Revised: 14 April 2023

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# Significant heterogeneity in structural asymmetry of the habenula in the human brain: A systematic review and metaanalysis

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#### Funding information

National Natural Science Foundation of China, Grant/Award Number: 32171031; STI 2030-Major Projects, Grant/Award Number: 2021ZD0200409; Fundamental Research Funds for the Central Universities, Grant/Award Number: 2021XZZX006; Information Technology Center of Zhejiang University; U.S. National Institutes of Health, Grant/Award Number: R01 MH116147

#### Abstract

Understanding the evolutionarily conserved feature of functional laterality in the habenula has been attracting attention due to its potential role in human cognition and neuropsychiatric disorders. Deciphering the structure of the human habenula remains to be challenging, which resulted in inconsistent findings for brain disorders. Here, we present a large-scale meta-analysis of the left-right differences in the habenular volume in the human brain to provide a clearer picture of the habenular asymmetry. We searched PubMed, Web of Science, and Google Scholar for articles that reported volume data of the bilateral habenula in the human brain, and assessed the left-right differences. We also assessed the potential effects of several moderating variables including the mean age of the participants, magnetic field strengths of the scanners and different disorders by using meta-regression and subgroup analysis. In total 52 datasets (N = 1427) were identified and showed significant heterogeneity in the left-right differences and the unilateral volume per se. Moderator analyses suggested that such heterogeneity was mainly due to different MRI scanners and segmentation approaches used. While inversed asymmetry patterns were suggested in patients with depression (leftward) and schizophrenia (rightward), no significant disorder-related differences relative to healthy controls were found in either the left-right asymmetry or the unilateral volume. This study provides useful data for future studies of brain imaging and methodological developments related to precision habenula measurements, and also helps to further understand potential roles of the habenula in various disorders.

#### KEYWORDS

brain asymmetry, brain disorders, habenula, heterogeneity, laterality

### 1 | INTRODUCTION

Understanding the hemispheric specialization of the human brain is one of the long-standing questions in human brain research (Duboc et al., 2015; Kong, Postema, et al., 2022; Toga & Thompson, 2003). Altered asymmetry has been suggested as a potential biomarker in various cognitive and neuropsychiatric disorders, including dyslexia, Alzheimer disease, autism, and obsessive-compulsive disorder (Eyler et al., 2012; Kim et al., 2012; Leonard & Eckert, 2008; Menzies et al., 2008). Only recently, have large-scale neuroimaging studies

This is an open access article under the terms of the Creative Commons Attribution-NonCommercial License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited and is not used for commercial purposes. © 2023 The Authors. *Human Brain Mapping* published by Wiley Periodicals LLC. been conducted aiming to provide a definitive and normative reference of brain asymmetry in healthy and diseased populations (Kong, Postema, et al., 2022; Kong, ENIGMA Laterality Wroking Group, et al., 2022; Ocklenburg et al., 2021). Thus far, existing large-scale studies either focused on mapping of the cortex or relatively large subcortical structures (e.g., hippocampus and thalamus), and the results suggest that such structural asymmetries could explain only a fraction of the variability in functional lateralization (e.g., Dorsaint-Pierre et al., 2006; Guadalupe et al., 2014; Kong et al., 2021). Yet little is known about the nature of the asymmetry for smaller subcortical structures such as the habenula (whose volume is ~15–30 mm<sup>3</sup> in each hemisphere).

The habenula is a bilateral nuclear complex located on the dorsal tip of the thalamus, which is phylogenetically conserved from fish to human (Amo et al., 2010). It has attracted much interest due to its potential role in the evolution of the brain and in its functions. This structure is considered to be essential for allowing animals to switch responses as needed when environments and motivational stages change (Mizumori & Baker, 2017). The habenular asymmetries are widespread amongst vertebrates and have been repeatedly

characterized regarding size, shape, neuronal organization, and connectivity, but the laterality (in terms of amount and directions) of asymmetry is variable between species (e.g., hagfishes: larger size on the right; cartilaginous fishes: larger size on the left; frogs: larger size on the left; Agetsuma et al., 2010; Concha & Wilson, 2001; Wree et al., 1981). The asymmetry has also been associated with various functions such as sensory processing, fear responses, and other behaviors in different animals (e.g., zebrafish: leftward processing of light, rightward processing of odor, and correlates of the left with fear responses; Dreosti et al., 2014; Duboué et al., 2017; Guglielmotti & Fiorino, 1998). However, the functional role and the nature of the asymmetry of the habenula in humans remains largely unexplored (Batalla et al., 2017), in part, because of the extremely small size of the habenula and its low anatomical contrast on in vivo human brain imaging (Figure 1).

Understanding this evolutionarily conserved brain feature has been attracting attention in different research areas including neuroscience, psychology and psychiatry, due to its potential role in human cognition and a variety of neuropsychiatric disorders such as



**FIGURE 1** Anatomical location of the habenula in the human brain (top left) and a summary of related studies in the past decades (top right and bottom). The habenula is highlighted with the red box and arrows. Data for the visualization was obtained via Web of Science using search terms "habenula (Title) or habenular (Title)." The bottom panel indicates the top research areas of the 1550 relevant publications obtained, including Neuroscience/Neurology, Behavioral Sciences, Psychology, and Psychiatry. Depression and Schizophrenia are the top brain disorders revealed in the MeSH Headings from Web of Science. For a detailed description of the anatomical location of the habenula in the human brain, please refer to Germann et al. (2021).

depression and schizophrenia (see Figure 1). Some of the leading efforts to understand habenular asymmetry have used high-resolution functional magnetic resonance imaging (fMRI; Lawson et al., 2013). For example, functional activation in the habenula has been found during the prediction of aversive stimuli (Lawson et al., 2017), which could be, in part, explained by its role in avoidance learning processing (Yoshino et al., 2020). Interestingly, researchers found that the left habenula, compared with the right, seems to be more significantly activated during punishment outcomes, and this activation correlated with higher reward dependence scores and more severe depressive symptoms (Liu et al., 2017; Yoshino et al., 2020). Asymmetric activation of the habenula has also been reported in studies of aversive processing (Hennigan et al., 2015; Lawson et al., 2014). Further evidence in supporting the functional differentiation of the left and right habenula comes from functional connectivity studies. For example, restingstate activity recorded from the left and right habenula showed very low correlation (r < .10), suggesting a lack of functional connectivity between the bilateral habenula (Hétu et al., 2016). Distinct functional connectivity patterns of the left and right habenula with other subcortical and cortical regions have been repeatedly reported with data of variable quality (Erpelding et al., 2014; Hétu et al., 2016; Torrisi et al., 2017). Taken together, these findings support the functional laterality of the habenula in the human brain.

Regarding structural asymmetry of the habenula, only a few studies have been published, and these studies usually showed mixed results. For example, Ahumada-Galleguillos et al. (2017) found greater volume of the left habenula, in line with findings of multiple studies (e.g., Savitz, Nugent, et al., 2011; Torrisi et al., 2017). However, other studies had inconsistent results: some studies did not find any interhemispheric differences in habenula volume (e.g., Hétu et al., 2016; Lawson et al., 2013), while others even suggested reversed asymmetry patterns (i.e., rightward; e.g., Germann et al., 2020). Thus, we could not have a clear hypothesis regarding the direction of the asymmetry. In addition, these mixed results on habenular asymmetry may reflect differences in many factors, including sample heterogeneity such as age and sex distributions and sample sizes, as well as differences in brain data types (e.g., in vivo brain imaging vs. brain tissues), magnetic field strengths of the scanners (e.g., 3T vs. 7T), segmentation approaches (e.g., manual vs. automated), as we have observed in our large-scale cortical and subcortical asymmetry studies (Guadalupe et al., 2017; Kong et al., 2018; Kong, Postema, et al., 2022; Ocklenburg & Güntürkün, 2018). Thus, a large-scale survey using metaanalysis approaches would be beneficial, to provide a clearer picture of the habenular asymmetry.

Moreover, altered asymmetry of the habenula has been suggested in various disorders. For example, a study using postmortem brain tissues from two groups of subjects with major and bipolar depression indicates the presence of asymmetric changes (a rightsided decrease) in habenular structure (Ranft et al., 2010). However, in some more recent studies, no significant differences were found in the volume of either the left or the right habenula (Luan et al., 2019; Schmidt et al., 2016). The left habenula showed a tendency toward lower volume in patients with schizophrenia (Zhang et al., 2017), while such differences were not found in another study (Ranft et al., 2010). As the results are mixed, a meta-analysis of existing data would be a major step forward in achieving a more accurate description of the variations related to these disorders.

In this study, we present a large-scale analysis of structural asymmetry of the habenula in the human brain, with the bilateral volume data from 52 datasets comprising 1427 subjects, using meta-analytic methods. Our aim was to examine whether—and to what extent—the habenula shows left-right differences in volume, and to provide informative data on the habenular asymmetry with the largest sample size to date. We also estimated inter-dataset variability in the asymmetry effects, and assessed potential influences of age and sex on the variability in the asymmetry, as well as methodological factors such as MRI scanner field strength and the segmentation approaches used. Finally, we compared data on bilateral habenular volume and the leftright differences between patient datasets and controls, to test potential disorder-related effects.

### 2 | MATERIALS AND METHODS

#### 2.1 | Data sources

We searched PubMed, Web of Science, and Google Scholar for the articles that reported volume data on the bilateral habenula in the human brain, following PRISMA guidelines (see the PRISMA checklist in Data S1; Page et al., 2021). Specifically, the literature search was conducted on April 19, 2022, using the key terms: ("habenula volume\*" OR "habenular volume\*") AND (human) AND ("magnetic resonance imaging" OR "MRI" OR "tissue"). Records covered the years from 1986 to 2022.

#### 2.2 | Inclusion criteria and study selection

Studies were included based on two criteria: (1) habenula volumes were examined in human samples; (2) bilateral habenula volume data were both reported. To screen the articles that might be relevant for the present study, we first removed duplicate records using EndNote, and two authors (Y.A. and X.K.) identified irrelevant entries by screening the titles and abstracts, and further retrieved the full-text of the remaining articles for data extraction (see below). Articles that did not report bilateral habenula volume data were excluded in the following analyses. Reference lists were also screened for additional references.

### 2.3 | Data extraction

From the selected articles, we extracted data on the mean and standard deviation (SD) of the left and right habenula volumes, various information on the samples (i.e., the sample sizes, mean ages and sex 4\_\_\_\_WILEY\_

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distributions of the samples [sex proportion of the sample, %male], and the sample category [e.g., healthy controls and major depressive disorder patients], and the measurement and segmentation approaches used). Note that healthy controls might include samples with other diseases that are not related to cerebral illness or neuropathy. For the majority of the articles, the data were accessible from the article text. In a few cases where the volume data was only presented within figures, we contacted the corresponding authors to request the exact numbers. When possible (Germann et al., 2020; Lawson et al., 2017; Liu et al., 2017), we extracted the volume data from these figures using an automatic tool, WebPlotDigitzer. Data from each study were double-extracted to prevent transcription errors. In addition, a study by Ranft et al. (2010) only reported the volume of the habenula subregions (i.e., the lateral and medial habenula). In this case, we combined the subregional data as an estimate of the whole habenula volume. When data were not accessible via any of these approaches, the studies were excluded from the follow-up statistical analyses.

During data extraction, we only kept one version of the data when datasets were identified as from the same or overlapping samples and were assessed using the same measurement approaches. Data from the overlapping samples but assessed using different measurement approaches were kept in the analyses for examining the effects of measurement approaches. For instance, we identified that the healthy individuals and the measurement approach used were identical in Savitz, Bonne, et al. (2011) and Savitz, Nugent, et al. (2011), so we only used that data once in the meta-analyses.

#### 2.4 **Bias assessment**

Quality of meta-analysis is usually affected by potential publication bias and the quality of the studies included. Publication bias refers to the fact that studies that fail to find significant results usually could be not published, and could have no chance to be included in a following meta-analysis; the quality of studies that are included often depends on risk bias assessment, which refers to systematic errors caused by the study design, data analysis and/or other factors. Publication bias would not be a problem for the present study as majority of the studies included in the meta-analysis focused on the structural variation of the habenula per se and did not report any results on the asymmetry of the habenular volume. Regarding the risk of bias assessment, it is actually part of the research question of this present study. Specifically, we estimated the potential effects of sample variables including age and sex, magnetic field strength of MRI scanners, and segmentation approaches used.

#### 2.5 Meta-analyses

We used mean difference (MD) and pooled standard deviation (SD<sub>pooled</sub>) derived from a dataset to assess the inter-hemispheric

difference between the left and right habenula. Specifically, first the MD and pooled SD were calculated by following Equations (1) and (2) below, respectively. In the formula for calculating the pooled SD of each study, the correlation coefficient between the bilateral habenula volumes was taken into account for a closer estimation of the pooled SD. While such a correlation was not reported in the majority of these studies, we repeatedly used a range of various correlation values from 0.30 to 0.80 with a step size of 0.10 as suggested in recently study with a relatively large sample size (N > 100; Germann et al., 2020). These derived measures were fed into a meta-analysis for estimating the left-right differences, which represent the effect sizes used in this study. Also, we visualized meta-analytic results using forest plots where sample size, effect size, 95% confidence interval (CI), and the weight of each study (determined by CI) were shown.

$$MD = Mean_L - Mean_R$$
(1)

$$SD_{pooled} = \frac{\sqrt{\left(SD_{L}^{2} + SD_{R}^{2} - 2 * r * SD_{L} * SD_{R}\right)}}{\sqrt{N}}$$
(2)

While the main analyses were based on data from all datasets, we repeated the analysis for the datasets from healthy controls, to reduce the heterogeneity that might be related to brain disorders. We also ran the same meta-analyses for the datasets from patients with one of the specific disorders included (i.e., major depressive disorders [MDD], schizophrenia [SCZ], and bipolar disorder [BD]).

Heterogeneity was tested using Cochran Q statistic, and quantified by the  $l^2$  statistic and  $\tau^2$  statistic, which was calculated by the restricted maximum likelihood estimator (Viechtbauer, 2005). We also used Knapp-Hartung adjustments (Knapp & Hartung, 2003) to calculate the CI around the pooled effect, to reduce the chance of false positives.

Given the similar results even with different correlations, in the main test, we focused on the results with a correlation of 0.50 (also discuss the results with other correlation values in the sensitivity analyses).

#### 2.6 Moderator analyses with meta-regression and subgroup analysis

To further address the source of heterogeneity in the meta-analyses, we investigated the potential effects of several moderating variables regarding the samples of each dataset, that is, the sex ratio (%male) and mean age of the samples. In one case (Ahumada-Galleguillos et al., 2017), sex and age information were only available for the overall samples but not the subgroups, and thus the combined effect statistics were calculated. Other moderating factors were also considered, including brain data type (i.e., MRI images of post-mortem brain tissues), magnetic field strengths of the MRI scanners (i.e., 1.5, 3, and 7T), and the segmentation approaches used (i.e., manual, fully automated, or semi-automated). In addition, we investigated a



FIGURE 2 Flowchart of study selection.

potential moderating effect of the average volume of bilateral habenula on the left-right differences. Meta-regression mixed effects models and subgroup analyses were used for the moderator analyses of continuous and categorical variables, respectively. When information for these variables were not available in some cases, the datasets were excluded from the corresponding analyses.

### 2.7 | Sensitivity analyses

The estimation of the pooled SD largely depends on the correlation coefficient of the left and right habenula volume, which in turn could impact the meta-analysis results. Thus, we repeated the analyses with a range of correlations (from 0.3 to 0.8), and reported the results in each case. In addition, including results based on too few participants may reduce reliability, so we re-ran the main analyses on datasets with a sample size of larger than 15.

#### 2.8 | Software used

We used JASP (version 0.16) and R (version 4.2.0) for basic statistical analyses, and R package *meta* (version 5.2-0) for the meta-analyses, subgroup analysis, and meta-regressions.

## 3 | RESULTS

#### 3.1 | Study inclusion and data extraction

In total, we identified 244 records: 8, 12, and 222 from PubMed, Web of Science, and Google Scholar, respectively, and 2 additional records from reference lists (Figure 2). After excluding duplicate records, 205 articles were left. Two authors (Y.A. and X.K.) read the abstract or full text as required, and removed 183 articles that did not report bilateral volume data for the habenula. One additional article was

removed because of the duplicate sample issue that we identified (see Section 2). Thus, 21 articles remained and were included in the main meta-analysis (Table 1).

Data on sample categories, sample size, and the mean and standard deviation of the bilateral habenula volume were extracted from each article (see Section 2). This resulted in 52 separate datasets. These samples covered healthy individuals as well as patients with various diseases and conditions, including MDD, BD, and SCZ. The sample sizes of the extracted volume data were up to 75 (mean = 27; median = 22; in total N = 1427). Measurement approaches included brain imaging with 3 or 7T MRI scanners, and image segmentation of brain tissues.

# 3.2 | Relationship between the volumes of the left and right habenula

The mean volume varied, ranging from 11.61 to 39.40 mm<sup>3</sup>, and from 11.21 to 35.65 mm<sup>3</sup> for the left and right habenula, respectively, suggesting considerable variability of the habenula volume reported in the literature (Figure 3). As expected, the left and right mean volume at the study level showed a high correlation when correlation was assessed across all the studies (r = .960, p < .001, number of datasets [k] = 52), as well as the various subgroups of healthy samples (r = .967, p < .001, k = 25), and patients with MDD (r = .957, p < .001, k = 11), SCZ (r = .960, p = .010, k = 5), but the correlation was relatively low in the BD patients (r = .361, p = .550, k = 5). These results suggested that little, if any, inter-hemispheric differences would exist in the habenula volume.

An initial synthesis of the volume data from the left and right habenula using a paired samples *t*-test also supported this point: no significant interhemispheric differences for either the healthy samples ( $t_{(24)} = 1.443$ , p = .162, Cohen d = 0.289), or the patients with SCZ ( $t_{(4)} = -0.550$ , p = .612, d = -0.246); there appeared to be a nominally significant difference in MDD subgroup ( $t_{(10)} = 2.403$ , p = .037, d = 0.725), but it did not survive correction of multiple testing. In the BD subgroup, no significant differences ( $t_{(4)} = -0.352$ , p = .743, d = -0.157) were observed, perhaps due to the limited number of studies. Note that this initial synthesis analysis was conducted at the study level, and potential individuallevel variance and important variations across studies (e.g., sample size) were not considered. To further confirm these findings, a more formal synthesis was conducted using a meta-analysis approach below.

## 3.3 | Inter-hemispheric differences in the habenula volume

#### 3.3.1 | Meta-analysis of all datasets

To take into account individual-level variances and the various samples of these studies, we performed a meta-analysis by integrating the MD and pooled SD data. The habenula showed positive MDs in 30 datasets (maximum MD: 3.90 mm<sup>3</sup>; Left > Right; N = 720), and negative MDs in the remaining 22 datasets (MD of the maximum difference:

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 $-3.88 \text{ mm}^3$ ; Left < Right; N = 707). The meta-analysis of the left-right differences in all the datasets showed an average difference of 0.41 mm<sup>3</sup>, with a 95% CI of -0.13 to 0.94 mm<sup>3</sup>. This observed difference was not significant with a significance threshold of p < .05 ( $t_{(51)} = 1.52$ , p = .134) (Figure 4). While publication bias could not be a problem for the present study as stated in the *Bias assessment*, we ran a statistical test for funnel-plot asymmetry in main meta-analysis. As expected, we found no significant effect (metabias Z = 1.26, p = .21) which indicated no publication bias related to left-right differences.

#### 3.3.2 | Meta-analysis of different groups of patients

We also ran meta-analyses of the inter-hemispheric differences in the volumes for the datasets of healthy samples and each patient group (i.e., patients with MDD, SCZ, and BD; Figure 5). In line with the overall findings above, no differences were found in either of these groups: healthy (MD = 0.40 mm<sup>3</sup>; 95% Cl: -0.37 to 1.16 mm<sup>3</sup>;  $t_{(24)} = 1.07$ , p = .294), SCZ (MD = -0.54 mm<sup>3</sup>; 95% Cl: -3.44 to 2.36 mm<sup>3</sup>;  $t_{(4)} < 1.00$ ), BD (MD = -0.54 mm<sup>3</sup>; 95% Cl: -3.44 to 2.85 mm<sup>3</sup>;  $t_{(4)} < 1.00$ ). Interestingly, the meta-analysis also showed a significant left-right difference in the MDD samples (MD = 1.23 mm<sup>3</sup>; 95% Cl: 0.14-2.33 mm<sup>3</sup>; Left > Right:  $t_{(10)} = 2.51$ , p = .031), which is similar to that of the initial comparison above. We did not find any significant differences in the left-right difference of the habenula volume between either of the disorder groups and the healthy control samples (ps > .05).

#### 3.4 | Moderator analyses

Moderate to substantial heterogeneity in the MD was observed across the datasets. The heterogeneity remained when analyzing only datasets from either the healthy controls ( $l^2 = 87\%$ ,  $\tau^2 = 2.756$ , p < .01), or specific disorder samples (MDD:  $l^2 = 80\%$ ,  $\tau^2 = 2.056$ , p < .01; SCZ:  $l^2 = 93\%$ ,  $\tau^2 = 5.031$ , p < .01; BD:  $l^2 = 92\%$ ,  $\tau^2 = 6.629$ , p < .01). To further address the heterogeneity in the meta-analyses, we investigated the potential effects of several moderating variables, including the sex ratio and age of samples in each dataset, brain data type (in vivo vs. post-mortem), magnetic field strengths of the MRI scanners, and the segmentation approaches used.

#### 3.4.1 | Effects of sex ratio and mean age

Information on the sex distribution and mean age of the samples was available for 50 and 42 datasets, respectively. Meta-regression analyses with either sex ratio or mean age as moderators showed no significant effects (Sex:  $R^2 = 5.25\%$ ,  $F_{(1,40)} = 2.81$ , p = .102; Age:  $R^2 = 5.54\%$ ,  $F_{(1,48)} = 2.81$ , p = .100). We also repeated the analyses within the healthy control datasets, and did not find any significant effects (Sex: p = .961; Age: p = .570; given the limited number of datasets available, we did not run separate moderator analyses for each patient group, the same as below).

### TABLE 1 Data extracted from the habenula studies.

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				Left volur	ne (mm <sup>3</sup> )	Right volu	olume (mm <sup>3</sup> )	
References	Subset	Sample category	Ν	Mean	SD	Mean	SD	
I. In vivo studies								
Lawson et al. (2013)		Healthy	24	29.40	4.70	29.30	3.70	
Hétu et al. (2016)		Healthy	34	27.88	8.49	28.03	8.18	
Kim et al. (2016)	а	Healthy	49	21.10	5.20	21.30	4.50	
	b	Healthy	6	18.30	2.30	17.90	2.10	
Torrisi et al. (2017)		Healthy	32	18.80	6.00	14.90	4.00	
He et al. (2020)		Healthy	50	23.90	5.80	23.40	5.40	
Furman and Gotlib (2016)	а	Healthy	13	28.30	3.50	28.70	2.50	
	b	MDD	15	28.80	3.80	28.30	4.20	
Schmidt et al. (2016)	а	Healthy	20	17.63	5.49	17.29	6.12	
	b	unmed.MDD	20	17.18	5.94	17.80	6.41	
	с	med. MDD	20	16.45	2.49	17.70	2.78	
Luan et al. (2019)	а	Healthy	15	22.73	1.87	18.87	2.47	
	b	TRD	15	22.33	3.06	18.73	2.49	
Savitz, Bonne, et al. (2011)	а	Healthy	75	19.80	5.10	17.10	4.60	
	b	PTSD	22	18.80	3.60	16.40	3.70	
Savitz, Nugent, et al. (2011)	а	MDD	28	19.10	4.60	15.60	4.80	
	b	RD	32	19.30	4.50	16.70	4.20	
	с	unmed. BD	22	16.30	3.20	14.10	3.00	
	d	med. BD	15	19.40	5.90	16.80	5.00	
Cho et al. (2021)	а	Healthy	36	18.47	5.04	18.54	3.91	
	b	MDD	33	18.37	3.39	16.43	3.24	
Lawson et al. (2017)	а	Healthy	25	22.72	2.06	22.90	3.71	
	b	MDD	25	20.40	3.22	21.15	3.71	
Liu et al. (2017)	а	Healthy	17	25.79	2.84	25.15	2.75	
	b	MDD	21	30.09	2.57	28.63	2.29	
Zhang et al. (2017)	а	Healthy	16	24.02	3.20	20.42	3.46	
	b	Schizophrenia	15	21.83	2.16	18.27	2.63	
Schafer et al. (2018)	а	Healthy	40	14.91	3.78	15.41	3.86	
	b	BD	32	14.74	4.91	15.93	3.99	
	с	Schizophrenia	68	15.10	3.51	16.93	3.92	
Bocchetta et al. (2016)	а	Healthy	15	23.60	2.20	23.30	2.20	
	b	bvFTD	15	16.90	2.40	16.40	2.70	
	с	AD	15	23.60	3.10	23.00	2.90	
Lim et al. (2021)	а	Healthy	36	12.40	4.00	12.21	3.57	
	b	Healthy	36	11.98	4.06	11.79	3.26	
	с	MDD	33	12.52	3.08	11.21	3.16	
	d	MDD	33	12.58	3.34	11.69	2.82	
Germann et al. (2020)	а	Healthy	26	12.29	2.37	14.20	2.73	
	b	BD	50	12.00	2.31	13.91	2.75	
	с	Schizophrenia	20	12.43	2.49	13.91	2.36	
	d	Healthy	55	14.09	3.38	16.87	4.09	
	e	BD	32	12.81	4.90	16.69	3.76	
	f	Healthy	73	11.61	2.97	13.48	3.25	
	g	Schizophrenia	45	13.88	3.24	15.20	3.41	

(Continues)

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### TABLE 1 (Continued)

					ne (mm <sup>3</sup> )	Right volume (mm <sup>3</sup> )		
References	Subset	Sample category	N	Mean	SD	Mean	SD	
II. Post-mortem studies								
Ahumada-Galleguillos et al. (2017)	а	Healthy	16	39.40	12.60	35.65	11.40	
	b	Healthy	22	35.55	11.50	33.08	10.70	
Ranft et al. (2010)	а	Healthy	13	30.92	5.22	33.23	4.93	
	b	Schizophrenia	17	32.15	5.79	33.93	5.11	
	С	Depression	14	26.78	4.59	25.49	6.02	
Müller et al. (2021)	а	Healthy	12	23.01	4.05	21.92	4.26	
	b	People addicted to heroin	12	17.61	3.73	19.02	4.85	
Kim et al. (2016)	С	Heart diseases	2	34.45	1.77	34.90	1.70	

Abbreviations: AD, Alzheimer disease; BD, bipolar disorder; bvFTD, behavioural variant frontotemporal dementia; MDD, major depressive disorder; *med*, medicated; PTSD, Post-traumatic stress disorder; RD, fully remitted patients with MDD; TRD, treatment-resistant depression; *unmed*, unmedicated.



**FIGURE 3** Scatter plot of the volume of the left (*x*-axis) and right (*y*-axis) habenula (mm<sup>3</sup>). Each dot indicates data from one dataset, with the size indicating its sample size and the color indicating the subgroups shown in the figure legend.

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First author and year (subgroup)	Sample size	Mean Difference	L – R	[95% CI]	Weight
Germann et al., 2020 (e)	32	<b></b>	-3.88	[−5.42 to −2.34]	2.0%
Germann et al., 2020 (d)	55		-2.78	[-3.78 to -1.78]	2.2%
Ranft et al., 2010 (a)	13		-2.31	[-5.07 to 0.45]	1.5%
Germann et al., 2020 (a)	26		-1.91	[-2.90 to -0.92]	2.2%
Germann et al., 2020 (b)	50		-1.91	[-2.62 to -1.20]	2.3%
Germann et al., 2020 (f)	73	-	-1.87	[-2.59 to -1.15]	2.3%
Schafer et al., 2018 (c)	68		-1.83	[-2.72 to -0.94]	2.2%
Ranft et al., 2010 (b)	17		-1.78	[-4.38 to 0.82]	1.5%
Germann et al., 2020 (c)	20		-1.48	[-2.54 to -0.42]	2.2%
Muller et al., 2021 (b)	12		-1.41	[-3.90 to 1.08]	1.6%
Germann et al., 2020 (g)	45		-1.32	[-2.29 to -0.35]	2.2%
Schmidt et al., 2016 (c)	20		-1.25	[-2.41 to -0.09]	2.2%
Schafer et al., 2018 (b)	32		-1.19	[-2.76 to 0.38]	2.0%
Lawson et al., 2017 (b)	25		-0.75	[-2.12 to 0.62]	2.1%
Schmidt et al., 2016 (b)	20		-0.62	[-3.33 to 2.09]	1.5%
Schafer et al., 2018 (a)	40		-0.50	[-1.68 to 0.68]	2.1%
Kim et al., 2016 (c)	2		-0.45	[-2.85 to 1.95]	1.6%
Furman & Gotlib, 2016 (a)	13		-0.40	[-2.10 to 1.30]	1.9%
Kim et al., 2016 (a)	49		-0.20	[-1.57 to 1.17]	2.1%
Lawson et al., 2017 (a)	25		-0.18	[-1.44 to 1.08]	2.1%
Hetu et al., 2016	34	<u>_</u>	-0.15	[-2.95 to 2.65]	1.4%
Cho et al., 2021 (a)	36		-0.07	[-1.57 to 1.43]	2.0%
Lawson et al., 2013	24		0.10	[-1.62 to 1.82]	1.9%
Lim et al., 2021 (a)	36	<u> </u>	0.19	[-1.05 to 1.43]	2.1%
Lim et al., 2021 (b)	36		0.19	[-1.03 to 1.41]	2.1%
Bocchetta et al., 2016 (a)	15		0.30	[-0.81 to 1.41]	2.2%
Schmidt et al., 2016 (a)	20		0.34	[-2.22 to 2.90]	1.5%
Kim et al., 2016 (b)	6		0.40	[-1.37 to 2.17]	1.9%
He et al., 2020	50		0.50	[-1.06 to 2.06]	2.0%
Furman & Gotlib, 2016 (b)	15		0.50	[-1.53 to 2.53]	1.8%
Bocchetta et al., 2016 (b)	15		0.50	[-0.80 to 1.80]	2.1%
Bocchetta et al., 2016 (c)	15	- <u>i</u>	0.60	[-0.92 to 2.12]	2.0%
Liu et al., 2017 (a)	17	- <u>-</u>	0.64	[-0.69 to 1.97]	2.1%
Lim et al., 2021 (d)	33		0.89	[-0.17 to 1.95]	2.2%
Muller et al., 2021 (a)	12		1.09	[-1.27 to 3.45]	1.6%
Ranft et al., 2010 (c)	14		1.29	[−1.57 to 4.15]	1.4%
Lim et al., 2021 (c)	33	÷ • -	1.31	[ 0.25 to 2.37]	2.2%
Liu et al., 2017 (b)	21	÷ • •	1.46	[ 0.42 to 2.50]	2.2%
Cho et al., 2021 (b)	33		1.94	[ 0.81 to 3.07]	2.2%
Savitz et al., 2011b (c)	22		2.20	[ 0.90 to 3.50]	2.1%
Savitz et al., 2011a (b)	22	<b></b>	2.40	[0.87 to 3.93]	2.0%
Ahumada-Galleguillos et al., 2017 (b)	22		2.47	[-2.18 to 7.12]	0.8%
Savitz et al., 2011b (b)	32		2.60	[ 1.09 to 4.11]	2.0%
Savitz et al., 2011b (d)	15	÷ •	2.60	[-0.19 to 5.39]	1.4%
Savitz et al., 2011a (a)	75		2.70	[ 1.60 to 3.80]	2.2%
Savitz et al., 2011b (a)	28		3.50	[ 1.76 to 5.24]	1.9%
Zhang et al., 2017 (b)	15		3.56	[ 2.33 to 4.79]	2.1%
Zhang et al., 2017 (a)	16		3.60	[ 1.96 to 5.24]	2.0%
Luan et al., 2019 (b)	15		3.60	[ 2.17 to 5.03]	2.0%
Anumada-Galleguillos et al., 2017 (a)	16		- 3.75	[-2.15 to 9.65]	0.6%
Luan et al., 2019 (a)	15		3.86	[ 2.73 to 4.99]	2.2%
Iorrisi et al., 2017	32		3.90	[ 2.07 to 5.73]	1.9%
Random effects model		₩	0.40	[-0.13 to 0.94]	100.0%
Heterogeneity: $I^2 = 88\%$ , $\tau^2 = 3.0492$ , $p < .07$	1				
		-5 U 5			

**FIGURE 4** Forest plot of the left-right differences in the habenula volume.

#### 3.4.2 | Effects of brain data type

In terms of brain data types used for the measurement of the habenula volume, 44 datasets were based on in vivo studies (i.e., using MRI), and 8 were based on post-mortem studies (i.e., via brain tissues). Subgroup analysis showed no significant differences associated with the brain data types ( $\chi^2_{(1)} = 1.07$ , p = .302), and the analysis with only the healthy control datasets showed similar results ( $\chi^2_{(1)} = 0.04$ , p = .850).

# 3.4.3 | Effects of the magnetic field strength of the MRI scanners

Within the samples using MRI, we further investigated the effects of various MRI scanners. Here we focused on the magnetic field strength of the scanners, which is perhaps the main factor that could affect the brain imaging measures. Among the 44 datasets, 7 were based on 1.5T MRI scanners, 26 were based on 3T, and 11 were based on 7T

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(d) BD patients

(a) Healthy samples					
First author and year (subgroup)	Sample size	Mean Difference	L – R	[95% CI]	Weight
Germann et al., 2020 (d)	55	-	-2.78	[-3.78 to -1.78]	4.8%
Ranft et al., 2010 (a)	13		-2.31	[-5.07 to 0.45]	3.0%
Germann et al., 2020 (a)	26		-1.91	[-2.90 to -0.92]	4.8%
Germann et al., 2020 (f)	73		-1.87	[-2.59 to -1.15]	5.0%
Schaler et al., 2018 (a)	40		-0.50	[-1.68 to 0.68]	4.0%
Kim et al. 2016 (a)	49		-0.20	[-1.57 to 1.17]	4.170
Lawson et al., 2017 (a)	25		-0.18	[-1.44 to 1.08]	4.5%
Hetu et al., 2016	34		-0.15	[-2.95 to 2.65]	3.0%
Cho et al., 2021 (a)	36		-0.07	[-1.57 to 1.43]	4.3%
Lawson et al., 2013	24		0.10	[-1.62 to 1.82]	4.1%
Lim et al., 2021 (a)	36		0.19	[-1.05 to 1.43]	4.5%
Lim et al., 2021 (b)	36		0.19	[-1.03 to 1.41]	4.6%
Sobmidt et al., 2016 (a)	15	<u> </u>	0.30	[-0.81 to 1.41]	4.7%
Kim et al. $2016$ (b)	6		0.40	[-1.37 to 2.17]	4.0%
He et al., 2020	50		0.50	[-1.06 to 2.06]	4.2%
Liu et al., 2017 (a)	17		0.64	[-0.69 to 1.97]	4.5%
Muller et al., 2021 (a)	12		1.09	[-1.27 to 3.45]	3.4%
Ahumada-Galleguillos et al., 2017 (b)	22		2.47	[-2.18 to 7.12]	1.7%
Savitz et al., 2011a (a)	75		2.70	[ 1.60 to 3.80]	4.7%
Zhang et al., 2017 (a)	16		3.60	[ 1.96 to 5.24]	4.2%
Ahumada-Galleguillos et al., 2017 (a)	16 15		3.75	[-2.15 to 9.65]	1.2%
Torrisi et al. 2017	32		3.80	[2.73 to 4.99]	4.0%
	02	_	0.00	[ 2:07 10 0:70]	1.070
Random effects model			0.40	[-0.37 to 1.16]	100.0%
Heterogeneity: $I^2 = 87\%$ , $\tau^2 = 2.7560$ , $p < .0$	01				
		-5 0 5			
(b) MDD nationts					
(b) MIDD patients					
First author and year (subgroup)	Sample size	Mean Difference	I - R	[95% CI]	Weight
First author and year (subgroup)	Sample size	Mean Difference	L – R	[95% CI]	Weight
First author and year (subgroup) Schmidt et al., 2016 (c)	Sample size	Mean Difference	<b>L – R</b> –1.25	<b>[95% CI]</b> [-2.41 to -0.09]	Weight 9.9%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b)	Sample size 20 25	Mean Difference	L – R -1.25 -0.75	<b>[95% CI]</b> [-2.41 to -0.09] [-2.12 to 0.62]	9.9% 9.4%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b)	20 25 20	Mean Difference	L - R -1.25 -0.75 -0.62	<b>[95% CI]</b> [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09]	Weight 9.9% 9.4% 6.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b)	20 25 20 15 22	Mean Difference	L – R -1.25 -0.75 -0.62 0.50	<b>[95% CI]</b> [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.05]	Weight 9.9% 9.4% 6.0% 7.6%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c)	Sample size 20 25 20 15 33 33	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [-0.25 to 2.37]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2017 (b)	Sample size 20 25 20 15 33 33 21	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50]	Weight           9.9%           9.4%           6.0%           7.6%           10.2%           10.1%           10.2%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2017 (c) Liu et al., 2017 (b) Cho et al., 2021 (b)	Sample size 20 25 20 15 33 33 21 33	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94	[95% C]] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07]	Weight           9.9%           9.4%           6.0%           7.6%           10.2%           10.2%           10.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Cho et al., 2021 (b) Savitz et al., 2011b (b)	20 25 20 15 33 33 21 33 32	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60	[95% C]] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11]	Weight           9.9%           9.4%           6.0%           7.6%           10.2%           10.2%           10.2%           10.0%           9.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a)	Sample size 20 25 20 15 33 33 21 33 32 28	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50	[95% C]] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b)	Sample size 20 25 20 15 33 33 21 33 21 33 32 28 15	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4% 9.2%
<b>First author and year (subgroup)</b> Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2017 (b) Cho et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b)	Sample size 20 25 20 15 33 33 21 33 21 33 22 28 15	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07] [ 1.09 to 4.11] [ 1.76 to 5.24] [ 2.17 to 5.03]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 8.4% 9.2%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Schmidt et al., 2017 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model	Sample size 20 25 20 15 33 33 21 33 22 28 15 32 28 15	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b>	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07] [ 1.09 to 4.11] [ 1.76 to 5.24] [ 2.17 to 5.03] [ 0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 8.4% 9.2% <b>100.0%</b>
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 1000$	Sample size 20 25 20 15 33 21 33 32 28 15 .01	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b>	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07] [ 1.09 to 4.11] [ 1.76 to 5.24] [ 2.17 to 5.03] [ 0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 8.4% 9.2% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2017 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 0.0000$	Sample size 20 25 20 15 33 33 21 33 32 28 15 .01	Mean Difference	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b>	[95% C]] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.53] [-0.17 to 1.95] [0.25 to 2.537] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4% 9.2% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Cho et al., 2021 (b) Savitz et al., 2011 b (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$	Sample size 20 25 20 15 33 21 33 32 28 15 .01	Hean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b>	[95% C]] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07] [ 1.09 to 4.11] [ 1.76 to 5.24] [ 2.17 to 5.03] [ 0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (d) Lim et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$ (c) SCZ patients First author and year (subgroup)	Sample size 20 25 20 15 33 33 21 33 32 28 15 .01 Sample size	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b>	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 8.4% 9.2% 100.0% Weight
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p <$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c)	Sample size 20 25 20 15 33 33 21 33 32 28 15 .01 Sample size 68	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 8.4% 9.2% 100.0% Weight 21.2%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011 b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p <$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2010 (b)	Sample size 20 25 20 15 33 32 21 33 32 28 15 .01 Sample size 68 17	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [ 0.25 to 2.37] [ 0.42 to 2.50] [ 0.81 to 3.07] [ 1.09 to 4.11] [ 1.76 to 5.24] [ 2.17 to 5.03] [ 0.14 to 2.33] [ 0.14 to 2.33]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0% Weight 21.2% 16.4%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p <$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (c)	Sample size 20 25 20 15 33 32 21 33 32 28 15 .01 .01 Sample size 68 17 20	Mean Difference	L - R -1.25 -0.75 -0.62 0.50 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b> L - R -1.83 -1.78 -1.48	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.30] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [95% CI] [-2.72 to -0.94] [-4.38 to 0.82] [-2.54 to -0.42]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0% 8.4% 9.2% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 0$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (c) Germann et al., 2020 (g)	Sample size 20 25 20 15 33 32 28 15 .01 Sample size 68 17 20 45	Mean Difference	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 <b>1.23</b> L - R -1.83 -1.78 -1.48 -1.32	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [-2.72 to -0.94] [-4.38 to 0.82] [-2.54 to -0.42] [-2.29 to -0.35]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4% 9.2% 100.0% 8.4% 9.2% 100.0% 21.2%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (c) Germann et al., 2020 (g) Zhang et al., 2017 (b)	Sample size 20 25 20 15 33 32 28 15 .01 Sample size 68 17 20 45 15 .01	Mean Difference	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78 -1.48 -1.32 3.56	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.42 to 2.50] [0.42 to 2.50] [0.42 to 2.30] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [-2.72 to -0.94] [-4.38 to 0.82] [-2.54 to -0.42] [-2.29 to -0.35] [2.33 to 4.79]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0% 8.4% 9.2% 100.0% 2.12% 16.4% 20.5%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (c) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2017 (b) Cho et al., 2021 (b) Savitz et al., 2011b (a) Luan et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (c) Germann et al., 2020 (g) Zhang et al., 2017 (b)	Sample size 20 25 20 15 33 32 28 15 .01 Sample size 68 17 20 45 15 .15 .01	Mean Difference	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78 -1.83 -1.78 -1.48 -1.32 3.56	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [-2.72 to -0.94] [-4.38 to 0.82] [-2.54 to -0.42] [-2.33 to 4.79] [-3.34 to 2.25]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4% 9.2% 100.0% Veight 21.2% 16.4% 20.9% 21.1% 20.5%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2017 (b) Cho et al., 2021 (b) Savitz et al., 2011b (a) Luan et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$ Schafer et al., 2018 (c) Ranft et al., 2018 (c) Ranft et al., 2020 (c) Germann et al., 2020 (g) Zhang et al., 2017 (b) Random effects model Heterogeneity: $I^2 = 93\%$ , $\tau^2 = 5.0308$ a s	Sample size 20 25 20 15 33 33 21 33 32 28 15 .01 .01 Sample size 68 17 20 45 15 .01	Mean Difference	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78 -1.48 -1.55 -1.	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [95% CI] [-2.72 to -0.94] [-4.38 to 0.82] [-2.34 to -0.42] [-2.29 to -0.35] [2.33 to 4.79] [-3.44 to 2.36]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0% Weight 21.2% 16.4% 20.9% 21.1% 20.5% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2021 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $I^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < 100$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (g) Zhang et al., 2017 (b) Random effects model Heterogeneity: $I^2 = 93\%$ , $\tau^2 = 5.0308$ , $p < 100$	Sample size 20 25 20 15 33 32 28 15 .01 .01 Sample size 68 17 20 45 15 .01	Mean Difference $ \begin{array}{cccc}  & & & & \\  & & & & \\  & & & & \\  & & & &$	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78 -1.48 -1.32 3.56 -0.54	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [0.14 to 2.33] [-2.72 to -0.94] [-4.38 to 0.82] [-2.54 to -0.42] [-2.29 to -0.35] [2.33 to 4.79] [-3.44 to 2.36]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.0% 9.0% 8.4% 9.2% 100.0% Weight 21.2% 16.4% 20.5% 100.0%
First author and year (subgroup) Schmidt et al., 2016 (c) Lawson et al., 2017 (b) Schmidt et al., 2016 (b) Furman & Gotlib, 2016 (b) Lim et al., 2021 (c) Liu et al., 2021 (c) Liu et al., 2011 (b) Savitz et al., 2011b (b) Savitz et al., 2011b (a) Luan et al., 2019 (b) Random effects model Heterogeneity: $l^2 = 80\%$ , $\tau^2 = 2.0557$ , $p < d$ (c) SCZ patients First author and year (subgroup) Schafer et al., 2018 (c) Ranft et al., 2020 (g) Germann et al., 2020 (g) Zhang et al., 2017 (b) Random effects model Heterogeneity: $l^2 = 93\%$ , $\tau^2 = 5.0308$ , $p < d$	Sample size 20 25 20 15 33 32 28 15 .01 Sample size 68 17 20 45 15 .01	Hean Difference $ \begin{array}{c}                                     $	L - R -1.25 -0.75 -0.62 0.89 1.31 1.46 1.94 2.60 3.50 3.60 1.23 L - R -1.83 -1.78 -1.48 -1.48 -1.32 3.56 -0.54	[95% CI] [-2.41 to -0.09] [-2.12 to 0.62] [-3.33 to 2.09] [-1.53 to 2.53] [-0.17 to 1.95] [0.25 to 2.37] [0.42 to 2.50] [0.81 to 3.07] [1.09 to 4.11] [1.76 to 5.24] [2.17 to 5.03] [0.14 to 2.33] [0.14 to 2.33] [95% CI] [-2.72 to -0.94] [-4.38 to 0.82] [-2.34 to -0.42] [-2.29 to -0.35] [2.33 to 4.79] [-3.44 to 2.36]	Weight 9.9% 9.4% 6.0% 7.6% 10.2% 10.1% 10.2% 10.0% 9.0% 8.4% 9.2% 100.0% 100.0% 21.2% 16.4% 20.9% 21.1% 20.5% 100.0%

First author and year (subgroup)	Sample size	Mean Difference	L – R	[95% CI]	Weight
Germann et al., 2020 (e)	32	— <b>·</b>	-3.88	[-5.42 to -2.34]	20.3%
Germann et al., 2020 (b)	50		-1.91	[-2.62 to -1.20]	21.7%
Schafer et al., 2018 (b)	32		-1.19	[-2.76 to 0.38]	20.2%
Savitz et al., 2011b (c)	22		2.20	[0.90 to 3.50]	20.8%
Savitz et al., 2011b (d)	15	-	2.60	[-0.19 to 5.39]	17.0%
Random effects model			-0.54	[-3.94 to 2.85]	100.0%
Heterogeneity: $I^2 = 92\%$ , $\tau^2 = 6.6285$ , p	< .01				
		-4 -2 0 2 4			

**FIGURE 5** Forest plots of inter-hemispheric habenula volume differences in different subgroups. (a) Healthy samples. (b) MDD patients. (c) SCZ patients. (d) BD patients.



FIGURE 6 Forest plots for the moderator analysis of magnetic field strength of the MRI scanners.

(Figure 6). As expected, subgroup analyses showed significant moderating effects ( $\chi^2_{(2)} = 65.37$ , p < .001),—effects that were mainly contributed by the significant left-right differences observed in the 3T datasets (MD = 1.15 mm<sup>3</sup>; 95% CI: 0.45 to 1.84; left > right:  $t_{(25)} = 3.41$ , p = .002) and 1.5T datasets (MD = -2.02 mm<sup>3</sup>; 95% CI: -2.66 to -1.37; left < right:  $t_{(6)} = -7.63$ , p = .0003;  $t_{(31)} = -4.952$ ,

p < .001, d = -2.109). Note that no differences were found when comparing the sample sizes of the datasets in each pair of three subgroups (ps > .05). In addition, with only the healthy control datasets, the results were similar in the subgroup analyses of both 3T scanners (MD = 0.90 mm<sup>3</sup>; 95% Cl: -0.12 to 1.92; left > right:  $t_{(11)} = 1.95$ , p = .077); and 1.5T scanners (MD = -2.12 mm<sup>3</sup>; 95% Cl: -3.31 to -0.92; left < right:  $t_{(2)} = -7.62$ , p = .017;  $t_{(13)} = -3.203$ , p = .007, d = -2.067). Note that the samples of the 3 and 1.5T datasets showed inversed asymmetry patterns which might also relate to different measurement approaches used such as segmentation (see below and also Section 4). In addition, no significant differences were found in the 7T datasets ( $t_{(10)} = 1.68$ , p = .125).

#### 3.4.4 | Effects of segmentation methods

The various segmentation methods used could be another vital factor contributing to the observed heterogeneity. Among the 44 MRI datasets, 28 were identified as using manual segmentation approaches, 9 used fully automated segmentation, and 7 used semi-automated segmentation (Figure 7). Subgroup analyses by segmentation method showed significant moderating effects ( $\chi^2_{(2)} = 52.32$ , p < .001). Specifically, we found that in the manual subgroup, the habenula volume was significantly larger in the left hemisphere compared with the right (MD = 1.49 mm<sup>3</sup>; 95% Cl: 0.91–2.07; left > right:  $t_{(27)} = 5.27$ , p < .001), while inverse patterns were found in the semi-automated (MD = -0.99; 95% Cl: -1.65 to -0.33; left < right:  $t_{(6)} = -3.68$ , p = .010) and fully-automated subgroups (MD = -1.54; 95% Cl: -2.61 to -0.48; left < right:  $t_{(6)} = -3.35$ , p = .010).

# 3.4.5 | Effects of the average volume of the bilateral habenula

We also investigated the moderating effects of the volume of the habenula on the left-right difference estimation. Meta-regression analyses showed no significant effects for either analysis across all the datasets, or for that within the healthy control datasets (all:  $R^2 = 1.10\%$ ,  $F_{(1,50)} = 1.11$ , p = .298; healthy:  $R^2 = .45\%$ ,  $F_{(1,23)} = 0.96$ , p = .338).

#### 3.5 | Sensitivity analyses

To rule out the potential confounding effects of the correlation of the bilateral habenula volume on the meta-analysis results, we repeated all analyses with a range of correlation values from 0.30 to 0.80. The results remained similar in all cases (Figure 8).

In addition, to evaluate the potential effects of the datasets with too few samples in the meta-analyses, we excluded these small datasets (i.e., those with no more than 15 individuals) and re-ran all analyses. The number of datasets included in the metaanalyses decreased remarkably, at least in some cases, but the main results remained similar except for the subgroup analyses of the left-right differences in the SCZ samples (Figure 8). Specifically, the inter-hemispheric differences in the SCZ samples became highly significant after excluding one small dataset (p < .002).

# 3.6 | Disorder-related differences in the habenula volume

We also used the data generated to investigate disorder-related differences in habenular volume. Meta-analyses showed no significant differences in the unilateral habenula volume of either samples with MDD (left:  $t_{(10)} = 0.01$ ; right:  $t_{(10)} = -0.53$ ), SCZ (left:  $t_{(4)} = 0.43$ ; right:  $t_{(4)} = 0.57$ ), or BD (left:  $t_{(4)} = -2.07$ , p = 0.107; right:  $t_{(4)} = -1.08$ , p = .341), compared with healthy control datasets from the same study. Cross-disorder comparisons showed no significant differences (ps > .05) except for the differences in the left habenula between the MDD subgroup and the BD subgroup (MDD > BD;  $Q_{(1)} = 7.70$ , p = .006).

#### 4 | DISCUSSION

This study presents a large-scale analysis of structural asymmetry of the habenula in the human brain. Meta-analyses of 52 datasets (N = 1427) revealed significant heterogeneity in the left-right differences in the habenula volume across datasets. The magnetic field strength of the MRI scanners (e.g., 3 or 1.5T) and segmentation methods (e.g., manual or automated) applied were found to be two of the key contributing factors. While little evidence was found for volume asymmetry in the meta-analysis of all of the datasets, the most pronounced left-right differences (i.e., left > right) were found in more homogeneous subgroup of datasets with 3T scanners and the manual segmentation approach. We did not find significant disorder-related differences, relative to healthy controls in either the left-right differences or the unilateral volume per se.

# 4.1 | Left-right differences in the volume of the habenula

The habenula volume of the left and right hemispheres showed a high correlation (r = .96) across datasets (i.e., at the study level), suggesting high cross-hemisphere similarity for the bilateral structure in the human brain. This seems to be inconsistent with the correlations reported at the individual level (around 0.50; e.g., Germann et al., 2020), which suggests a considerable extent of distinctness. Such inconsistency could be due to higher measurement errors of individual data, and/or larger variance across studies at the study-level analysis. Moreover, mixed asymmetry patterns of the habenula volume have been suggested in the previous studies (see Section 1). The present study, with a large number of samples, revealed little evidence



FIGURE 7 Forest plots for the moderator analysis of segmentation approaches.

to support left-right differences in the habenula volume. Interestingly, we successfully identified several significant moderating factors that could impact the left-right comparison results, such as the field

strength of the scanners and the segmentation approaches. In terms of scanners, we found that datasets from 3T scanners tended to show a leftward asymmetry in the habenular volume, while datasets from of use; OA

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		k	<i>r</i> = .3	r = .4	r = .5	<i>r</i> = .6	<i>r</i> = .7	r = .8	_		k	<i>r</i> = .3	r = .4	r = .5	r = .6	r = .7	r = .8
Inter-hemispheric difference	All participants	52	0.8353	0.8539	0.8735	0.8955	0.9215	0.9551		4	36	0.0982	0.1135	0.1306	0.1503	0.1741	0.2052
Inter-hemispheric difference	Health subgroup	25	0.4947	0.5122	0.5318	0.5547	0.5834	0.6216			19	0.2695	0.2950	0.3239	0.3579	0.3998	0.4544
Inter-hemispheric difference	MDD subgroup	11	1.5258	1.5186	1.5086	1.4989	1.4855	1.4711			9	1.1002	1.0931	1.0846	1.0747	1.0635	1.0501
Inter-hemispheric difference	SCZ subgroup	5	0.1940	0.1957	0.19 <b>78</b>	0.2002	0.2029	0.2061		0	4	2.9208	2.9208	2.9208	2.9208	2.9208	2.9208
Inter-hemispheric difference	BD subgroup	5	0.1819	0.1748	0,1676	0.1601	0.1526	0.1448			4	0,3803	0,3802	0,3802	0.3802	0,3802	0,3802
Moderator effect	Mean age	50	1.0650	1.0325	0.9991	0.9650	0.9311	0.8972			35	1.8508	1.8239	1.7986	1.7721	1.7447	1.7235
Moderator effect	Sex ratio	42	0.9851	0.9889	0.9931	0.9983	1.0044	1.0123			28	1.3019	1.2381	1.1733	1.1062	1.0362	0.9610
Moderator effect	Segmentation method manual	28	>4.0000	>4.0000	>4.0000	>4.0000	>4.0000	>4.0000			17	2.9586	2.9208	2.9208	2.9208	2.8861	2.8861
Moderator effect	Segmentation method semi-automated	7	2.0757	2.0315	1.9830	1.9281	1.8633	1.7747			7	2.0757	2.0315	1.9830	1.9281	1.8633	1.7747
Moderator effect	Segmentation method fully automated	9	2.0177	2.0044	1.9957	1.9830	1.9747	1.9666			9	2.0177	2.0044	1.9957	1.9830	1.9747	1.9666
Moderator effect	Brain data type in vivo	44	0.8986	0.9190	0.9397	0.9610	0.9825	1.0044			33	0.0812	0.0914	0.1016	0.1118	0.1219	0.1319
Moderator effect	Brain data type post-mortem	8	0.1948	0.1836	0.1514	0.1103	0.0588	0.0017			3	0.1111	0.1350	0.1605	0.1878	0.2175	0.2502
Moderator effect	1.5T MRI	7	3.6990	3.6990	3.5229	3.3979	3.3979	3.3010			7	3.6990	3.6990	3.5229	3.3979	3.3979	3.3010
Moderator effect	3.0T MRI	26	2.6383	2.6383	2.6576	2.6778	2.6778	2.6990			16	1.0297	1.0391	1.0477	1.0565	1.0640	1.0711
Moderator effect	7.0T MRI	11	0.9212	0.9129	0.9038	0.8931	0.8804	0.8655			10	0.8419	0.8342	0.8256	0.8159	0.8044	0.7905
Moderator effect	Average volume of the bilateral habenula	52	0.5702	0.5467	0.5262	0.5105	0.5025	0.5084			36	0.8935	0.8918	0.8965	0.9097	0.9370	0.9884

**FIGURE 8** Results of the sensitivity analyses. Color indicates the statistical significance ( $-\log(p)$ ) in each analysis: the left panel for analyses with all dataset, the right for analyses after excluding the datasets with no more than 15 samples.

1.5T scanners tended to show an inverse pattern (the 1.5T datasets were from the same study, i.e., Germann et al., 2020). Datasets from 7T scanners showed no significant asymmetry although we would expect better imaging quality with such a high field strength. The scanner effects seem to be unique to the habenular volume asymmetry as we did not find significant effects on the asymmetry of cortical thickness or area (Kong et al., 2018), or the volume asymmetry of larger subcortical structures such as the hippocampus or thalamus (Guadalupe et al., 2017). In terms of segmentation approaches, MRI studies using a manual approach tended to report a leftward asymmetry, while those using a semi- or fully-automated approach tended to report a rightward asymmetry. There could be some interesting interactions between scanner-related factors and the segmentation approaches used in measuring the hemisphere-related differences, which could be addressed in future studies.

To our knowledge, this is the first large-scale empirical investigation of habenular structural asymmetry. The results provided robust data on the moderating effects of multiple factors including scanners and habenula segmentation. Such moderating effects suggested that when developing new segmentation algorithms for future studies, the potential asymmetrical nature of the brain structures should be taken into consideration if possible.

### 4.2 | Disorder-related differences in the habenula

The habenula has been implicated in a variety of psychiatric disorders such as depression and anxiety disorders (Boulos et al., 2017;

Fakhoury, 2017), and also considered as a promising target for the treatment of intractable conditions (Sartorius et al., 2010: Sartorius & Henn, 2007). A few brain structural studies have suggested asymmetrical alterations in the habenular volume in patients with, for example, MDD (e.g., Ranft et al., 2010) or SCZ (e.g., Zhang et al., 2017). However, the present meta-analyses showed little differences when comparing the unilateral habenular volume data of patients with that of the healthy controls. Interestingly, we found that patients with MDD tended to show a larger volume in the left hemisphere than the right, and patients with SCZ seemed to possess a larger habenular volume in the right hemisphere. Note that these findings were obtained using a relatively smaller number of datasets and meanwhile could be confounded by variability in brain imaging approaches (e.g., all datasets with 7 or 3T scanners for MDD versus half of the datasets with 1.5T scanners for SCZ). Thus such differences should be interpreted with caution. At best, these observations could serve as potential hypotheses for additional experiments that include independent data.

#### 4.3 | Limitations and future directions

This study has several limitations that could be overcome in future studies. First, while this is to our best knowledge the largest study on the habenular asymmetry, the findings warrant further investigation with a larger number of individuals. The UK Biobank cohort (Sudlow et al., 2015) and the ENIGMA working groups (Thompson et al., 2020) could provide great opportunities for achieving a more definitive picture of the nature of human habenula asymmetry. Second, an

automated and unbiased segmentation approach is necessary for analyzing such large-scale individual data. Currently, a few approaches have been proposed such as MAGeTbrain (Chakravarty et al., 2013; Pipitone et al., 2014), myelin content-based segmentation (Kim et al., 2016, 2018), and deep learning-based U-Net segmentation (Lim et al., 2021). It remains to be determined whether such approaches could detect any hemispheric differences in the habenula. In addition, the habenula can be divided into medial and lateral habenula both structurally and functionally in animal studies (Fakhoury & Domínguez López, 2014; Hikosaka, 2010). Ultra-high-resolution in vivo MR imaging could provide new data on the functional lateralization of the habenula in the human brain and its structural basis. Finally, it would be of interest to investigate whether the results are specific to the habenula, or would be found in general in similar small, ill-defined anatomical structures. The operator's laterality, for example, in hand and eye preference could also be an important factor for manual segmentation of small brain structures.

#### 4.4 | Summary

In summary, the present study presents a large-scale analysis of structural asymmetry of the habenula in the human brain. Results showed significant heterogeneity in both the habenula volume and its leftright differences across different studies. The magnetic field strength of the MRI scanners and segmentation methods used were found to be two of the key contributing factors to such heterogeneity. The most pronounced left-right differences (i.e., left > right) were found in datasets with 3T scanners and manual segmentation approach. While inversed asymmetry patterns were suggested in MDD (left > right) and S'CZ (left < right) patients, little evidence for the differences was found when comparing with healthy samples or other disorder samples. This study provides useful data for future studies of brain imaging and methodological developments related to precision habenula segmentation, and also contributes to the understanding of potential roles of habenular laterality in health and disorders.

#### AUTHOR CONTRIBUTIONS

Study conception and design, data preparation, analysis, visualization, and preparing the first draft: Yilamujiang Abuduaini. Writing and editing: Yi Pu and Paul M. Thompson. Study conception and design, data preparation, analysis, visualization, preparing the first draft, and editing: Xiang-Zhen Kong.

#### ACKNOWLEDGMENTS

Xiang-Zhen Kong is supported by the STI 2030-Major Projects (2021ZD0200409), National Natural Science Foundation of China (32171031), Fundamental Research Funds for the Central Universities (2021XZZX006), and Information Technology Center of Zhejiang University. Paul M. Thompson is supported in part by the U.S. National Institutes of Health under grant R01 MH116147. We thank Prof.

#### CONFLICT OF INTEREST STATEMENT

The authors indicate no competing interest.

#### DATA AVAILABILITY STATEMENT

Our study primarily relies on the dataset extracted from the published papers. We provide the data and code at https://osf.io/f7tpz/.

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### SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article. **How to cite this article:** Abuduaini, Y., Pu, Y., Thompson, P. M., & Kong, X.-Z. (2023). Significant heterogeneity in structural asymmetry of the habenula in the human brain: A systematic review and meta-analysis. *Human Brain Mapping*, 1–18. <u>https://doi.org/10.1002/hbm.26337</u>