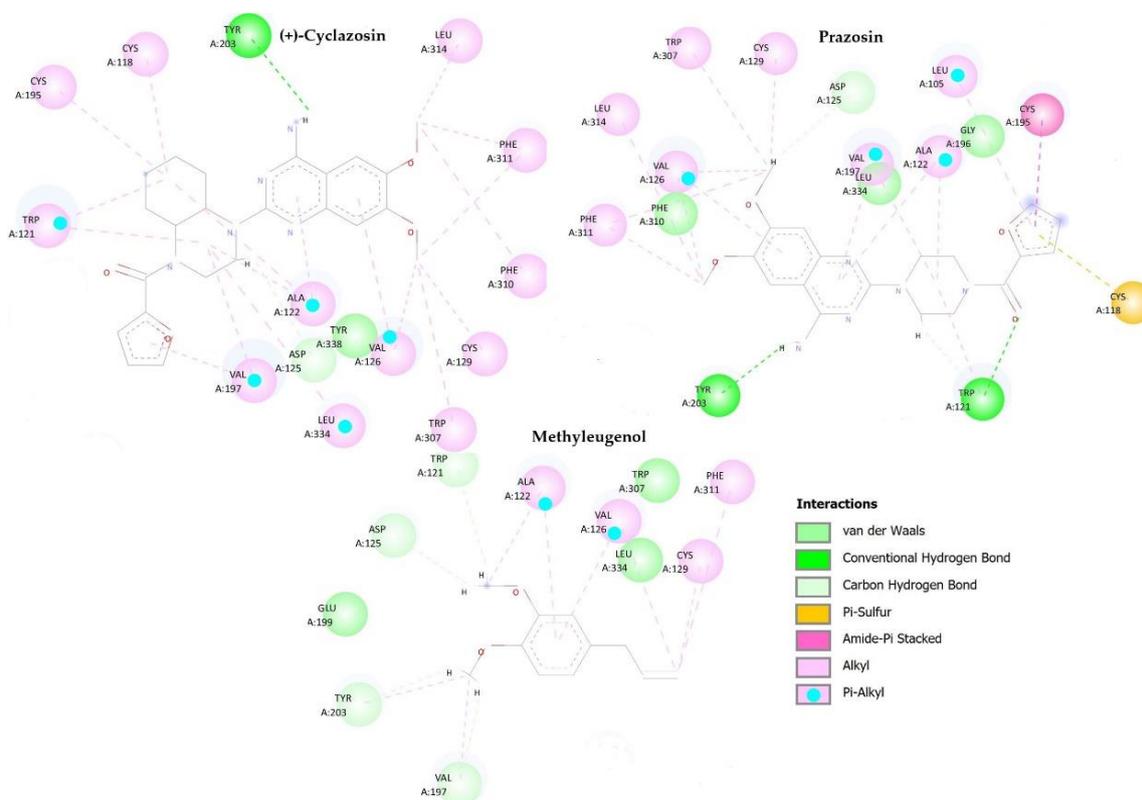
The open field test is widely used for assessing spontaneous locomotor activity and provides clues to the emotional state of rodents [44–46]. In our study, in the open field test, only the animals that received the positive control (diazepam) and the dexamethasone group reduced the number of crosses. Methyleugenol (50 mg/kg) and animals receiving diazepam increased the number of rearing and dwell time in the center, related to Dex. A methyleugenol dose of 25 mg/kg decreased the latency to rearing. This set of results

suggests that methyleugenol did not alter spontaneous locomotor activity and showed an effect indicative of anxiolytic activity, providing clues to the emotional state of the mice.

Once we identified the antidepressant-similar activity of ME in female mice, we needed to understand how this action occurs. Accordingly on the principles of the 3Rs (Replacement, Reduction, and Refinement) in non-clinical research (Russel and Burch, 1992), we conducted the study *in silico* to evaluate *in vivo* by which route(s) the ME was exerting the antidepressant effect.

In molecular docking studies, we found interaction probabilities between methyleugenol and receptors of the D1,  $\alpha$ 1 adrenergic, and tryptophan hydroxylase pathways. The  $\alpha$ 1 adrenergic protein (PDB ID 7B6W) obtained from PDB Data Bank is crystallized with the inverse agonist (+)-cyclazosin. The analysis focused on the binding site, examining the following amino acid residues: Trp 121, Asp 125, Val 126, Cys 129, Val 197, Tyr 203, and Phe 311 were found to be the most influential for enzyme activity.

(+)-cyclazosin makes hydrogen interactions with Tyr 203, while forming van der Waals interaction with Asp 125. Additionally, it establishes Alkyl-type interactions with Val 126, Phe 311, Val 127, Trp 121, and Cys 129 amino acid residues. The antagonist Prazosin participates in hydrogen interactions with Tyr 203 and Trp 121, and carbon–hydrogen bond-type interactions with Asp 125. Alkyl-type interactions emerge with Phe 311, Val 197, Val 126, and Cys 129 residues. Methyleugenol establishes carbon–hydrogen bonding interactions with Val 197, Tyr 203, Asp 125, and Trp 121, while also forming Alkyl-type interactions with Val 126, Phe 311, and Cys 129. The visual representation of these interactions between these molecules and the protein's binding site is displayed in Figure 5.



**Figure 5.** Molecular interactions between the compounds (+)-cyclazosin, prazosin, and methyleugenol, and active site's amino acid residues of the  $\alpha$ 1 adrenergic protein (PDB ID: 7B6W).

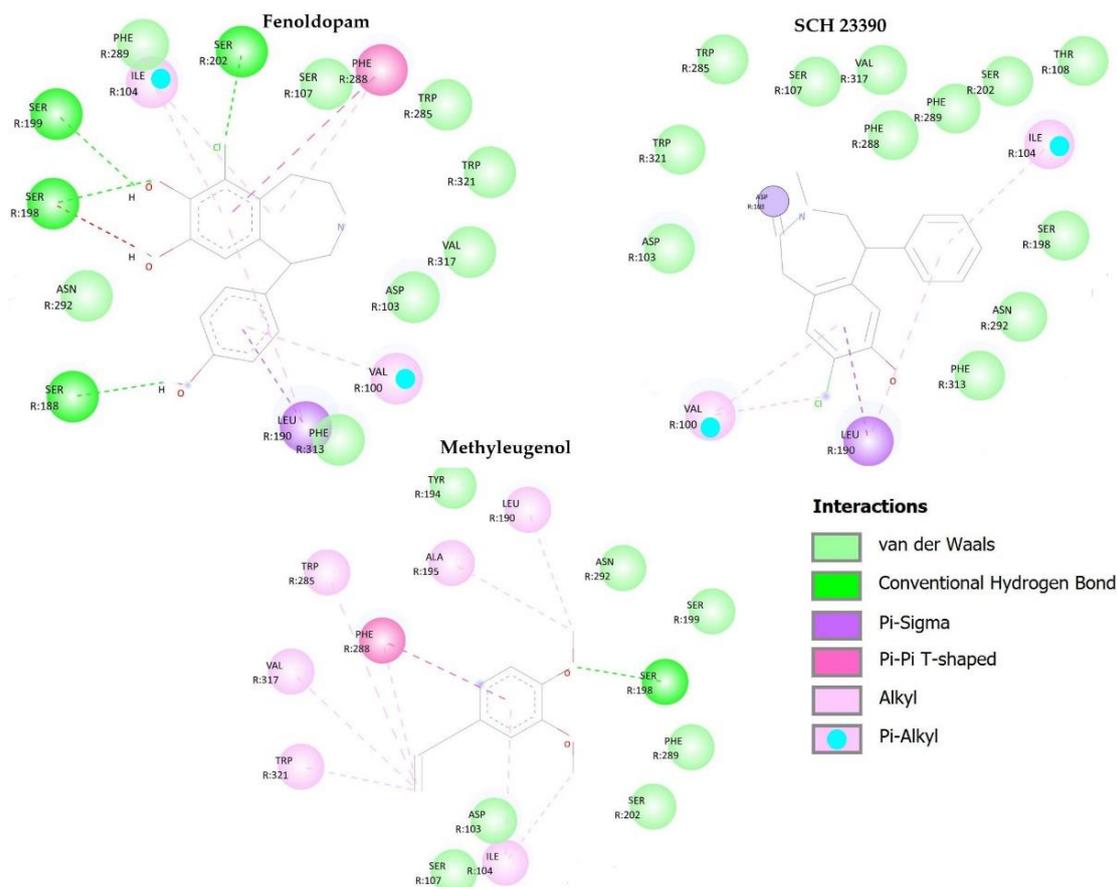
These interactions explain the energetic behavior of the connections within the compound and the binding site of the  $\alpha$ 1 adrenergic protein. Prazosin had the strongest interaction, followed by methyleugenol and (+)-cyclazosin. Imipramine was not evalu-

ated for its interaction with any of the proteins in this study as its mode of operation involves inhibiting the reuptake of serotonin and noradrenaline, rather than influencing these proteins.

The dopaminergic D1 receptor (PDB ID 7CKW), obtained from the PDB Data Bank, is crystallized with the catechol agonist fenoldopam. According to the literature, the active site of D1 receptor agonists and antagonists act on the same active site, and residues Asp103, Trp 321, Phe 289, Phe 288, Trp 285, Asn 292, and Ser 202 are considered the most influential for antagonist activity at the D1 receptor [47,48].

The interactions occurring between the agonist fenoldopam and the active site's amino acid residues of the D1 receptor are stronger than the interactions between the antagonists SCH 23390 and methyleugenol. Fenoldopam makes hydrogen-type interactions with the Ser 202; engages in van der Waals contacts with residues Asp 103, Asn 292, Trp 321, Phe 289 and Trp 285; and Pi-Pi interaction with the amino acid residue Phe 288.

Both the control antagonist used in this study (SCH 23390) and the target molecule (methyleugenol) share similarities in interactions with active sites' amino acid residues within the D1 receptor. Both perform van der Waals interactions with residues Asp 103, Asn 292, Phe 289, and Ser 202. The SCH 23390 also performs van der Waals interactions with residues Phe 288, Trp 285, and Trp 321, while methyleugenol performs alkyl or Pi-Pi type interactions with residues Trp 321, Trp 285, and Phe 288. The similarities between the antagonists are also observed in the interaction energy (Table 1). Figure 6 shows the interactions between these molecules and the D2 receptor.

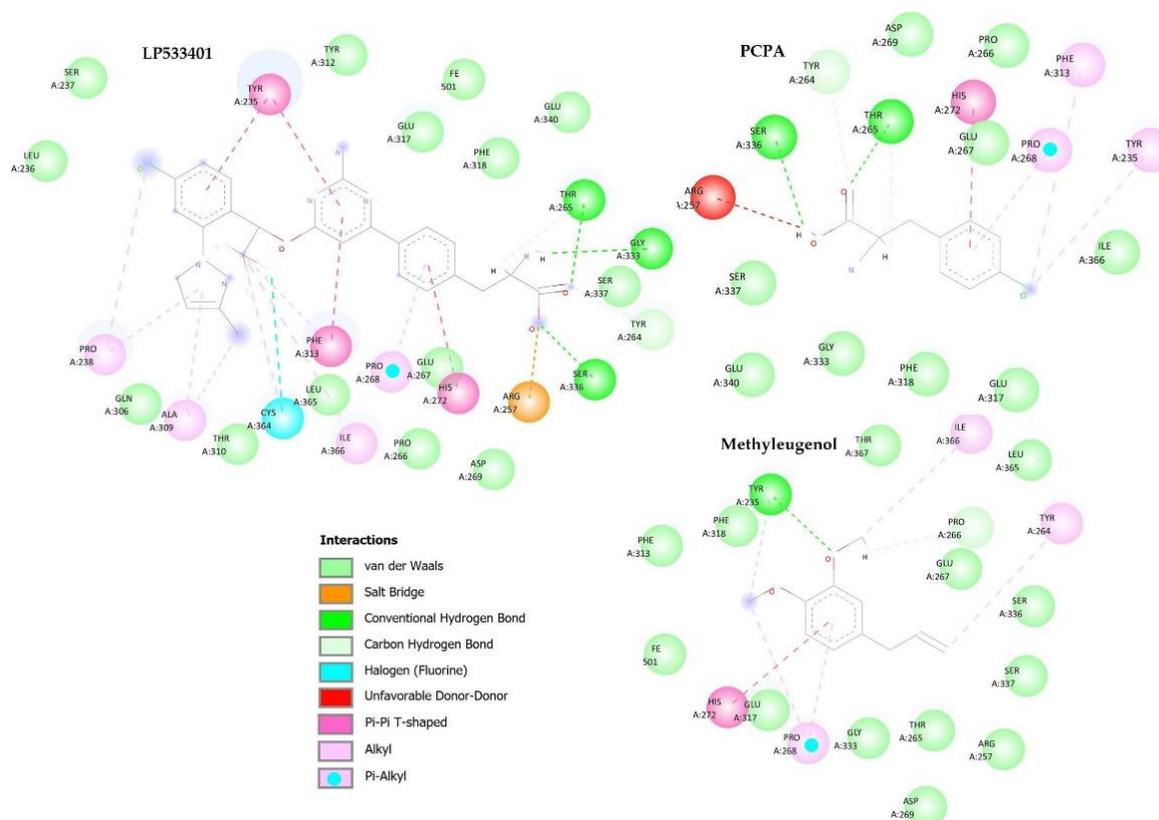


**Figure 6.** Molecular interactions between the compounds fenoldopam, SCH 23390, and methyleugenol and active site's amino acid residues of the dopaminergic protein D1 (PDB ID: 7CKW).

The tryptophan hydroxylase protein (PDB ID 7ZIK) obtained from the PDB dataBank is crystallized with the LP53340 inhibitor. Based on the analysis of the active site, the

amino acid residues Thr 265, Glu 317, Tyr 235, Ser 336, Ser 337, Gly 333, and Arg 257 were considered the most influential to inhibit the action of the protein [49].

The LP533401 inhibitor crystallized with the protein and the inhibitor used in the *in vitro* analysis, PCPA, performs hydrogen interactions with amino acid residues Thr 265 and Ser 336, and van der Waals interactions with Glu 317, Ser 337. Methyleugenol interacts with hydrogen only with the amino acid residue Tyr 235, and with the other six amino acid residues considered to be most important, methyleugenol interacts through van der Waals interactions. The LP533401 inhibitor also performs Pi-Pi-type interactions with the Tyr 235 and a salt bridge interaction with the Arg residue 257. In addition, the PCPA inhibitor performs alkyl-type interactions with the Tyr residue 235 and an unfavorable hydrogen interaction with the amino acid residue Arg 257. Figure 7 shows the interactions between these molecules in the active site of the tryptophan hydroxylase protein.



**Figure 7.** Molecular interactions between the compounds LP533401, PCPA, and methyleugenol, and active site's amino acid residues of the tryptophan hydroxylase protein (PDB ID: 7ZIK).

We assessed the possibility of interaction through the noradrenergic system *in vivo*. Because of their involvement in the pathophysiology of depression, some antidepressants increase the availability of noradrenaline at synapses [50].  $\alpha$ 1- and  $\alpha$ 2-adrenergic receptors are linked to antidepressant drug responses in behavioral models of depression [51,52].

In the *in vivo* evaluation, methyleugenol had its antidepressant effect blocked by pretreatment with prazosin ( $\alpha$ 1-adrenergic antagonist), increasing immobility time in the tail suspension test. Thus, we suggest that ME interacts with the  $\alpha$ 1-adrenergic receptors of the noradrenergic system to exert its antidepressant-similar effect. Similar to our findings, [53] demonstrated that the structural analog of eugenol ( $\alpha$ -asarone) exhibited antidepressant-sympathetic activity reversed by prazosin pretreatment.

Another important system implicated in depression is the serotonergic system. The increase in the level of 5-hydroxytryptamine (serotonin) would be related to the combatting of depression [54]. We used PCPA, a selective inhibitor of serotonin synthesis that inhibits the enzyme tryptophan hydroxylase and depletes the level of serotonin without

affecting the levels of noradrenaline and dopamine in the brain. In the present study, PCPA administered alone showed a significant difference regarding immobility time in the tail suspension test compared to methyleugenol. When the animals were pretreated with PCPA and then with methyleugenol, they also increased the immobility time, thus reversing the anti-immobility effect of methyleugenol. Similar to our results, the study conducted by [55] found that pretreatment with PCPA altered the effect on the immobility time of *Carthamus tinctorius* L. in the tail suspension test.

To continue the understanding of the mechanism of action of methyleugenol, we evaluated the possibility that its action is still through the dopaminergic system. This is especially involved in mood regulation [56], and it is known that dysregulation of the mesolimbic and mesocortical dopaminergic pathway is related to signs of melancholy and altered cognition in depression [57–59]. In addition, there are already records of the efficacy of antidepressants that act on the dopaminergic pathway [60].

Thus, involvement of the dopaminergic (D1) system in the antidepressant effect of ME was found, as pretreatment of the mice with SCH23390 (D1 receptor antagonist) blocked the effect induced by methyleugenol, thus increasing the immobility time in the tail suspension test. Only the administration of SCH23390 increased immobility, as was also observed in the findings of [35], who by administering SCH23390 to mice, reversed the antidepressant effect of gallic acid.

Our findings obtained in the present study allow us to conclude that the administration of methyleugenol in animals submitted to the dexamethasone-induced depression model produces a relevant antidepressant effect without altering locomotor activity. Its action on depressive-like behavior seems to be associated with the involvement of the D1 dopaminergic,  $\alpha 1$  adrenergic, and serotonergic systems.

## 4. Materials and Methods

### 4.1. Chemicals and Drugs

Methyleugenol (ME) was purchased from Sigma-Aldrich® Chemical (St. Louis, MO, USA), Diazepam (Cristália, Itapira, SP, Brazil), Distilled water (Federal University of Paraíba [UFPB], João Pessoa, PB, Brazil), Dexamethasone (Sigma, St. Louis, MO, USA), Ethanol (Vetec, Duque de Caxias, Brazil), R (+)-SCH23390 Hydrochloride (Sigma, USA), Imipramine (Cristália, Itapira, SP, Brazil), p-chlorophenylalanine (PCPA) (Sigma, USA), Prazosin (Sigma, USA), Sucrose (Vetec, Brazil), Tween 80 (polyoxyethylene sorbitan monoelate, Sigma, USA).

We prepared the substances in their doses minutes before use. The decimal concentrations used allowed the injection of 0.1 mL/10 g of animal weight. Methyleugenol was dissolved in distilled water with the help of tween 80, in doses 25 mg/kg, 50 mg/kg, and 100 mg/kg, for intraperitoneal (i.p.) administration. For the negative control, Tween 80 1% in distilled water (i.p.) was used.

We diluted the other drugs in distilled water or 0.9% saline solution. We administered them i.p. (methyleugenol, saline, imipramine, prazosin, and PCPA) or subcutaneously (s.c.) (dexamethasone and SCH23390).

### 4.2. Animals

Female albino Swiss mice (*Mus musculus*), weighing 25–35 g, approximately 3 months old, were housed in polyethylene cages and kept under controlled temperature ( $21 \pm 1$  °C). They had free access to pellet feed (Quimtia®) and water available in polyethylene bottles, and were kept under a 12 h light/dark cycle (the light phase from 6 am to 6 pm). We used mice for all experimental protocols. The animals were from the Animal Production Unit (UPA), Institute for Research in Drugs and Medicines (IPeFarM), UFPB. All procedures were conducted according to the guidelines proposed by the National Institutes of Health (NIH) Animal Care and Use Committee and approved by the Animal Ethics Committee (CEUA), UFPB (certificate no. 045/2017).

### 4.3. In Vivo Methodologies

#### 4.3.1. Methyleugenol Acute Toxicity Estimation and Pharmacological Screening

We performed the determination of acute toxicity in the procedures adapted from the Brazilian Health Regulatory Agency [61] and the Organization for Economic Cooperation and Development “Guidelines for testing of chemicals” [62] with modifications.

Groups of Swiss mice (consisting of three mice per group, including the control) were administered a single dose of 2000 mg/kg of ME. The control group received treatment with 0.9% saline + 0.5% Tween (i.p.). Following administration, we conducted observations at intervals of 30 min, 1st hour, 2nd hour, and 4th hour; and subsequently once daily up to the fourteenth day. After the fourteenth day, the animals were weighed and euthanized with an anesthetic overdose injection. From this step, we established the doses from the estimated LD50 value.

#### 4.3.2. Induction of Depressive-like Behavior

To evaluate the antidepressant activity, we initially induced depressive-like behavior in the animals, which was adapted from the studies of Wrobél and colleagues (2014). Dexamethasone (64 µg/kg s.c.) was administered to all groups, except the saline group, 3 h 30 min before we performed the behavioral tests. We ended the experiments by noon because of peak endogenous corticosterone release.

#### Tail Suspension Test

After the time of administration of dexamethasone (180 min our 165 min), we performed the test according to the protocol described previously [63]. Groups of mice were used (n = 8). The groups Vehicle (control group: Tween 80 0.5%), imipramine (30 mg/kg i.p.), and ME (25, 50 and 100 mg/kg i.p.) had their substances administered 30 (imipramine) and 45 (ME) minutes before the beginning of the test. After the treatment period, we suspended the mice 58 cm above the ground with tape placed approximately 1 cm from the tip of the tail. Each animal remained in this position for 6 min. The behavioral parameters observed were latency to immobility and total immobility time. The animal was classified as immobile when it showed no body movement and remained totally still.

#### Splash Test

To analyze the self-cleaning behavior (grooming) of the animals, a 10% sucrose solution was sprayed on the animal's back. The sucrose solution is thick and induces an unpleasant sensation in the animal, prompting it to engage in body cleaning, a typical self-care action. Animal cleanliness represents a form of self-care behavior, and its decrease is linked to depressive symptoms (10). After administration of dexamethasone, groups of mice were used (n = 8). The groups Vehicle (control group: Tween 80 0.5%), imipramine (30 mg/kg i.p.), and ME (25, 50 and 100 mg/kg i.p.) had their substances administered 30 (imipramine) and 45 (ME) minutes before the beginning of the test. The animal was individually placed inside an acrylic box (9 × 7 × 11 cm) and the latency parameters for the onset of self-cleaning and the time the animal remained in the self-cleaning behavior were observed for 5 min. Between observations, we sanitized the apparatus with 10% alcohol.

#### Open Field Test

The aim is to observe the locomotion and exploratory behavior of animals in the open field apparatus, similar to that described by an early study [64]. The field is manufactured of polyethylene measuring 30 cm × 30 cm, and the floor (50 cm × 50 cm) has 12 equal squares. The groups Vehicle (control group: Tween 80 0.5%), diazepam (DZP) (1 mg/kg i.p.), and ME (25, 50 and 100 mg/kg i.p.) had their substances administered 30 (DZP) and 45 (ME) minutes before the beginning of the test. The mice were positioned in the central region of the arena, and the number of animal crossings in the quadrants, dwell time in the center of the arena, amount of lifting (rearing), and latency to 1st rearing were observed for 5 min. The dwell time in the central area (4 central quadrants) was considered a measure of anxiety.

We cleaned the base of the arena with 10% ethanol between tests to avoid the effects of odors left by previous subjects.

#### Investigation of the Mechanisms of Action of the Antidepressant Activity of Methyleugenol

The dose of 50 mg/kg of methyleugenol was used because of its good performance in behavioral tests. The experimental groups received dexamethasone (64 µg/kg s.c.) about 3 h 30 min before the beginning of the evaluation; we chose the tail suspension test for this step.

To ascertain the participation of the noradrenergic system in the antidepressant activity of methyleugenol, Prazosin, an  $\alpha_1$  receptor antagonist [53,55], was used. Afterward, the different groups received the following treatments: vehicle (Tween 80 1% i.p.), ME (50 mg/kg i.p.), or prazosin (1 mg/kg i.p.). The other group received ME administration 15 min after the prazosin application. After 45 min, we performed the tail suspension test.

To investigate the serotonergic pathway, p-chlorophenylalanine (PCPA 100 mg/kg i.p.), an inhibitor of tryptophan hydroxylase-serotonin synthesis [55], was used. We also organized two other groups that received the administration of PCPA daily for 5 days; thus, on the day of the experiment, 30 min prior to the test, one group received the methyleugenol (50 mg/kg i.p.) and the other group the saline. After 30 min, we submitted the animals to the tail suspension test.

We used SCH23390 (a D1 receptor antagonist) to investigate the dopaminergic system [35]. The groups used received vehicle (Tween 80 1% i.p.), methyleugenol (50 mg/kg i.p.), or SCH23390 (1 mg/kg s.c.). Another group received SCH23390, and after 15 min, received methyleugenol. After 30 min, we started the tail suspension test.

#### 4.4. Video Recording and Analysis

All behavioral tests were engraved, and experienced researchers analyzed the behaviors in a double-blind scheme.

#### 4.5. Statistical Analysis

The results obtained during the *in vivo* experiments passed the Shapiro–Wilk normality test. Parametric tests were used, as the data showed normal distribution. Differences for single-factor group data were ruled out by one-way analysis of variance (ANOVA) followed by Dunnett or Tukey (post hoc). Data were expressed as mean  $\pm$  SEM (standard error of the mean). A 5% confidence level was adopted. All data analyses were performed with GraphPad Prism 9.0 free version (GraphPad Software, San Diego, CA, USA).

#### 4.6. In Silico Methodology

##### Investigation of the Mechanisms of Action of the Antidepressant Activity of Methyleugenol

We performed molecular docking analysis to investigate the mechanisms of the action of methyleugenol via D1 dopaminergic,  $\alpha_1$  adrenergic, and tryptophan hydroxylase receptors with antidepressant effects. Both proteins in the complex with the catechol agonist for D1 dopaminergic and inverse agonist for  $\alpha_1$  adrenergic and inhibitor LP533401 for tryptophan hydroxylase were obtained from the Protein Data Bank (PDB) at [www.rcsb.org](http://www.rcsb.org) (accessed on 1 October 2023): D1 dopaminergic (PDB ID 7CKW, ligand fenoldopam) [65]  $\alpha_1$  adrenergic (PDB ID 7B6W, ligand (+)-Cyclazosin) [66] and tryptophan hydroxylase (PDB ID 7ZIK) [49].

We employed redocking as an assessment technique for the docking protocol. In the redocking approach, we utilized the original arrangement of a crystallized ligand bound to the protein and juxtaposed it with the positioning of a ligand affixed to the active site of the identical protein through molecular docking. To gauge the efficacy, we employed the RMSD (root-mean-square deviation) metric, which measures the average spatial disparity between the crystallized ligand and the docked ligand. A docking outcome was considered acceptable if the RMSD measured up to 2.0 Å [67]. This evaluation procedure was applied to both proteins under scrutiny in the current investigation.

Methyleugenol underwent molecular docking through Molegro Virtual Docker, version 6.0.1 (MVD). During the docking procedure, elimination of all water molecules was carried out to facilitate the docking process. Preparation of enzyme and compound structures was executed using the default parameter settings within the same software package, adhering to the following configurations: ligand evaluation criteria encompassing internal electrostatics (ES), internal hydrogen bonding, Sp<sup>2</sup>-Sp<sup>2</sup> twists—all of which were verified; a total of 10 runs were conducted; MolDock SE algorithm was employed; interactions capped at 1500; population size limited to 50; a maximum of 300 steps; a neighbor distance factor of 1.00; and retrieval of up to 5 poses. The fitting process employed a GRID with a radius of 10 Å and resolution of 0.30 to encompass the ligand-binding site across both PDB files.

The central point of the target site served as the origin for the grid box, aligning with the ligand, which was then juxtaposed against the remaining trio of proteins, utilizing RMSD as the evaluative yardstick. We generated models incorporating pertinent attributes from both ligands that bore relevance to ligand binding. Employing the MolDock scoring algorithm in conjunction with the MolDock search algorithm [68], we harnessed Molegro Virtual Docker (CLC bio Company, Aarhus, Denmark) to generate five distinct poses for each alkaloid within the active site of every individual protein. Among these, the most stable conformation, characterized by the lowest interaction energy, was cherry-picked and integrated into the Discovery Studio 2021 software for visual scrutiny [69]. By employing the MolDock scoring mechanism, we calculated the energy of the crystallized ligand for each protein. Leveraging the MolDock score algorithm, the energy of the ligand from its crystallized configuration was seamlessly converted to the energy scale employed by the algorithm.

## 5. Conclusions

The investigation into the antidepressant-like effects of methyleugenol has yielded important results. In this study utilizing female mice, methyleugenol demonstrated a significant improvement in depressive symptoms induced by Dex, without altering locomotor activity. Notably, the observed effects of methyleugenol were associated with modulation of the D1 dopaminergic,  $\alpha$ 1 adrenergic, and serotonergic systems.

This suggests a potential therapeutic value of methyleugenol in mitigating depression-related behaviors in model animals. While these findings hold promise, further research is needed to elucidate the precise molecular mechanisms of methyleugenol's antidepressant action, and explore potential interactions with existing antidepressant treatments.

**Author Contributions:** M.C.N.O., I.L.C., M.G.D.S.S.S. and M.T.S.: Conceptualization. M.C.N.O., I.L.C. and R.P.B.d.M.: Methodology. M.C.N.O., R.P.B.d.M., I.L.C., A.N.d.A., A.M.F.d.S., C.H.-A. and N.F.d.S.: Investigation. M.G.D.S.S.S., M.T.S., R.D.d.C., L.S. and J.M.B.-F.: Writing—review and editing, resources. R.P.B.d.M., C.H.-A. and N.F.d.S.: In silico methodologies. J.d.S.A., R.N.A. and M.G.D.S.S.S.: Writing—original. All contributors discussed the results, commented on the manuscript, and approved the author list. All data were generated in-house and no paper mill was used. All authors have read and agreed to the published version of the manuscript.

**Funding:** This work was supported by Public Call no. 03 of Research Productivity PROPESQ/PRPG/UFPB PVB13281-2020, the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq) grant 302469/2022-2 and grant 17/2022 Paraíba State 602 Research Foundation (FAPESQ).

**Institutional Review Board Statement:** All procedures were conducted according to the guidelines proposed by the National Institutes of Health (NIH) Animal Care and Use Committee and approved by the Animal Ethics Committee (CEUA), UFPB (certificate no. 045/2017).

**Data Availability Statement:** Data is contained within the article.

**Conflicts of Interest:** The authors declare no conflict of interest.

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