

# 1 Convergent functional effects of antidepressants in major depressive 2 disorder: a neuroimaging meta-analysis

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38 **Running title:** Meta-analytic functional effects of antidepressants in depression

## 39 Abstract

40 **Background:** Neuroimaging studies have provided valuable insights into the macroscale  
41 impacts of antidepressants on brain functions in patients with major depressive disorder.  
42 However, the findings of individual studies are inconsistent. Here, we aimed to provide a  
43 quantitative synthesis of the literature to identify convergence of the reported findings at  
44 both regional and network levels and to examine their associations with neurotransmitter  
45 systems.

46 **Methods:** Through a comprehensive search in PubMed and Scopus databases, we reviewed  
47 5,258 abstracts and identified 37 eligible functional neuroimaging studies on antidepressant  
48 effects in major depressive disorder. Activation likelihood estimation was used to investi-  
49 gate regional convergence of the reported foci of consistent antidepressant effects, followed  
50 by functional decoding and connectivity mapping of the convergent clusters. Additionally,  
51 utilizing group-averaged data from the Human Connectome Project, we assessed conver-  
52 gent resting-state functional connectivity patterns of the reported foci. Next, we compared  
53 the convergent circuit with the circuits targeted by transcranial magnetic stimulation (TMS)  
54 therapy. Last, we studied the association of regional and network-level convergence maps  
55 with the selected neurotransmitter receptors/transporters maps.

56 **Results:** We found regional convergence of the reported treatment-associated increases of  
57 functional measures in the left dorsolateral prefrontal cortex, which was associated with  
58 working memory and attention behavioral domains. No regional convergence was found  
59 across foci of alterations in functional imaging associated with antidepressants. Moreover,  
60 we found network-level convergence of functional alterations in a circuit that was promi-  
61 nent in the frontoparietal and salience networks. This circuit was co-aligned with a circuit  
62 targeted by anti-subgenual TMS therapy. We observed no significant correlations between  
63 our meta-analytic findings with the maps of neurotransmitter receptors/transporters.

64 **Conclusion:** Our findings highlight the importance of the left dorsolateral prefrontal cortex,  
65 as well as frontoparietal network and the salience network in the therapeutic effects of anti-  
66 depressants, possibly associated with their role in improving executive functions and emo-  
67 tional processing.

## 68 Introduction

69 Major depressive disorder (MDD) is the most common psychiatric disorder and a leading  
70 cause of disability worldwide [1]. Despite decades of research and the development of vari-  
71 ous pharmacological, psychological, and stimulation-based treatments optimal treatment of  
72 MDD remains a challenge [2]. The conventional antidepressant medications, which are the  
73 mainstay of MDD treatment, can only achieve clinical response after several weeks of treat-  
74 ment [3] and only in around half the patients [4]. The challenges in the treatment of MDD  
75 are partly due to our limited understanding of the mechanisms by which antidepressants  
76 interact with the complex and heterogeneous neurobiology of MDD.

77 First-generation antidepressants were the monoamine oxidase inhibitors (MAOIs)  
78 and tricyclic antidepressants (TCAs) which were discovered accidentally, originally intend-  
79 ed for treating tuberculosis and schizophrenia [5]. The discovery of the antidepressant ef-  
80 fects of these medications, which possess monoaminergic properties, constituted the founda-  
81 tion of the neurotransmitter hypothesis of MDD. This hypothesis postulated that decreased  
82 levels of serotonin and norepinephrine in certain brain regions are responsible for depres-  
83 sive symptoms [6]. The neurotransmitter hypothesis of MDD led to the development of oth-  
84 er classes of antidepressants, including selective serotonin reuptake inhibitors (SSRIs) and  
85 serotonin-norepinephrine reuptake inhibitors (SNRIs), which increase the availability of  
86 synaptic monoamine neurotransmitters by inhibiting their reuptake or degradation [5].  
87 While this hypothesis has dominated the field of MDD research and treatment for decades,  
88 it is increasingly being questioned, as the supporting evidence for a decreased concentra-  
89 tion/activity of serotonin in MDD has been found inconclusive [7]. This, together with the  
90 discovery of rapid antidepressant effects of ketamine, a glutamate receptor antagonist [8, 9],  
91 suggests that the therapeutic effects of antidepressants cannot be simply explained as re-  
92 balancing the synaptic levels of the monoamine neurotransmitters. Thus, it is imperative to  
93 study the macroscale effects of antidepressant medications on the brain regions and net-  
94 works beyond their neurochemical and cellular effects. Understanding these macroscale ef-  
95 fects may help in better understanding their clinical effects on various symptoms of MDD  
96 which is ultimately needed for improving treatment outcomes.

97 Neuroimaging techniques such as functional magnetic resonance imaging (fMRI)  
98 and positron emission tomography (PET) have been used to study macroscale effects of an-  
99 tidepressants on brain activity, metabolism, or connectivity [10, 11]. However, findings of  
100 individual neuroimaging studies have been largely inconsistent, which can be attributed to  
101 their methodological and analytic flexibility, center-specific idiosyncrasies, clinical hetero-

geneity of included patients, and the small number of participants, which may make their findings less generalizable and/or reproducible [12, 13]. Neuroimaging meta-analysis is a promising tool to identify the most consistent brain findings by synthesizing the previously published literature [14, 15]. The common approach in neuroimaging meta-analyses, i.e., coordinate-based meta-analysis (CBMA), aims to find potential regional convergence across the peak coordinates of the reported effects in individual studies [16]. Several neuroimaging meta-analyses have previously used this approach to study the regional convergence of the brain effects associated with the treatment of MDD, focusing on various therapeutic approaches and different neuroimaging modalities [17–23]. However, MDD is increasingly being recognized as a brain network disorder with distributed abnormalities across the whole brain, and similarly, the antidepressants' effects could be distributed across the brain rather than localized [10]. Such distributed effects may be overlooked by the CBMA approaches which are inherently intended for regional localization of effects. Recently, a novel meta-analytic approach has been introduced which aims to identify the convergence of reported findings at the level of networks, by characterizing the normative convergent connectivity of the reported foci tested against random foci [24]. Using this approach, it was shown that despite a lack of regional convergence of reported abnormalities in MDD [25], there is a convergence of their connectivity in circuits which recapitulates clinically meaningful models of MDD [24].

Here, we aimed to identify how the findings of the previous functional neuroimaging studies on the effects of antidepressants converge on both regional and network levels by performing an updated CBMA as well as a network-level meta-analysis on the reported findings. Following, we compared our meta-analytic findings with the targets of transcranial magnetic stimulation (TMS) therapy and their associated circuits. Last, we asked whether the pattern of the observed meta-analytic effects of antidepressant medications on functional imaging can be potentially explained by the regional distribution of the neurotransmitter receptors/transporters (NRT) linked to these medications, leveraging the publicly available PET maps of neurotransmitter receptors and transporters [26].

## Methods

This meta-analysis was performed according to the best-practice guidelines for neuroimaging meta-analyses [14, 15] and is reported adhering to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement [27]. The protocol for this study

was pre-registered on the International Prospective Register of Systematic Reviews (PROSPERO, CRD42020213202).

## Search and study selection

We searched the PubMed and Scopus databases to identify peer-reviewed eligible neuroimaging studies investigating the effects of antidepressants on MDD. The search was performed in July 2022, using the keywords reported in Table S1. In addition, we searched the BrainMap annotated database of neuroimaging experiments using Sleuth by setting the diagnosis to MDD and pharmacology to the antidepressants [28–31]. Further, to avoid missing any additional relevant studies, we traced the references of relevant neuroimaging reviews/meta-analyses. Next, the duplicated records were removed, and the resulting 5258 unique records were assessed for eligibility by two reviewers (S.RJ. and S.SN.) independently. The eligibility of records was assessed first using their titles or abstracts and then, for the potentially relevant records, by examining their full texts. Any disagreements between the reviewers were resolved by another author (A.E. and A.S.).

As suggested previously [14, 15], original studies were included if: 1) they studied patients with MDD, excluding patients with other major psychiatric or neurological comorbidities and adolescent or late-life patients, 2) the patients were treated with antidepressants, 3) the antidepressants effects on the function of gray matter structures were investigated using eligible neuroimaging modalities, i.e., functional magnetic resonance imaging (including task-based [tb-fMRI], resting-state [rs-fMRI] and arterial spin labeling [ASL-fMRI]), fluorodeoxyglucose positron emission tomography (FDG-PET), or single-photon emission computed tomography (SPECT), 4) the results of pre- vs. post-treatment, treated vs. placebo/untreated, or group-by-time interaction contrasts were reported as peak coordinates of significant clusters in standard spaces (Montreal Neurological Institute [MNI] or Talairach) or were provided by the authors at our request, 5) the analysis was performed across the whole brain, was not limited to a region of interest (ROI), and small volume correction (SVC) was not performed, as these approaches are biased toward finding significance in the respective areas, hence violating the assumption of ALE method that all voxels of the brain have a unified chance of being reported [14, 15], and 6) at least six subjects were included in each group (Fig. 1).

## Data extraction and preprocessing

From the eligible studies, we extracted demographic and clinical data (number of participants, age, sex, response to treatment, medications, treatment duration), methodological details (imaging modality, scanner field strength, task paradigm, software package, statistical

contrast, and the multiple comparisons correction method), as well as the peak coordinates/foci (x, y, z) of experiments' findings. Of note, we use the term "study" to refer to an individual publication, and the term "experiment" to refer to the individual group-level contrasts reported within each "study" (e.g., Treated > Untreated). Following the data extraction, the coordinates reported in Talairach space were transformed into MNI space [32], so that all the experiments are in the same reference space. If the applied reference space was not explicitly reported or provided by authors after our request, we assumed the default settings of the software packages were used for normalization [14, 15]. In addition, to avoid spurious convergence over the experiments performed on the same/overlapping samples (reported within or across studies), in each meta-analysis, we merged the coordinates from multiple experiments pertaining to the same/overlapping samples, to make sure that each study contributes once per analysis, as suggested previously [14, 15, 33].

## Activation likelihood estimation

The revised version of the activation likelihood estimation (ALE) method [16] was used to test the regional convergence of the reported differences against the null hypothesis of randomly distributed findings across the brain. In this method, the peak coordinates are convolved with 3D Gaussian probability distributions that have a full width at half maximum inversely proportional to the sample size. This allows experiments with larger samples to have a greater statistical certainty in the meta-analysis. Next, for each experiment, the convolved foci are combined to generate per-experiment "modeled activation" (MA) maps. Subsequently, the MA maps for all the experiments included in the meta-analysis were combined into an ALE score map, representing the regional convergence of results at each location of the brain. The ALE score map was then statistically tested against a null distribution reflecting randomly distributed findings, to distinguish true convergence from by-chance overlap [16, 33]. Finally, to avoid spurious findings [16], the resulting p-values are corrected for multiple comparisons using the family-wise error correction at the cluster level (cFWE), thresholded at  $p < 0.05$ .

In addition to an ALE meta-analysis on all included experiments (the 'all-effects' ALE) we performed several complementary ALE meta-analyses based on the direction of the effect (treatment contrast i.e., Treated > Untreated [Tr+] or Untreated > Treated [Tr-]), imaging modality, study design, and type of the antidepressants. The analyses were performed only if a sufficient number of experiments were included in each category, as ALE analyses with too few experiments are likely to be largely driven by a single experiment, and therefore lack sufficient statistical power to provide valid results [34].



## **Functional decoding of the convergent clusters**

We applied the data from task-based functional neuroimaging experiments and their annotated behavioral domains (BD) included in the BrainMap database [28–31] to identify BDs that were significantly associated with the convergent clusters identified in the main ALE analyses [35]. In particular we used binomial tests to assess whether the probability of each cluster activation given a particular BD, i.e.,  $P(\text{Activation} | \text{BD})$ , is significantly higher than the overall a priori chance of its activation across all BDs, i.e.,  $P(\text{Activation})$ .

## **Meta-analytic coactivation mapping of the convergent clusters**

We investigated the task-based functional connectivity of the convergent clusters identified in the main ALE analyses using meta-analytic coactivation mapping (MACM) [36]. We used the data from task-based functional neuroimaging experiments on healthy individuals included in the BrainMap database [28–31]. For each identified convergence cluster from the main ALE analyses, we identified all the experiments that reported at least one focus of activation therein, and performed a meta-analysis across those experiments, thresholded at  $p < 0.05$  and cFWE-corrected. This approach identifies brain regions that are consistently co-activated with the convergent cluster across all task-based functional neuroimaging experiments.

## **Resting-state functional connectivity of the convergent clusters**

We obtained the group-averaged dense resting-state functional connectivity (RSFC) matrix of the Human Connectome Project (HCP) dataset ( $n=1003$ ) available in *Cifti* format [37, 38]. The MNI coordinates of the convergent cluster peaks in volumetric space were mapped to their closest ‘grayordinate’ (cortical vertex or subcortical voxel) in the MNI *Cifti* space based on Euclidean distance. Subsequently, the whole-brain RSFC maps of the foci were extracted from the HCP dense RSFC and were plotted.

## **Network-based meta-analysis**

In addition to the conventional CBMA, we performed network-based meta-analyses [24], to identify convergent connectivity maps of the reported foci compared to randomly distributed foci. We used the normative group-averaged dense RSFC matrix of the HCP dataset in these analyses. For the given set of experiments in the all-effects, Tr+ and Tr- analyses, we first extracted the MNI coordinates of all the reported foci in the included experiments. The MNI coordinates of the foci in volumetric space were then mapped to their closest grayordinate based on Euclidean distance. The foci with no grayordinate in their 10 mm radius were excluded (19 out of 534). Of note, the median distance of the mapped grayordinates from the MNI coordinates of foci was 2.44 mm. Following, the whole-brain

RSFC maps of the foci were extracted from the HCP dense RSFC and averaged to create the RSFC map of the set of experiments. The observed RSFC map was compared to a permutation-based null distribution of RSFC maps to create a z-scored map. Specifically, in each of 1000 permutations we randomly sampled an equal number of foci as reported in the included experiments and averaged their RSFC maps, resulting in a set of 1000 null RSFC maps. Following, we subtracted the mean of the null RSFC maps from the observed RSFC map and divided the result by the standard deviation across the null RSFC maps, to compute the Z-scored RSFC map of the given condition. These maps, referred to as ‘convergent connectivity maps’, reflect greater- or lower-than-chance connectivity of the reported foci to the rest of the brain, and may indicate circuits affected by antidepressant treatment.

Subsequently, we investigated the distribution of the convergent connectivity maps of each condition across the seven canonical resting-state networks (RSNs). The statistical significance of these associations was assessed using a spin test, which accounts for the spatial autocorrelation in the brain. In this approach, we first calculated the observed mean of convergent connectivity map of a given condition within each RSN, and then tested whether the observed means are more extreme than null means based on 1000 surrogate maps created by randomly spinning the convergent connectivity map on the cortical sphere. The spin surrogate maps were generated as implemented in the *neuromaps* toolbox [39, 40].

## Association with transcranial magnetic stimulation targets

We compared the location of the ALE convergent clusters with four TMS target coordinates, including the anatomical 5-CM rule site (MNI -41, 16, 54) [41] and the anti-subgenual site (MNI -38, 44, 26) [42] used in clinical trials as well as the peak sites associated with the clinical improvement of dysphoric symptoms (MNI -32, 44, 34) and anxiosomatic symptoms (MNI -37, 22, 54), as reported previously [43]. In addition, we extracted the RSFC maps of grayordinates corresponding to these coordinates from the HCP dense RSFC and evaluated their spatial correlations with the convergent connectivity maps of antidepressant effects as well as the RSFC maps of ALE convergent clusters by using spin permutation, as described above.

## Association of meta-analytic findings with neurotransmitter receptor/transporter densities

The PET maps of tracers associated with NRT were obtained from a previous study [26], which curated these maps from various sources [44–61]. These maps were based on tracers for serotonergic and noradrenergic receptors/transporters (5HT1a, 5HT1b, 5HT2a, 5HT4, 5HT6, 5HTT, NAT) as well as the NMDA receptor. The PET maps were available in MNI



volumetric space and were parcellated using Schaefer-400 parcellation in the cortex (400 parcels) and Tian S2 parcellation in the subcortex (32 parcels), and were subsequently Z-scored across parcels. In case multiple maps were available for a NRT we calculated an averaged map weighted by the sample size of the source studies.

We then calculated the correlation of parcellated NRT maps with the convergent connectivity maps while accounting for spatial autocorrelation by using variogram-based permutation. In this approach, random surrogate maps were created with variograms that were approximately matched to that of the original map, as implemented in *BrainSMASH* [62]. Furthermore, we used spin permutation to test for over-/underexpression of the NRTs in the ALE convergent clusters. We first rank-normalized the z-scored and parcellated maps of NRTs and after projecting them to the cortical surface, calculated the median rank-normalized density of each NRT within the convergent cluster. Next, we compared the observed median densities against a null distribution calculated based on surrogate NRT maps with preserved spatial autocorrelation. These surrogate maps were created by spinning the parcels on the cortical sphere, as implemented in the *ENIGMA Toolbox* [63]. In all the tests, the resulting p-values were corrected for multiple comparisons across the NRT maps by using false discovery rate (FDR).

## Results

### Experiments included in the meta-analyses

The study selection process is depicted in Fig. 1. We screened 5258 records resulted from our broad and sensitive search, and assessed 586 full texts for eligibility to finally include 37 studies and 31 experiments with non-overlapping samples (Table 1 and Table S2) [64–100]. Collectively 862 MDD patients were included in the experiments. The patients were treated using SSRIs (n=18), ketamine (n=7), S/NRIs (n=7), mirtazapine (n=2), clomipramine (n=1), amesergide (n=1), quetiapine (n=1), or bupropion (n=1). In seven experiments the patients received variable medications. The imaging modalities included were tb-fMRI (n=18), FDG-PET (n=4), H<sub>2</sub>O-PET (n=1), rs-fMRI (n=4), ASL-fMRI (n=2) and <sup>99m</sup>Tc-HMPAO SPECT (n=3).

297

## 298 Convergent localized effects of antidepressants in the dorsolateral pre- 299 frontal cortex

300 No significant regional convergence was found in our ALE meta-analysis on all the included  
301 experiments. Furthermore, the subgroup analyses limited to specific types of treatments or  
302 modalities showed no significant convergence (Table 2). However, among the Tr+ experi-  
303 ments (n=21) showing increases in functional imaging measures associated with the treat-  
304 ment, we observed a significant cluster of convergence in the left middle frontal gyrus with-  
305 in the dorsolateral prefrontal cortex (DLPFC) (MNI -38, 30, 28; 129 voxels) (Fig. 2). The con-  
306 vergence in this cluster was driven by contributions from seven experiments [67, 78–81, 83,  
307 85, 99]. The relative contribution of experiments using different medications included SSRIs  
308 (58.3%), ketamine (25.4%), and variable classes (16.1%). The contribution of experiments us-  
309 ing PET (56.1%) was the highest, followed by fMRI (43.6%) and SPECT (0.3%). The scanning  
310 paradigms of contributing experiments included resting state (81.2%) and emotional tasks  
311 (18.8%). In the complementary ALE analyses on subgroups of the Tr+ experiments, we ob-  
312 served additional/different clusters of convergence (Fig. S1). The subgroup analysis limited  
313 to the 19 Tr+ experiments that only reported pre- versus post-treatment effects revealed  
314 clusters in the left (MNI -38, 30, 28; 132 voxels) and right DLPFC (MNI 44, 26, 24; 92 voxels).  
315 Similarly, the meta-analysis among Tr+ experiments using treatments other than ketamine  
316 (15 experiments) showed clusters of convergence in the left (MNI -38, 30, 30; 95 voxels) and  
317 right DLPFC (MNI 44, 26, 24; 106 voxels). The meta-analysis on Tr+ effects reported after  
318 more than 4 weeks of treatment (13 experiments) revealed a convergent cluster in the medial  
319 superior frontal gyrus (MNI 8, 54, 30; 104 voxels). In the meta-analysis on the Tr+ experi-  
320 ments reporting • 50% rate of clinical response, we found a cluster of convergence in the left  
321 supramarginal gyrus (MNI -48, -44, 40; 123 voxels) in addition to the right DLPFC (MNI 44,  
322 26, 24; 123 voxels).

323 We observed no significant convergence in the ALE meta-analyses performed across  
324 Tr- experiments showing decreases of functional imaging features associated with the  
325 treatment (22 experiments) as well as their more specific subgroups (Table 2).

326

## Functional decoding and MACM of the dorsolateral prefrontal cortex cluster

Next, we studied the behavioral relevance and connectivity of the convergent cluster identified in the ALE meta-analysis on Tr+ experiments within the left DLPFC. Using the data from BrainMap database we observed that the behavioral domains of working memory (likelihood ratio = 1.85) and attention (likelihood ratio = 1.43) were significantly associated with the activity of this cluster.

The MACM of the left DLPFC cluster showed its significant co-activation with regions in the prefrontal cortex, superior parietal lobule, insula and anterior cingulate and paracingulate cortices ( $p_{\text{cFWE}} < 0.05$ ; Fig. 3A). In addition, the RSFC of the left DLPFC cluster center based on the HCP dataset dense connectome showed its connectivity with widespread regions in the prefrontal cortex, superior frontal gyrus, insula, anterior cingulate, paracingulate cortices, supramarginal gyrus, inferior temporal gyrus and basal ganglia, in contrast to its negative resting-state anti-correlation with regions in the subgenual anterior cingulate cortex, orbitofrontal cortex, posterior cingulate, angular gyrus, and temporal pole (Fig. 3B).

## Convergent connectivity of antidepressant effects

Having characterized the regional convergence of antidepressant effects using ALE, next, we aimed to investigate meta-analytic effects of antidepressants at a circuit level, following a recently introduced approach [24]. To do so, we used the group-averaged dense functional connectome obtained from the HCP dataset and quantified the convergent connectivity of the reported coordinates of antidepressant effects, which was compared against null connectivity patterns of random points.

The peak coordinates of all the included experiments, indicating alterations in functional imaging measures associated with antidepressants (515 foci from 31 experiments), showed greater-than-chance connectivity of these coordinates with regions in the dorsolateral and medial prefrontal cortex, anterior insula, posterior cingulate cortex, supramarginal gyrus, inferior temporal gyrus, primary visual cortex, and basal ganglia, in contrast to their lower-than-chance connectivity with regions in the subgenual anterior cingulate, posterior cingulate, angular gyrus, temporal pole and superior frontal gyrus (Fig. 4A). We observed significantly higher values of convergent connectivity in the frontoparietal (FPN;  $\langle Z \rangle = 4.23$ ,  $p_{\text{spin}} < 0.001$ ) and salience (SAN;  $\langle Z \rangle = 3.16$ ,  $p_{\text{spin}} = 0.047$ ) networks (Fig. 4B). Accordingly,

evaluating the distribution of the foci across RSNs revealed significantly higher than chance number of foci in the FPN ( $n=84$ ,  $p_{\text{spin}} < 0.001$ ; Fig. 4B).

The network-level analyses separately performed on the Tr+ (180 foci from 21 experiments) and Tr- effects (206 foci from 22 experiments) revealed convergent connectivity maps which were anti-correlated with each other ( $r = -0.45$ ,  $p_{\text{variogram}} < 0.001$ ; Fig. S2A). The convergent connectivity map of the Tr+ effects was significantly more prominent in the FPN ( $\langle Z \rangle = 5.41$ ,  $p_{\text{spin}} < 0.001$ ) and default mode network (DMN;  $\langle Z \rangle = 1.71$ ,  $p = 0.047$ ). On the other hand, the convergent connectivity map of Tr- effects was significantly more prominent in the visual (VIS;  $\langle Z \rangle = 3.17$ ,  $p_{\text{spin}} < 0.001$ ) and somatomotor (SMN;  $\langle Z \rangle = 2.79$ ,  $p_{\text{spin}} = 0.016$ ) networks.

### The association of antidepressants effects with TMS targets

The left DLPFC is suggested to be the optimal stimulation target in the TMS treatment of MDD [101, 102]. We next explored whether our meta-analytic findings on the convergent effects of antidepressants might correspond with the different TMS targets. We compared the location of the left DLPFC cluster identified in the Tr+ ALE meta-analysis and observed it was closer to the “anti-subgenual” (14 mm) and “dysphoric” (16 mm) TMS targets than the “5-CM” (29 mm) and “anxiosomatic” (27 mm) targets (Fig. S3A). Moreover, the RSFC map of the Tr+ cluster and the network-level meta-analysis convergent connectivity maps were positively correlated with the RSFC maps of anti-subgenual and dysphoric TMS targets but negatively correlated with the RSFC maps of 5-CM and anxiosomatic stimulation sites (Fig S3B).

### The association between neurotransmitter receptor/transporter maps and meta-analytic effects of antidepressants

Lastly, we studied whether the regional and network-level convergence of antidepressant effects colocalizes with the spatial distribution of serotonergic and noradrenergic NRTs as well as NMDA receptor (Fig. 5a) [26]. We first focused on the cluster of convergence of Tr+ effects in the left DLPFC and quantified the median density of each NRT (rank-normalized) in this region, showing the varying density of the NRTs. However, none of the NRTs were significantly over-/underexpressed in this cluster (Fig. 5b). Next, we evaluated the correlation of parcellated convergent connectivity map with the NRT maps but observed no signifi-

cant correlations (Fig. 5c). Similarly, the NRT maps were not significantly correlated with the parcellated convergent connectivity maps of Tr+ and Tr- experiments (Fig. S4).

## Discussion

In the present study, we synthesized findings of the neuroimaging literature on the brain effects associated with pharmacotherapy of MDD at regional and circuit levels. At the regional level, our meta-analysis showed no significant convergence across all the included experiments, though we found convergence of the reported treatment-associated increases of functional measures in the left dorsolateral prefrontal cortex. This convergent cluster was associated with working memory and attention behavioral domains and showed meta-analytical coactivation with regions in the prefrontal cortex, superior parietal lobule and insula. Extending our meta-analysis to the circuit level, we investigated the convergent connectivity of the reported foci and found a circuit that was most prominent in the frontoparietal and salience resting-state functional networks. Following, we found that this circuit was co-aligned with a circuit targeted by anti-subgenual TMS therapy. Last, we studied whether the spatial pattern of the observed regional and network-level meta-analytic effects co-align with the maps of receptors and transporters related to the studied antidepressants and found no significant associations.

### Convergent effects of antidepressants on the frontoparietal and salience networks

The pathology in MDD is increasingly thought to be distributed across brain regions and circuits, rather than being localized [10]. In fact, previous ALE meta-analyses aimed at localizing the convergent abnormalities in MDD have revealed minimal or no regional convergence [25, 103, 104]. However, a recent study revisited the functional imaging literature on the brain abnormalities in MDD and, using a connectomic approach, showed that the reported abnormalities in MDD are connected to circuits involving regions such as DLPFC, insula, cingulum, pre-supplementary motor area and precuneus [24]. These circuits were shown to recapitulate clinically meaningful models of MDD, such as a lesion-derived MDD circuit [24]. Here, following a similar approach, we also found network-level convergence of the reported findings in a circuit of brain regions most prominent in the frontoparietal and the salience resting-state networks.

The prominence of the convergent connectivity map in the FPN, together with the convergent cluster found in the ALE meta-analysis on the Tr+ experiments, highlights the

importance of DLPFC and FPN in the therapeutic effects of antidepressants. These regions play pivotal roles in higher executive and cognitive functions of the brain, which are shown to be impaired in patients with MDD [105–108]. More severe deficits of executive functions are linked with higher severity of depressive symptoms [107]. The executive and cognitive dysfunction in MDD is thought to contribute to emotional dysregulation, which is a hallmark of MDD psychopathology [109, 110]. Specifically, patients with MDD might have impairments in cognitive control when processing negative emotions, deficits in the inhibition of mood-incongruent material, and difficulties in attentional disengagement from negative stimuli, which are among the mechanisms that are thought to contribute to emotional dysregulation [109, 110]. Indeed, antidepressant medications are shown to improve the executive functioning of patients with MDD, in the domains of attention and processing speed [111], psychomotor speed [112] and cognitive interference inhibition [108], and can lead to better emotional regulation strategies [113]. Hypoactivity of the prefrontal cortex in MDD is thought to contribute to the deficits of executive functioning [107, 114, 115] and emotional regulation [110, 114, 116, 117]. For instance, patients with MDD have shown a reduced activity of the DLPFC during an attentional interference task using emotional distracters [118], which can be normalized by antidepressants [119]. Furthermore, the FPN in patients with MDD shows reduced within-network connectivity and decreased connectivity with the parietal regions of the dorsal attention network, as reported by a meta-analysis on seed-based RSFC studies [120]. In addition, hypoconnectivity of the FPN with the rest of the brain has been observed in relation to depressive symptoms in the general population [121]. The treatment of MDD using various therapeutic approaches can affect intra- and inter-network connectivity of the FPN [10] with the DMN [122] and SAN [123–125]. The abnormalities of SAN connectivity in MDD include decrease of within-network as well as SAN-FPN connectivity [10, 124, 126]. The SAN consists of regions such as the anterior insula and dorsal anterior cingulate and is involved in guiding behavior in response to salient events and the processing of emotional information and rewards [10, 120, 127]. Accordingly, the deficits in within- and between-network connectivity of SAN is suggested to contribute to the symptoms of depressed mood, anxiety, and anhedonia in MDD [10, 128–130]. Overall, MDD is characterized by altered function and connectivity of distributed networks, importantly including the FPN and SAN, but also the DMN and limbic networks, which can be modulated by the treatment (see review in Chai et al. [10]). Of note, our network-level meta-analysis was performed using resting-state imaging data of healthy subjects, and therefore, provides an indirect view on network-level actions of antidepressants. Further large-scale studies are



needed to investigate these effects using connectomic approaches on MDD patients treated with antidepressants.

## Similar networks may be modulated by antidepressants and TMS

We found convergent network-level and regional effects prominent in the FPN and more specifically the DLPFC. The importance of DLPFC and FPN in MDD treatment has further been observed in non-pharmacological therapeutic approaches. Psychotherapy of patients with MDD and PTSD is shown to normalize the activity of DLPFC and increase within-network connectivity of FPN [131]. In addition, the left DLPFC is suggested to be the optimal target of the stimulation in TMS therapy of MDD [101, 102]. High-frequency TMS applied to this region increases its activity, which in turn is thought to have therapeutic effects by modulating the activity of a network of connected regions [41, 101]. A recent retrospective analysis of the clinical effects of the different TMS locations revealed that the improvement in distinct clusters of depressive symptoms, i.e., anxiousomatic and dysphoric symptoms, relates to stimulating targets that engage distinct circuits [43]. Interestingly, we observed that the convergent connectivity map of the antidepressant effects as well as the RSFC map of the left DLPFC cluster were positively correlated with the “dysphoric” peak target circuit and negatively correlated with the circuit of “anxiousomatic” peak target. While the anxiousomatic circuit corresponds to the RSFC map of the anatomical “5 cm” TMS target used in the early clinical trials, the dysphoric cluster circuit resembles that of the more recent connectivity-based “anti-subgenual” TMS targets [42, 43, 101, 102]. The latter circuit is characterized by negative connectivity to sgACC, which was also found in our meta-analytic convergent connectivity map and the RSFC map of the left DLPFC cluster identified in the ALE meta-analysis. Indeed, hyperactivity of the subgenual anterior cingulate cortex in MDD is thought to contribute to increased processing of negative stimuli [115]. Therefore, both the anti-subgenual TMS and antidepressant treatment of MDD might modulate the activity of a similar circuit including DLPFC and sgACC. This circuit, based on the findings from TMS studies, seems to be more effective on dysphoric symptoms [43]. This is particularly interesting given that antidepressant medications have been shown to be more effective for the core emotional symptoms (e.g., sadness) than for sleep and atypical symptoms (e.g., psychomotor agitation) [132]. Future research is needed to more directly address the question of how the brain function changes in association with the effects of antidepressants on the different subsets of depressive symptoms, especially from a meta-analytical perspective.

## Lack of association between neurotransmitters and the system-level effects of antidepressants

The neurotransmitter hypothesis of MDD suggests that the dysregulation of the monoaminergic neurotransmitter systems is central to the pathophysiology of MDD, and antidepressants act by normalizing the dysregulations of these neurotransmitter systems [5, 6, 133]. In our analyses, we found that the PET-based maps of serotonergic and noradrenergic receptors and transporters were not significantly correlated with the regional and network-level meta-analytic effects of antidepressants. This suggests a divergence between the antidepressant effects on brain function, as observed in functional imaging studies, and the regions where their target NRTs are highly expressed. The observed divergence raises the question of what mechanisms may relate the micro-scale actions of antidepressants on the NRTs to their system-level effects on brain function. Molecular imaging techniques combined with functional imaging might provide some clues to this link. The findings of molecular imaging studies in MDD and its treatment are diverse (see a comprehensive review by Ruhé et al. [133]). For example, there has been some evidence of decreased serotonin synthesis rate in the prefrontal and cingulate cortex of patients with MDD [134–136]. However, a recent umbrella review summarizing the research on the serotonin hypothesis of MDD concluded that there is a lack of convincing evidence for the association of MDD with serotonergic deficits such as a lower serotonin concentration or changes in the receptors [7]. Moreover, the antidepressive effects of ketamine, an NMDA receptor antagonist, highlight the importance of the other, non-monoaminergic neurotransmitters in the pathophysiology and treatment of MDD [8, 9]. These findings have increasingly led to the belief that the monoaminergic neurotransmitter hypothesis of MDD may not provide a full understanding of the disease [7, 137, 138]. However, these neurotransmitter systems are indeed involved in the pathophysiology and treatment of MDD, yet their role needs to be revisited in a broader context. One promising area for future research on this matter is the computational modeling of the changes in brain activity in response to pharmacological interventions using biophysical network models [139] coupled with biological models of neurotransmitter systems [140].

## Research in context, strengths, and limitations

The neuroimaging effects of antidepressant treatment in MDD has been previously investigated in a number of CBMAs [17–23] (Table 3). These studies have focused on various types of treatment, with more specific (e.g., only SSRI medications [20]) or broader (e.g., pharmacotherapy, psychotherapy, and ECT [22]) scopes compared to our study. In addition, various

imaging modalities under different conditions have been investigated, from focusing on fMRI experiments during emotional processing tasks [23] to a broader multimodal investigation of the functional and structural imaging experiments [22]. Given the differences in the scope and methodology of the previous CBMAs, it is not unexpected to find that they have reported different meta-analytic findings. However, it is important to note that according to the current guidelines [14, 15], there are a few methodological issues to consider in some of the (earlier) CBMAs which may have influenced their findings. These issues include: i) a small number of experiments included in the main or subgroup analyses, which can limit the power and increase the risk of a single experiment dominating the findings [34], ii) including explicit or hidden ROI-based experiments which are biased to inflate significance in the selected region, iii) using less stringent methods of multiple comparisons correction, e.g., thresholding clusters simply by applying a lenient cluster extent and height, or by using FDR, or iv) performing ALE using the earlier versions of GingerALE (< 2.3.3), in which a software bug was reported that can lead to more lenient multiple comparisons correction [141]. Here, we set out to avoid such methodological issues by following the best-practice guidelines for the CBMAs [14, 15]. Furthermore, we provided network-level accounts of the effects of antidepressants reported in the literature [24], which acknowledges that the effects may be distributed rather than localized, and in doing so, complements the conventional CBMA approach of identifying regional convergence.

Our study had a few limitations as well. The heterogeneity of the included experiments, in terms of imaging modalities, conditions, medications, and the clinical characteristics of the patients, limits our findings, but at the same time, enables identifying convergence of the effects that may be robust to such variability. To identify convergence among more selected, harmonized, subsets of the experiments, we planned subgroup analyses. Yet, it was not possible to perform some of the planned analyses, such as a comparison of different medication classes, due to a limited number of experiments identified in each subgroup. Moreover, here we studied the neuroimaging effects of antidepressant medications on patients with MDD who had been treated but not necessarily responded to the treatment. Of note, in a small subgroup analysis focused on experiments reporting clinical response in at least half the patients, we observed convergent clusters in the left supramarginal gyrus and the right DLPFC (Fig. S1d). It is possible that the neuroimaging effects of antidepressants vary across individual patients, and in turn, relate to their variability in response to treatment. Further original and meta-analytic neuroimaging studies are needed to evaluate the

individual variability of treatment-induced changes in brain function and its relevance to clinical response.

## Conclusion

This comprehensive meta-analysis of the functional neuroimaging studies on the regional and network-level convergence of the effects of antidepressant medications in MDD underscores the importance of the FPN (and particularly DLPFC) and SAN in the pharmacotherapy of MDD. This observation may be attributed to the key roles of these regions in executive functions and emotional processing, which may transcend to other therapeutic approaches. In particular, the convergent connectivity map of antidepressant effects engaged a circuit similar to the circuits of TMS targets associated with the improvement of dysphoric symptoms. This may hint at symptoms-specific effects of antidepressants that need to be further investigated in the future. Notably, we identified no associations between our regional and network-level meta-analytic findings with the spatial maps of neurotransmitter receptors/transporters. We highlight the need for future research integrating the multiple levels of antidepressant actions at the micro- and macroscale.

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## Conflicts of Interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

## 585 Data Availability

586 The coordinates of the foci reported in the included experiments are available in  
587 <https://doi.org/10.6084/m9.figshare.24592539>. The group-averaged dense resting-state  
588 functional connectivity matrix from the Human Connectome Project can be accessed at  
589 <https://db.humanconnectome.org>.

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## Figure Legends

### **Fig. 1. Study selection flowchart.**

MDD: major depressive disorder, LLD: late-life depression, ROI: region of interest, SVC: small volume correction.

### **Fig. 2. Treatment-induced increase of voxel-based physiology in the left middle frontal gyrus.**

A. Peak coordinates of the included experiments in Treated > Untreated (red) and Untreated > Treated (blue) comparisons. Each dot represents a peak coordinate. B. Activation likelihood estimation showed significant convergence of Treated>Untreated comparisons in the left dorsolateral prefrontal cortex (DLPFC) after family-wise error correction at cluster level.

### **Fig. 3. Connectivity mapping of the left dorsolateral prefrontal cortex cluster.**

Using the center of convergent cluster at the left dorsolateral prefrontal cortex as the seed (outlined patch), the meta-analytical co-activation (A) and resting-state functional connectivity (B) maps are shown.

### **Fig. 4. Network-level convergence of antidepressant effects.**

A. The cortical and subcortical map (left) represent z-scored convergent connectivity map of the foci from all experiments. The distribution of convergent connectivity map across the resting-state networks (RSNs) is shown (right). Asterisk denotes networks with mean convergent connectivity significantly more extreme than a null distribution based on surrogate spun maps. B. The radar plot (left) shows the number of foci in each RSN. Asterisk denotes networks with statistically significant counts of observed foci compared to a null based on random foci.

VIS: visual network, SMN: somatomotor network, DAN: dorsal attention network, SAN: salience network, LIM: limbic network, FPN: frontoparietal network, DMN: default mode network.

### **Fig. 5. Association of meta-analytic findings with neurotransmitter receptor/transporter maps.**

A. The parcellated and z-scored PET maps of neurotransmitter receptor/transporter (NRT) are shown. Red outline indicates the left dorsolateral prefrontal cortex (L DLPFC) convergent cluster. B. Median rank-normalized density of NRTs in L DLPFC cluster. C. Correlation of parcellated convergent connectivity map with the PET maps.

1052 Table 1. Characteristics of studies included in this meta-analysis

#	First Author, Year <sup>a</sup>	N (female ) Treated / Untreated	Responded %	Age Treated / Untreated	Wash out period	Antidepressant medi- cation	Duration between Scans	Modality
1	Abdallah, C. G., 2017	18 (44%)	55.5%	43	1 w	Ketamine	1 d	rs-fMRI (GBC)
	Murrough, J. W., 2015							tb-fMRI
2	Bremner, J. D., 2007	13 (84%)	100%	40.0 <sup>c</sup>	4 w	Fluoxetine or Venla- faxine	6 +/- 3 m	H <sub>2</sub> O-PET
3	Carlson, P. J., 2013	20 (30%)	30.0%	48	2 w	Ketamine	1-3 d	FDG-PET
4	Cheng, Y., 2017	38 (70%)	60%	28 <sup>c</sup>	Drug-naive	Escitalopram	5 h, 4 w, 8 w	rs-fMRI (fALFF)
5	Downey, D., 2016	21 (62%) / 19 (58%)	NR	27.1 / 25.7	NR	Ketamine	45 min	rs-fMRI
6	Fonzo, G. A., 2019	96 (72%) / 105 (64%)	NR	37 / 36	NR	Sertraline	8 w	tb-fMRI
7	Frodl, T., 2011	24 (33%)	NR	39	1 y	Venlafaxine or Mirtazapine	4 w	tb-fMRI
8	Fu, C. H., 2004, 2007	19 (68%)	NR	43	NR	Fluoxetine	8 w	tb-fMRI
9	Fu, C. H., 2015	24 (41%)	79%	40.2	4 weeks	Duloxetine	12 w	tb-fMRI
10	Gonzalez, S., 2020	11 (27%)	45.4%	48	None	Ketamine	1 h, 6 h, 24 h	ASL-fMRI
11	Jiang, W., 2012	21 (57%)	100%	29	Drug-naive	Escitalopram	8-12 w	tb-fMRI
12	Joe, A.Y., 2006	35 (72%)	53.8%	45.3	NR	Citalopram	3 w	<sup>99m</sup> Tc- HMPAO SPECT
13	Keedwell, P., 2008	12 (50%)	66%	49.0	Drug-naive	Variable	6 – 18 w	tb-fMRI

14	Kennedy, S. H., 2001	13 (0%)	100%	37	4 w	Paroxetine	6 w	FDG-PET
15	Kohn, Y., 2008	11 (54%)	100%	49	Variable	Paroxetine, Fluoxetine or Clomipramine	2 y	<sup>99m</sup> Tc-HMPAO SPECT
16	Komulainen, E., 2018	17 (53%) / 15 (60%)	NR	27 / 23	4 m	Escitalopram	1 w	tb-fMRI
	Komulainen, E., 2021	15 (53%) / 14 (57%)	NR	29 / 24	NR			
17	Kraus, C., 2019	26 (73%) / 36 (63%)	84%	30.4 / 28.5	3 m	Escitalopram ± Venlafaxine or Mirtazapine	12 w	tb-fMRI
	Rütgen, M., 2019	29 (72%)	75.8%	30	3 m			
18	Li, C. T., 2016	32 (69%)	31.2%	44	NR	Ketamine	1 h	FDG-PET
19	Lopez-Sola, M., 2010	13 (85%)	69.2%	45	15 d	Duloxetine	1 w, 8 w	tb-fMRI
20	Mayberg, H. S., 2000	10 (0%)	50.0%	49 <sup>c</sup>	1 m	Fluoxetine	1 w, 6 w	FDG-PET
21	Reed, J. L., 2018, 2019	28 (64%)	NR	36 <sup>c</sup>	2 w	Ketamine	2d, 11d	tb-fMRI
22	Robertson, B., 2007	8 (75%)	75%	41 <sup>c</sup>	NR	Bupropion	8 w	tb-fMRI
23	Sankar, A., 2017	23 (56%)	78.2%	40	4 w	Duloxetine	12 w	tb-fMRI
24	Sterpenich, V., 2019	10 (60%)	NR	51	None	Ketamine	1 d, 7 d	tb-fMRI
25	Vlassenko, A., 2004	14 (57%)	100%	43.1	3 w	Fluoxetine, Paroxetine or Amesergide	12 w	<sup>99m</sup> Tc-HMPAO SPECT
26	Wagner, G., 2010	20 (90%)	50.0%	39	Variable	Citalopram or Reboxetine	6 w	tb-fMRI
27	Walsh, N. D., 2007	20 (70%)	75.0%	44	NR	Fluoxetine	8 w	tb-fMRI
28	Wang, L., 2014	14 (36%)	100%	33	Drug-naive	Escitalopram	8 w	rs-fMRI (ReHo)

	Wang, L., 2017	20 (55%)	100%	35				rs-fMRI (fALFF)
29	Wang, Y., 2012	18 (61%)	NR	32	Drug-naive	Fluoxetine	8 w	tb-fMRI
30	Williams, R. J., 2021	38 (68%)	57% (after 8 weeks)	36.2	NR	Citalopram or Quetiapine	1 w	tb-fMRI
31	Yin, Y., 2018	11 (36%)	100%	49.2	4 w	Variable	8 w	ASL-fMRI

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1054 <sup>a</sup>Publications with overlapping samples are grouped together, <sup>b</sup> Mean or median, <sup>c</sup> Reported for all the subjects rather than those in the includ-  
1055 ed experiment

1056 NR: not reported, SSRI: selective serotonin reuptake inhibitor, SNRI: serotonin-norepinephrine reuptake inhibitor, NRI: norepinephrine reuptake inhibitor,  
1057 NMDAR: N-methyl-D-aspartate receptor, NDRI: fMRI: functional magnetic resonance imaging, rs-fMRI: resting state fMRI, tb-fMRI: task based fMRI, ASL-  
1058 fMRI: arterial spin labeling fMRI, FDG-PET: fluorodeoxyglucose-positron emission tomography, SPECT: single-photon emission computed tomography,  
1059 GBC: global brain connectivity, fALFF: fractional amplitude of low-frequency fluctuations, ReHo: regional homogeneity, ASL: arterial spin labeling.

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1061 **Table 2.** Activation Likelihood Estimation (ALE) analyses on the effects of antidepressants in major depressive disorder

Experiments	Comparison	N	Min $p_{cFWE}^a$	Convergence
All	All	31	0.402	-
	Treated > Untreated	21	<b>0.007</b>	L DLPFC
	Treated < Untreated	22	0.624	-
<i>Based on modality</i>				
Rest	All	13	0.056	-
Task	All	19	0.970	-
<i>Based on treatment and clinical setting</i>				
Excluding ketamine	All	24	0.505	-
	Treated > Untreated	15	<b>0.024</b>	L DLPFC, R DLPFC
	Treated < Untreated	17	0.475	-
Pre- versus post-treatment effects	All	25	0.323	-
	Treated > Untreated	19	<b>0.005</b>	L DLPFC, R DLPFC
	Treated < Untreated	21	0.482	-
Treatment duration > 4 weeks	All	21	0.352	-
	Treated > Untreated	13	<b>0.014</b>	R mSFG
	Treated < Untreated	15	0.197	-
Response in •50% of subjects	All	20	0.636	-
	Treated > Untreated	13	<b>0.009</b>	L SMG, R DLPFC
	Treated < Untreated	16	0.371	-

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1063 <sup>a</sup> Bold p-values indicate statistical significance.

1064 cFWE: cluster-wise family-wise error, SSRI: selective serotonin reuptake inhibitor, L: left, R: right, DLPFC: dorsolateral prefrontal cortex, mSFG: medial supe-  
 1065 rior frontal gyrus, SMG: supramarginal gyrus.

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1068 **Table 3.** Comparison of the existing coordinate-based neuroimaging meta-analyses on the brain effects of antidepressants.

CBMA	N studies <sup>a</sup>	Treatment	Imaging	Method	Findings
Fitzgerald et al., 2008	9	SSRI	PET, SPECT (resting state)	ALE, FDR-corrected, BrainMap	<p>↑: middle frontal gyrus, inferior frontal gyrus, anterior cingulate cortex, precentral gyrus, supramarginal gyrus, posterior cingulate gyrus, inferior parietal lobe, mid-brain, putamen</p> <p>↓: middle and superior frontal gyri, medial frontal gyrus, subgenual and pregenual anterior cingulate, parahippocampal gyrus, hippocampus, insula, putamen</p>
Delaveau et al., 2011	9	Antidepressants	fMRI, PET (emotional activation)	ALE, FDR-corrected, GingerALE 2.0	<p>↑: dorsal medial prefrontal cortex, dorsolateral prefrontal cortex, cuneus, fusiform gyrus, lingual gyrus, middle temporal gyrus, putamen, caudate, thalamus, anterior insula, anterior cingulate cortex</p> <p>↓: thalamus, caudate, putamen, globus pallidus, hippocampus, amygdala, parahippocampal gyrus, anterior insula, anterior cingulate cortex, orbitofrontal cortex, posterior cingulate cortex, middle frontal gyrus,</p>

					pre/postcentral gyri, lingual gyrus, cerebellum, supramarginal gyrus, fusiform gyrus
Graham et al., 2013	4	Any	fMRI	ALE / GPR, FDR-corrected, GingerALE 2.1 / custom code	↑: - (ALE), precentral gyrus, precuneus, dorsolateral prefrontal cortex (GPR) ↓: superior temporal gyrus, cerebellum (ALE), precuneus, dorsal lateral prefrontal cortex, lateral occipital region (GPR)
Ma, 2015 <sup>b</sup>	22	SSRI, SNRI	fMRI (emotional processing)	ALE, FDR-corrected, GingerALE 2.3	↑: dorsolateral prefrontal cortex (negative emotions) ↓: amygdala, hypothalamus, putamen, middle temporal gyrus, ventromedial prefrontal cortex, posterior insula, middle frontal gyrus (negative emotions) ↑↓: amygdala, dorsolateral prefrontal cortex, hippocampus, ventromedial prefrontal cortex, anterior cingulate cortex, fusiform, anterior insula, precuneus (positive emotions)
Boccia et al., 2016	12	Antidepressants, Psychotherapy	fMRI	ALE, FDR-corrected, GingerALE 2.1	↑↓: insula, anterior cingulate cortex, precentral and postcentral gyri, middle frontal gyrus, precuneus, basal ganglia, putamen, cerebellum
Chau et al., 2017	7	SSRI, TMS, ECT	PET, SPECT, ASL-fMRI (resting	MLKD, cFWE-corrected	↑: - ↓: anterior insula

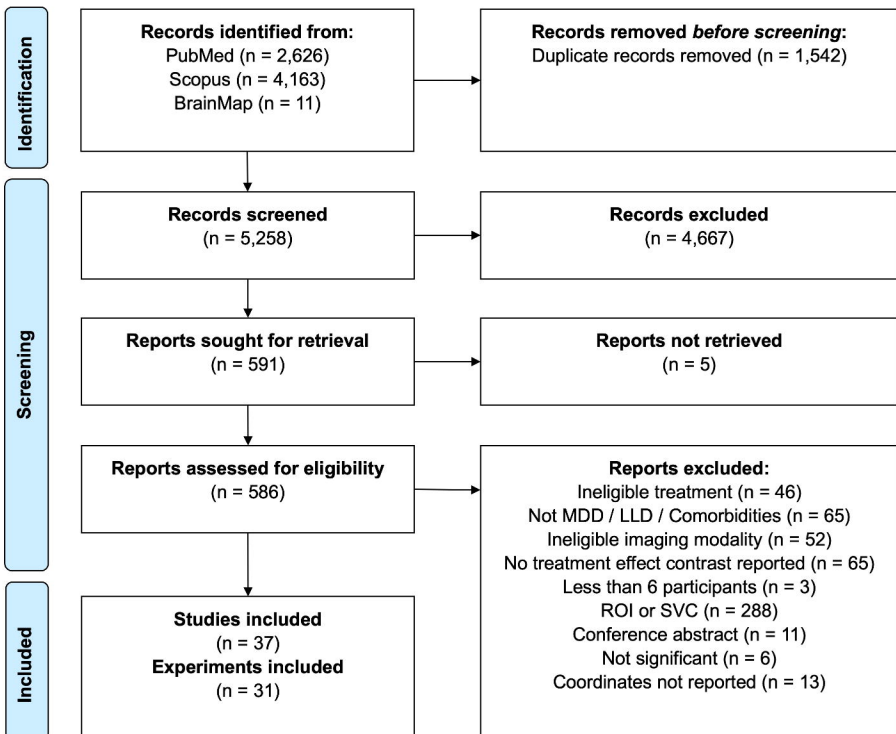
state)					
Li et al., 2022 <sup>c</sup>	33	Antidepressants (incl. ketamine), CBT, ECT	fMRI, PET, VBM	ALE, cFWE-corrected, GingerALE 3.0	↑: amygdala, parahippocampal gyrus, thalamus ↑↓: amygdala, parahippocampal gyrus, thalamus, anterior cingulate cortex, middle frontal gyrus, insula, claustrum
Current paper	37	Antidepressants (incl. ketamine)	fMRI, PET, SPECT	ALE, cFWE-corrected, pyALE	↑: dorsolateral prefrontal cortex ↓: - ↑↓: -

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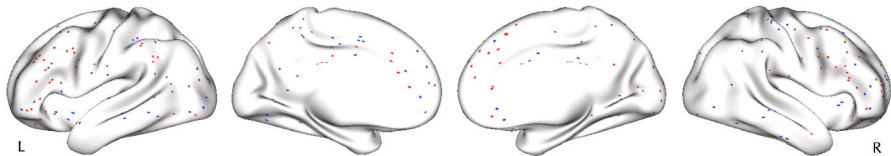
1070 <sup>a</sup> Only includes number of studies on the effects of pharmacotherapy on MDD, excluding other treatments and disorders, <sup>b</sup> Reported results indicate antidepressant effects in mood disorders including depression and anxiety. Results specific to MDD were not reported. <sup>c</sup> Reported results indicate effects of all treatments. Specific effects of antidepressant medications were not reported.

1073 ↑: increased imaging measures, ↓: decreased imaging measures, ↑↓: changed imaging measure in any direction, SSRI: selective serotonin reuptake inhibitor, SNRI: serotonin-norepinephrine reuptake inhibitor, TMS: transcranial magnetic stimulation, CBT: cognitive behavioral therapy ECT: electroconvulsive therapy, PET: positron emission tomography, SPECT: single-photon emission computed tomography, fMRI: functional magnetic resonance imaging, ASL: arterial spin labeling, VBM: voxel-based morphometry, ALE: activation likelihood estimation, FDR: false discovery rate, MKLD: cFWE: cluster-level family-wise error correction, multilevel kernel density.

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## A. Peak coordinates of antidepressant effects



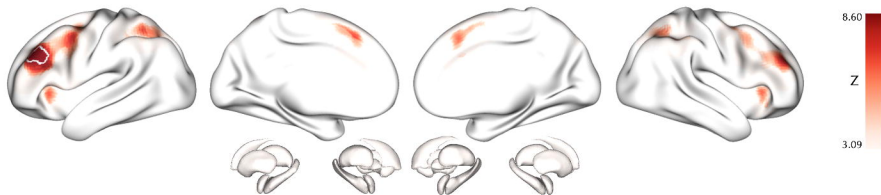
## B. Activation likelihood estimation (Treated > Untreated)



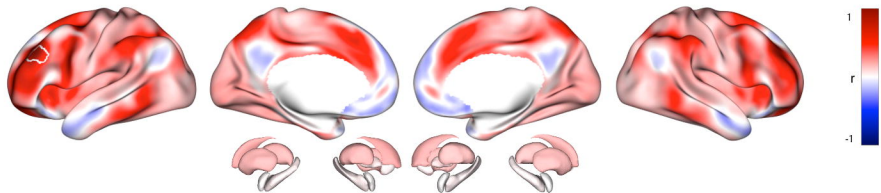
■ Treated > Untreated  
■ Untreated > Treated



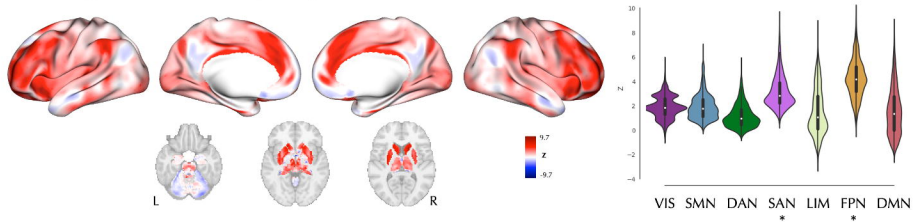
## A. Meta-analytical coactivation



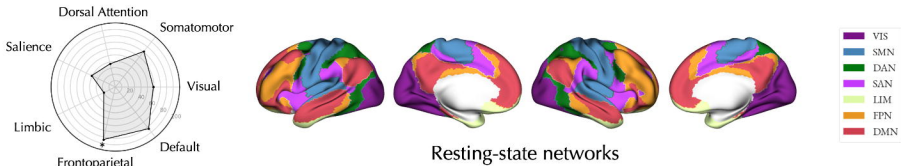
## B. Resting-state functional connectivity



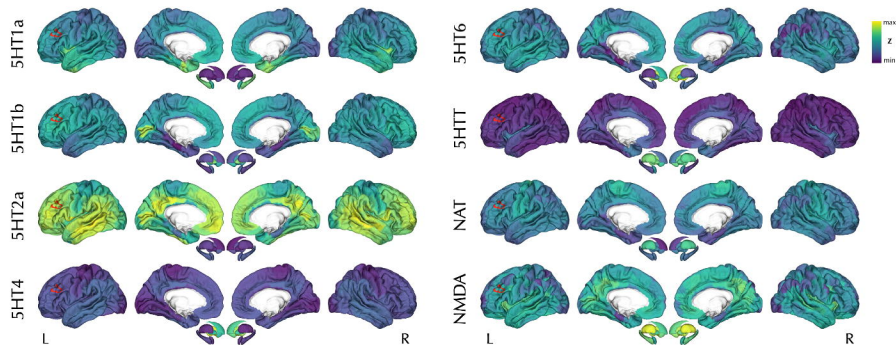
## A. Convergent connectivity map of foci



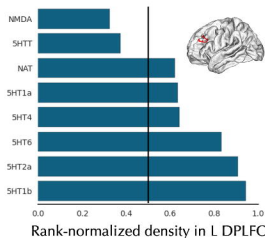
## B. Foci distribution across resting-state networks



## A. Receptor/transporter density PET maps



## B. Receptor/transporter density in L DLPFC



## C. Correlation of PET maps with convergent connectivity map

