

# Magnetic Resonance Imaging–Derived Markers of Acute and Chronic Inflammatory Processes in the Ventral Tegmental Area Associated With Depression

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## ABSTRACT

**BACKGROUND:** Depression is a leading cause of disability worldwide, with inflammation increasingly recognized as a contributing factor. Inflammatory processes can disrupt the brain's reward circuitry, particularly the ventral tegmental area (VTA), which is central to dopamine-mediated motivation and reward. This study investigates whether magnetic resonance imaging–derived markers sensitive to neuroinflammation and microstructure in the VTA are associated with depression diagnosis and symptom severity.

**METHODS:** We analyzed diffusion-weighted imaging and quantitative susceptibility mapping data from 32,495 UK Biobank participants, including 3807 individuals with ICD-10–diagnosed major depressive disorder (MDD). Metrics sensitive to neuroinflammation (free water [FW], isotropic volume fraction [ISOVF], magnetic susceptibility) and microstructure (intracellular volume fraction [ICVF], orientation dispersion index [ODI], volume) were extracted from the VTA. Group differences between the MDD group and healthy control (HC) group matched on body mass index, sex, and age were assessed using analysis of variance. Linear regression was used to predict acute symptom severity based on Recent Depressive Symptoms scores.

**RESULTS:** Participants with an MDD diagnosis had significantly higher FW ( $p < .001$ ) and ISOVF ( $p = .001$ ) compared with HCs, indicating increased extracellular processes such as inflammation in the VTA. Lower ISOVF ( $\beta = -0.28$ ,  $p = .033$ ) and higher ICVF ( $\beta = 0.29$ ,  $p = .017$ ) and ODI ( $\beta = 0.4$ ,  $p = .007$ ) were associated with higher depression severity, independent of depressive diagnosis history.

**CONCLUSIONS:** Our findings reveal distinct patterns of VTA microstructural changes associated with major depression history versus acute depressive symptom severity, suggesting different underlying pathophysiological mechanisms. Distinct patterns of neuroinflammation may differentiate acute from chronic depression, informing targeted interventions.

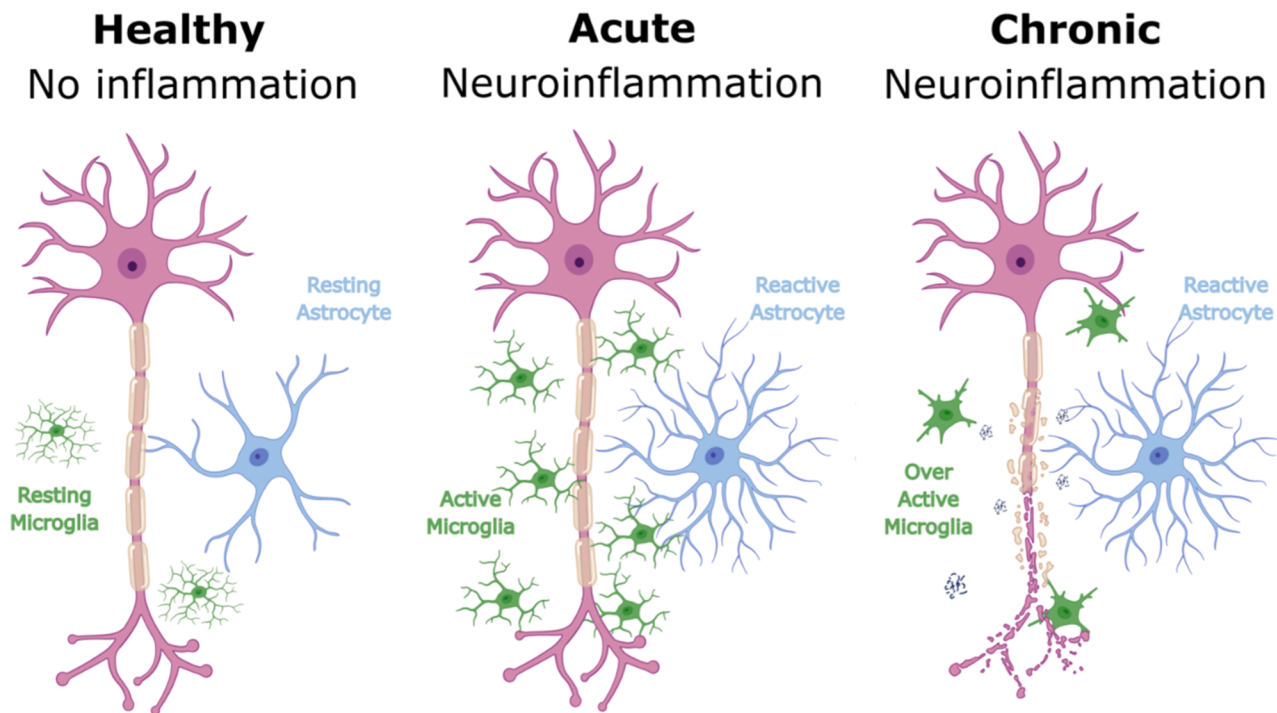
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According to the World Health Organization, depression affects approximately 300 million individuals worldwide, making it a leading cause of disability (1). Depression is a complex, multifactorial disorder, with growing evidence suggesting that inflammation plays a crucial role in its pathophysiology, contributing to both its onset and progression (2). Studies report that individuals with depression often exhibit inflammatory markers and cytokines in their cerebrospinal fluid and blood, suggesting the activation of microglia and astrocytes (see Figure 1) (3). These inflammatory responses can be triggered by various environmental factors, prolonged stress, infections, neurological conditions, and autoimmune diseases, among others (4,5). Chronic inflammation may lead to oxidative and nitrosative stress, causing neurotoxicity and neuronal damage, both of which have been linked to depression (6).

Inflammation disrupts the brain's reward network, which is central to regulating motivation and pleasure (7). Anhedonia

and lack of motivation, hallmark symptoms of depression, may be linked to the impact of inflammatory cytokines on mesolimbic dopamine signaling (8). Previous research suggests that inflammatory processes can disrupt the synthesis, release, and reuptake of dopamine, potentially reflecting an adaptive mechanism that prioritizes energy conservation over reward-seeking behaviors during inflammatory states (9). This notion is further supported by animal models demonstrating that inflammation reduces reward-seeking behaviors (10,11).

At the center of the reward network is the ventral tegmental area (VTA), a rich dopaminergic region that plays a critical role in regulating mood, motivation, and reward (12). The VTA is susceptible to ischemia and inflammation, which can lead to neuronal dysfunction in this region (13). Although previous studies have reported decreased VTA activation in depression, microstructural changes in the VTA remain underexplored, likely due to its small, complex structure (14).



**Figure 1.** Cellular changes in response to varying degrees of inflammation. The figure illustrates the progression of neuroinflammation from a healthy state to acute and chronic inflammation. In the absence of inflammation, microglia (in green) exhibit a resting morphology with fine, branched processes, and astrocytes (in blue) maintain their characteristic star-shaped structure. During acute neuroinflammation, microglia become activated, adopting an amoeboid shape, while astrocytes display hypertrophy, indicating reactivity. In chronic neuroinflammation, sustained elevated cytokine levels cause prolonged activation in both cell types; microglia become overactive, while astrocytes remain reactive. This results in cellular toxicity and potential neurodegeneration, characterized by increased iron deposition and myelin sheath breakdown.

However, recent advancements in neuroimaging, such as high-resolution parcellation atlases, now enable more precise VTA mapping (15).

Several neuroimaging studies have investigated relationships between inflammation and depression, using diverse imaging biomarkers. For example, positron emission tomography (PET) studies have revealed increased radiotracer uptake by microglia in participants with depression, indicating heightened inflammation (16). While the magnetic resonance imaging (MRI)-based technique quantitative susceptibility mapping (QSM) does not directly measure inflammation, it quantifies brain iron accumulation, which may reflect secondary effects of chronic neuroinflammatory processes, such as oxidative stress and prolonged microglial activation (17). QSM is sensitive to tissue magnetic susceptibility, particularly from paramagnetic substances such as iron, as well as diamagnetic components such as calcium and myelin (18). Therefore, abnormal iron distribution observed in depression and other psychiatric disorders (19,20) may provide indirect insights into inflammatory mechanisms over time. Diffusion-weighted imaging (DWI) has similarly revealed inflammation-related microstructural changes (21) and elevated free water (FW) in the extracellular space, commonly associated with inflammatory processes (22). For example, patients with poststroke depression exhibit elevated FW in reward-related regions such as the amygdala and hippocampus (23,24).

In this study, we utilize QSM and DWI to noninvasively examine whether putative markers sensitive to inflammation, as well as microstructural markers that may be altered as a secondary consequence of inflammation, are associated with depression in the VTA. Given the established relationship between inflammation, dopaminergic dysfunction, and depression, as well as the VTA's central role in reward circuitry, we hypothesized that individuals with a history of major depressive disorder (MDD) would exhibit significantly higher levels of QSM- and DWI-derived markers in the VTA compared with matched healthy control (HC) participants. Furthermore, based on the transient nature of inflammation, we hypothesized that acute depression severity would be predicted by MRI markers sensitive to both inflammation and microstructural changes in the VTA irrespective of diagnostic depression history.

## METHODS AND MATERIALS

### Participants

The UK Biobank (UKB) is a large-scale, prospective epidemiological cohort study comprising approximately 500,000 participants recruited from the general UK population (25). Here, we utilized a subsample of the UKB for whom DWI, QSM, and T1-weighted MRIs of brain anatomy were acquired ( $n = 46,703$ ). Participants with intellectual disabilities, psychotic or bipolar disorders, and a history of head injury or

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neurological conditions were excluded due to their known impact on brain structure (see [Table S1](#) for a detailed variable list). We did not exclude or control for immunometabolic or inflammatory comorbidities due to limited inflammatory data in the UKB at the time of scanning and to maintain the generalizability of findings to real-world clinical populations. Participants with missing data or outliers ( $>3$  SDs from the mean) were excluded, yielding a final sample of 32,495. Ethical approval was granted by the UKB (11/NW/0382; Application No. 100773) and ratified by the University of Queensland (2023/HE000221).

### Depression Measures

Of our total sample, 3807 had a history of MDD, based on ICD-10 diagnosis (UKB field ID: 41270). To compare individuals with and without a history of depression, we matched a subset of participants on body mass index (BMI), age, and sex. Descriptive statistics for both groups are presented in [Table 1](#).

To address the possibility that an ICD-10 diagnosis may not reflect current depressive state (some diagnoses were established years before the imaging assessment), we used the Recent Depressive Symptoms (RDS) questionnaire to quantify acute depressive symptoms on the imaging day. The RDS includes 4 items (depressed mood, unenthusiasm/disinterest, tenseness/restlessness, tiredness/lethargy) assessed over the previous 2 weeks, rated on a 4-point Likert scale (1 = not at all, 4 = nearly every day), yielding a total score of 4 to 16 (UKB field IDs: 2050, 2060, 2070, 2080). Higher scores indicate more frequent and severe depressive symptoms, with scores  $\geq 13$  representing individuals with particularly high symptom severity. The RDS has been validated against other depression scales, including the Patient Health Questionnaire-9 ([26](#)).

### Imaging Acquisition and Processing

DW, susceptibility-weighted, and T1-weighted MR images were acquired by the UKB at several sites on a 3T Siemens Skyra scanner with a 32-channel radio frequency receiver head coil, using a standardized acquisition protocol ([27](#)).

T1-weighted (1 mm isotropic) and DW (2 mm isotropic) images were acquired using magnetization-prepared rapid acquisition gradient-echo (MPRAGE) and multiband echo-planar imaging sequences, respectively ([27](#)). DWI used a multishell approach ( $b = 1000$  and  $2000$  s/mm<sup>2</sup> with 50 directions per shell,  $5 b = 0$  s/mm<sup>2</sup>, and 1 reverse-phase encoded  $b = 0$  s/mm<sup>2</sup>). Preprocessing included gradient distortion, slice outlier, head motion, and eddy current corrections ([27](#)). Further details on the preprocessing pipeline can be found in the [Supplement](#). FW-corrected tensor maps were derived from preprocessed DWI data ([28](#)), and neurite orientation dispersion and density imaging (NODDI) was estimated by the UKB using Accelerated Microstructure Imaging via Convex Optimization (AMICO) ([29,30](#)). NODDI measures included isotropic volume fraction (ISOVF), intracellular volume fraction (ICVF), and orientation dispersion index (ODI). QSM images from susceptibility-weighted imaging were minimally preprocessed by the UKB ([27](#)) and minimum-maximum normalized (0–1) for comparison.

Neuroimaging analyses were conducted on the University of Queensland's high-performance computer Bunya ([31](#)), utilizing Neurodesk ([32](#)). The VTA was delineated with the Levinson-Bari Limbic Brainstem Atlas ([15](#)), with T1 images coregistered to DWI using advanced normalization tools (ANTs) ([33](#)), followed by Montreal Neurological Institute (MNI)-to-T1 registration and transformation to DWI space ([34](#)) (see [Figure 2](#)). QSM images were similarly coregistered to DWI participant space. To validate the accuracy of the atlas-derived VTA mask, we manually segmented the VTA in an independent 7T dataset using high-resolution T1 and QSM images. The atlas was coregistered to each participant's T1 image in subject space using the same protocol as in the main analysis, and spatial overlap was quantified using the dilated Dice similarity coefficient. Full details and results, which validated that atlas-based VTA masks were well localized, are described in the [Supplement](#).

We extracted FW, ISOVF, and magnetic susceptibility from the VTA, as these metrics have previously been reported to be associated with neuroinflammation ([35–37](#)). While both FW

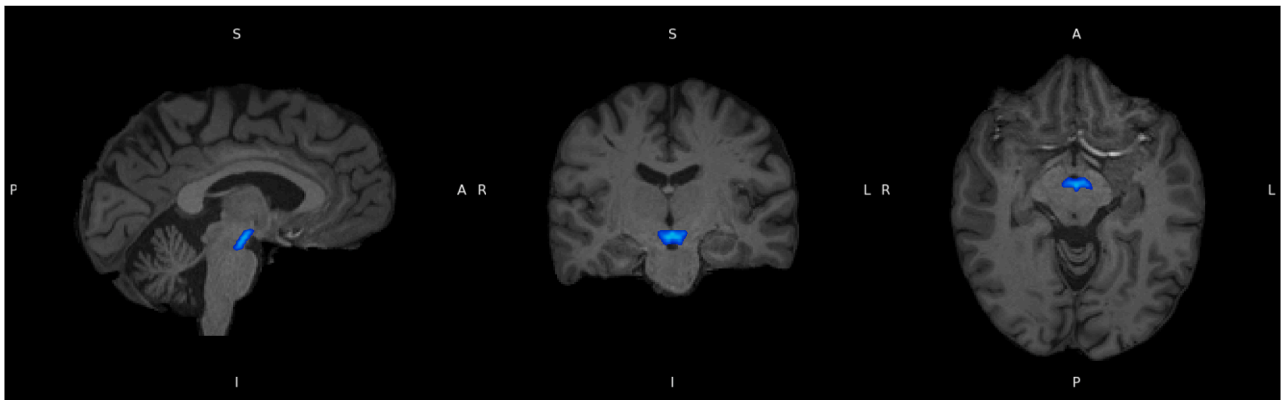
**Table 1. Demographic and VTA MRI Metrics in HC Participants and Individuals With ICD-10 MDD History**

Variable	Total, <i>N</i> = 32,495	HCs, <i>n</i> = 3807	MDD History, <i>n</i> = 3807	<i>p</i> Value
Sex, Female	16,806 (51.7%)	2435 (64%)	2431 (63.9%)	–
Age, Years	63.9 (7.7)	62.4 (7.6)	62.5 (7.6)	.788
BMI	26.2 (3.9)	26.9 (4.2)	26.9 (4.2)	.740
RDS Score	5.2 (1.7)	5.1 (1.6)	6.5 (2.6)	<.001*
MRI-Derived Metrics				
ICVF	0.533 (0.074)	0.531 (0.073)	0.527 (0.072)	.044**
ODI	0.295 (0.073)	0.294 (0.073)	0.298 (0.073)	.025**
ISOVF	0.374 (0.114)	0.366 (0.115)	0.375 (0.115)	.001***
FW	0.456 (0.101)	0.451 (0.102)	0.459 (0.103)	.001*
Magnetic susceptibility	0.395 (0.072)	0.393 (0.071)	0.394 (0.072)	.457
Volume ratio	$7.025 \times 10^{-4}$ ( $1.98 \times 10^{-4}$ )	$7.135 \times 10^{-4}$ ( $2.057 \times 10^{-4}$ )	$7.076 \times 10^{-4}$ ( $2.006 \times 10^{-4}$ )	.201

Values are presented as *n* (%) or mean (SD). Magnetic resonance imaging (MRI) metrics from the ventral tegmental area (VTA) provide insights into microstructural and inflammatory characteristics. *p* Values indicate the significance of differences between healthy control (HC) and major depressive disorder (MDD) history groups.

\**p* < .05, \*\**p* < .01, \*\*\**p* < .001.

BMI, body mass index; FW, free water; ICVF, intracellular volume fraction; ISOVF, isotropic volume fraction; ODI, orientation dispersion index; RDS, Recent Depressive Symptoms.



**Figure 2.** Ventral tegmental area (in blue), delineated using the Levinson-Bari Limbic Brainstem Atlas, displayed on coregistered participant T1 images. A, anterior; I, inferior; L, left; P, posterior; R, right; S, superior.

and ISOVF measure free water, ISOVF requires advanced protocols, while FW is more clinically accessible. ICVF and ODI were also extracted to investigate potential microstructural changes associated with acute and prolonged neuroinflammation. VTA volume was adjusted for total intracranial volume to account for individual brain size differences.

While our primary focus was on the VTA, we conducted exploratory analyses in additional regions implicated in MDD, including the orbitofrontal cortex, insula, amygdala, hippocampus, ventral anterior cingulate cortex (ACC), medial prefrontal cortex, posterior cingulate cortex, precuneus, angular gyrus, dorsolateral prefrontal cortex, dorsal ACC, prefrontal cortex, caudate, and nucleus accumbens (38). Full results are presented in Tables S3 to S5.

### Statistical Analysis

Group differences on demographic and clinical variables were tested using  $t$  tests and  $\chi^2$  tests. To examine associations between depression diagnosis and MRI-derived VTA metrics (FW, ISOVF, QSM, ICVF, ODI, volume), we conducted one-way analyses of variance (ANOVAs) with group (MDD/HC) as the between-subjects factor. Bonferroni correction was applied, setting the corrected alpha level at 0.0083 (0.05/6 comparisons), and effect sizes were reported as  $\eta_p^2$ . Groups were matched on age, sex, and BMI.

To assess whether VTA MRI metrics predicted current depressive symptom severity, we performed linear regression using MRI markers as predictors and RDS scores as the outcome. VTA volume was not included in the regression models due to concerns about the accuracy of atlas-based volume estimates in small brainstem structures and to avoid introducing additional noise into the analysis. To ensure adequate representation of individuals with high depressive symptom severity, we identified a reference group with RDS scores  $\geq 13$  (indicative of high symptom severity). Given the limited number of participants in this high-severity reference group (<100 participants), we limited the sampling of participants with lower RDS scores to a maximum of 100 participants per RDS score. The selection was pseudorandomized and iterated (50,000

permutations) to closely match age and sex with the high-symptom group. Residual confounding was addressed by controlling for age, age<sup>2</sup>, sex, BMI, and the log-transformed duration and number of depression episodes (capped at 100) (Tables S5 and S6). Log transformation was applied to reduce skew and enhance sensitivity to lower-range values. Standardized coefficients were reported for effect size comparison. Multicollinearity was checked using the variance inflation factor (VIF), and correlations among predictors were explored (Figure S2).

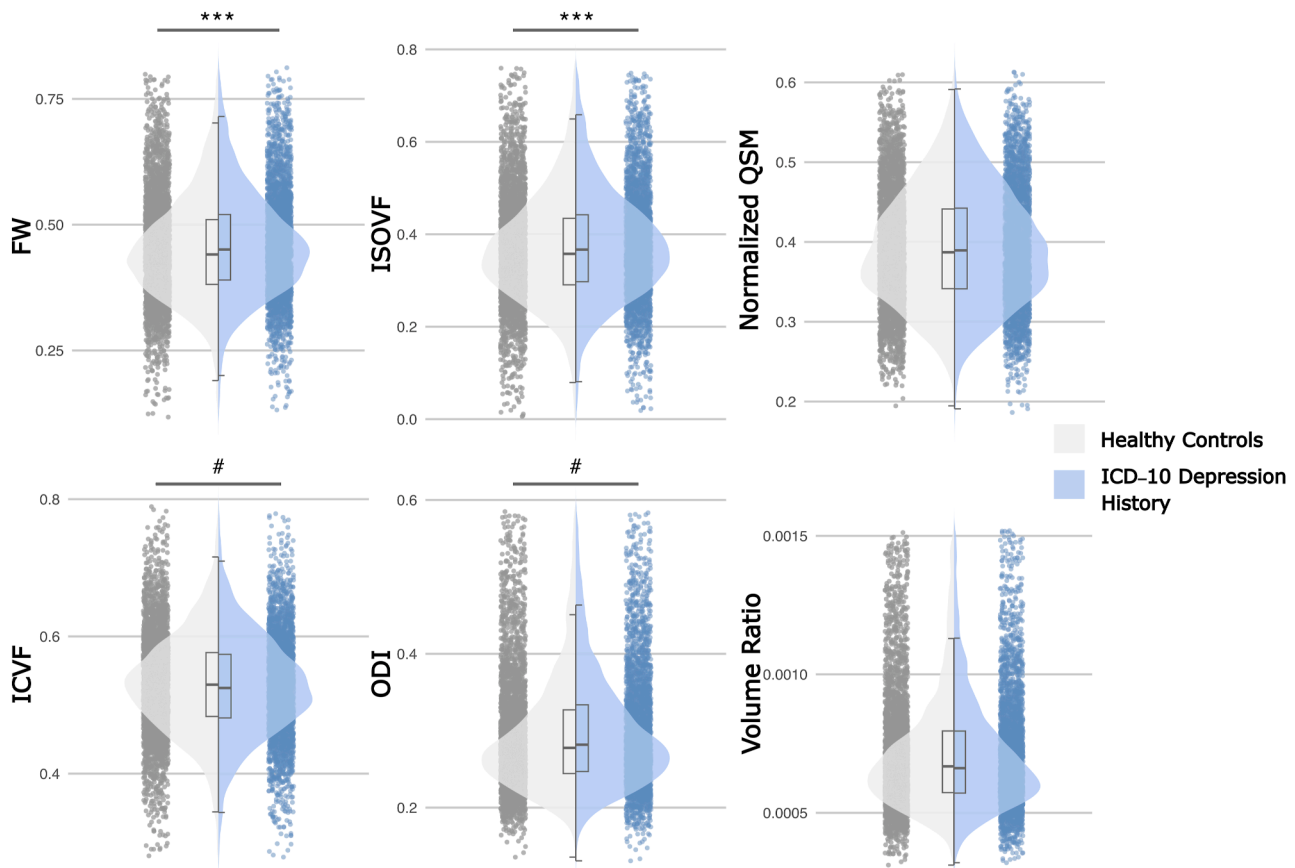
## RESULTS

### Association Between VTA-Derived MRI Markers and MDD Diagnosis

As shown in Table 1, our groups consisted of 64% females, were  $62.5 \pm 7.6$  years of age, and had a BMI of  $26.9 \pm 4.2$ . No significant differences were found in age, sex, or BMI, but the depression history group had significantly higher RDS scores than the HC group ( $t_{6327} = 28.3$ ,  $p < .001$ ).

ANOVAs revealed that the MDD group exhibited significantly higher FW ( $F_{1,7612} = 12.29$ ,  $p < .001$ ,  $\eta_p^2 = 0.002$ ) and ISOVF ( $F_{1,7612} = 10.26$ ,  $p = .001$ ,  $\eta_p^2 = 0.001$ ) compared with the HC group, suggesting increased extracellular inflammation-related processes in the VTA among individuals with a history of depression. The findings of higher ICVF ( $F_{1,7612} = 4.06$ ,  $p = .044$ ,  $\eta_p^2 = 5.34 \times 10^{-4}$ ) and ODI ( $F_{1,7612} = 4.99$ ,  $p = .025$ ,  $\eta_p^2 = 6.56 \times 10^{-4}$ ) in the MDD group compared with the HC group did not survive Bonferroni correction. No significant group differences were observed for volume ( $F_{1,7612} = 1.63$ ,  $p = .2$ ,  $\eta_p^2 = 0.0002$ ) or magnetic susceptibility ( $F_{1,7612} = 0.55$ ,  $p = .457$ ,  $\eta_p^2 = 7.2 \times 10^{-5}$ ). Group means and distributions are visualized in Figure 3. Across the additional brain regions examined, relative to HC participants, individuals with a history of MDD showed significantly higher ISOVF and FW in several cortical and subcortical areas, including the orbitofrontal cortex, insula, hippocampus, and caudate, indicating that greater extracellular water content and potential microstructural alterations associated with depression extend to these regions.

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**Figure 3.** Group comparisons between healthy control participants and individuals with an ICD-10 history of major depressive disorder. \*\*\* represents significant group differences that were sustained even after multiple comparison correction ( $p$  value  $\leq .001$ ), and # indicates significant group differences before multiple corrections. Magnetic resonance imaging (MRI) metrics from the ventral tegmental area (VTA), including intracellular volume fraction (ICVF), orientation dispersion index (ODI), isotropic volume fraction (ISOVF), free water (FW), quantitative susceptibility mapping (QSM), and standardized volume ratio.

### VTA-Derived MRI Markers as Predictors of Acute Depression Severity

The final subsample included 1151 participants, comprising individuals from the high-symptom reference group (RDS score  $\geq 13$ ) and the best-matched lower-symptom group (RDS score  $< 13$ ). Descriptive statistics for the full sample and subgroups are shown in Table 2. The total subsample consisted of 58% females, with a mean age of  $60.9 \pm 7.4$  years and a BMI of  $27.2 \pm 4.4$ .

FW (VIF = 34.12) and ISOVF (VIF = 33.08) showed high multicollinearity in the regression model, supported by their strong correlation ( $r = 0.96$ ,  $p < .001$ ), suggesting that they reflect similar processes, allowing them to be used interchangeably. To reduce redundancy and for consistency with other NODDI metrics, FW was excluded and ISOVF retained as a proxy for extracellular inflammation. After excluding FW, ISOVF VIF decreased to 1.82, and all other variables (except age and age<sup>2</sup>) had VIFs  $< 5$ , indicating that multicollinearity was resolved. See Figure S2 for correlation details.

The linear regression model was significant ( $F_{10,1140} = 19.73$ ,  $p < .001$ ), explaining 14% of the variance in depression severity

( $R^2 = 0.148$ , adjusted  $R^2 = 0.14$ ). As shown in Figure 4A, the model showed moderate fit with slight overestimation near the center of the distribution. Standardized beta coefficients are presented in Figure 4B. Both ICVF ( $\beta = 0.29$ ;  $p = .017$ ; 95% CI, 0.053 to 0.53) and ODI ( $\beta = 0.4$ ;  $p = .007$ ; 95% CI, 0.11 to 0.7) were positively associated with depressive symptom severity, indicative of an increase in microstructural changes with higher depression severity. Elevated ODI values suggest increased microstructural changes in the extracellular space, while higher ICVF values indicate greater neurite density associated with worsening depression symptoms. ISOVF showed a negative association with depression severity ( $\beta = -0.28$ ;  $p = .033$ ; 95% CI,  $-0.53$  to  $-0.02$ ), indicating that decreased NODDI FW was linked to greater symptom severity. In contrast, magnetic susceptibility was not significantly associated with depressive symptom severity.

Sex was a significant predictor of depression severity, with males exhibiting lower depressive symptom severity compared with females ( $\beta = -0.4$ ;  $p < .001$ ; 95% CI,  $-0.6$  to  $-0.2$ ). Additionally, BMI was positively associated with depressive severity ( $\beta = 0.38$ ;  $p < .001$ ; 95% CI, 0.18 to 0.57), suggesting that higher BMI may be linked to increased



**Table 2. Demographic and VTA MRI Metrics Across Subgroups With Varying Levels of Recent Depressive Symptoms**

Variable	Total Subsample, <i>n</i> = 1151	Best Matched, RDS Score < 13, <i>n</i> = 900	Reference, RDS Score ≥ 13, <i>n</i> = 251
Sex, Female	667 (57.9%)	506 (56.2%)	161 (64.1%)
Age, Years	60.9 (7.4)	61.3 (7.5)	59.5 (6.9)
BMI	27.2 (4.4)	26.9 (4.2)	27.9 (5)
RDS Score	9.4 (3.5)	8 (2.6)	14.4 (1.2)
MRI-Derived Metrics			
ICVF	0.529 (0.074)	0.527 (0.074)	0.533 (0.074)
ODI	0.297 (0.077)	0.297 (0.078)	0.296 (0.076)
ISOVF	0.371 (0.115)	0.375 (0.115)	0.356 (0.115)
FW	0.455 (0.103)	0.459 (0.103)	0.441 (0.104)
Magnetic susceptibility	0.392 (0.07)	0.391 (0.069)	0.394 (0.073)
Volume ratio	$7.128 \times 10^{-4}$ ( $2.01 \times 10^{-4}$ )	$7.125 \times 10^{-4}$ ( $2.01 \times 10^{-4}$ )	$7.139 \times 10^{-4}$ ( $2.016 \times 10^{-4}$ )

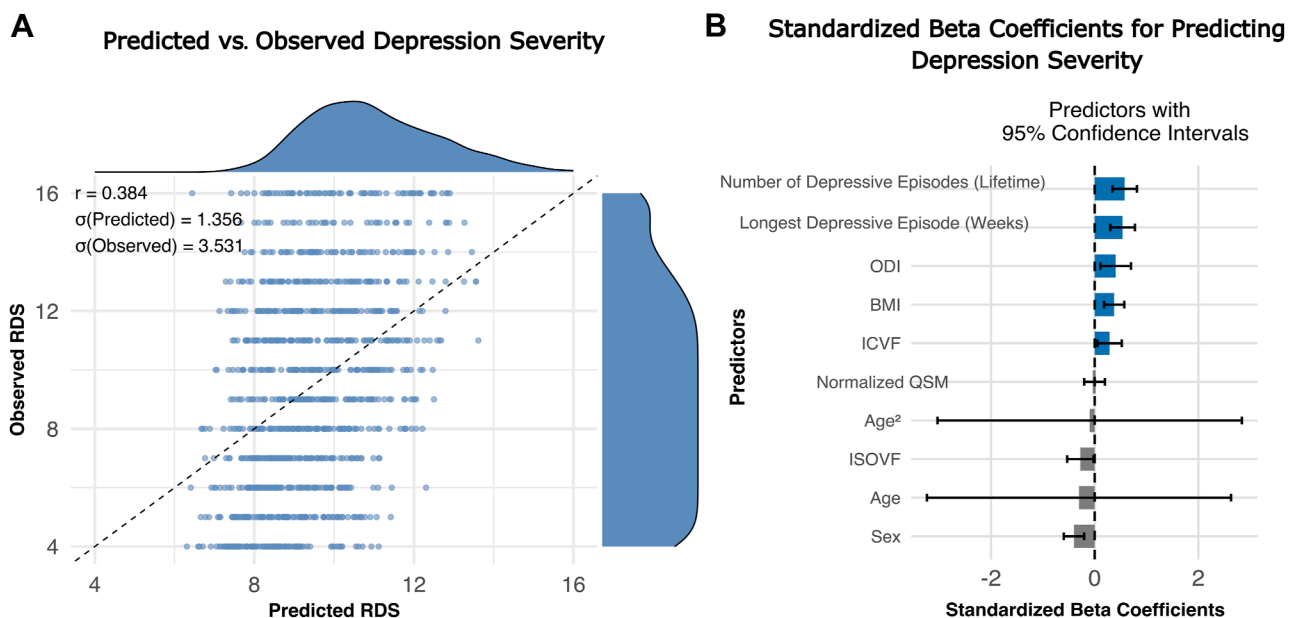
Values are presented as *n* (%) or mean (SD). Main analysis was run on the total subsample.

BMI, body mass index; FW, free water; ICVF, intracellular volume fraction; ISOVF, isotropic volume fraction; MRI, magnetic resonance imaging; ODI, orientation dispersion index; RDS, Recent Depressive Symptoms; VTA, ventral tegmental area.

severity of depressive symptoms. Notably, the number of lifetime depressive episodes ( $\beta = 0.58$ ;  $p < .001$ ; 95% CI, 0.34 to 0.82) and the duration of the longest depressive episode in weeks ( $\beta = 0.54$ ;  $p < .001$ ; 95% CI, 0.30 to 0.78) were also significant predictors, indicating that greater depression chronicity was associated with more severe current symptoms. The covariates age and age<sup>2</sup> were not significantly associated with depressive symptom severity.

## DISCUSSION

This study provides novel insights into the neurobiological correlates of depression, revealing a complex relationship between structural and inflammatory markers in the VTA and depressive symptomatology. Our findings demonstrate that individuals with a diagnosis of MDD exhibited significantly higher extracellular FW in the VTA compared with matched HCs, suggesting extraneuronal pathology potentially linked to



**Figure 4.** (A) Scatter plot depicting the relationship between predicted and observed depression severity scores (Recent Depressive Symptoms [RDS]) derived from the linear regression model. Each point represents an individual participant, with the dashed identity line ( $y = x$ ) included for reference. The marginal density plots illustrate the distribution of predicted and observed scores. The Pearson correlation coefficient ( $r$ ), together with the standard deviations of the predicted and observed RDS values, is annotated. (B) Standardized beta coefficients from the regression model predicting depression severity. Bars represent the effect size of each predictor, with 95% CIs indicated by error bars. Positive predictors (e.g., orientation dispersion index [ODI], body mass index [BMI], intracellular volume fraction [ICVF]) are shown in blue, while negative predictors are shown in gray. The dashed horizontal line at 0 represents the null effect. Variables are ordered by effect size for clarity. ISOVF, isotropic volume fraction.

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inflammatory processes. While these inflammatory markers were elevated, no structural alterations or differences in iron deposition (estimated via QSM) were observed between groups. Notably, MRI markers sensitive to inflammation and structural changes in the VTA predicted acute depressive symptom severity, independent of prior history of MDD diagnosis, with higher ICVF and ODI and lower ISOVF associated with more severe depressive symptoms. In contrast, magnetic susceptibility did not predict symptom severity, reinforcing the absence of chronic inflammation-related effects such as cell damage or iron accumulation.

ISOVF and FW are MRI markers sensitive to neuroinflammation and edema (28,29). Although derived from different diffusion models (3-compartment NODDI for ISOVF and 2-compartment tensor modeling for FW), both quantify freely diffusing water and can reflect extracellular fluid increases associated with inflammatory processes such as vasogenic edema and associated cellular responses (28). Therefore, elevated ISOVF and FW in individuals with a history of MDD are consistent with our prediction that major depression is associated with elevated inflammation markers in the VTA. In contrast, while FW and ISOVF were elevated in individuals with MDD, we observed that acute depression severity in the general population was associated with lower ISOVF. This dissociation suggests these two states of depression—historical diagnosis of major depression versus acute depressive symptoms across the general population—may reflect distinct biological processes and inflammatory states. Yi *et al.* (39) highlighted important distinctions between acute and chronic inflammation, showing that acute neuroinflammation is marked by increased hindered water diffusion due to increases in the number of activated hyperramified microglia, while chronic inflammation is marked by reduced hindered diffusion due to overactivation of microglia, which increases their density. Thus, our findings of lower ISOVF may reflect acute inflammation processes, while the elevated ISOVF and FW may be indicative of chronic inflammatory processes in major depression.

Increased FW is often associated with psychiatric symptom severity and neuroinflammation in psychiatric and neurological disorders (22,40–42), although findings in depression remain mixed. For example, elevated FW has been reported in patients with poststroke depression in reward-related regions (22), suggesting vulnerability of these areas to inflammation. Conversely, Bergamino *et al.* (40) found no FW differences between HCs and unmedicated patients with MDD. Consistent with our findings, Althubaiti *et al.* (43) reported reduced ISOVF in the gray matter of individuals with depression, particularly in the ACC of individuals with low peripheral inflammation, although this difference was not statistically significant across the entire depression group. The ACC, like the VTA, has shown decreased functional MRI activation and increased microglial expression in PET studies (44). However, no significant ISOVF differences were found in the prefrontal or insular cortices across inflammation subgroups or overall depressed versus control groups (43). These findings suggest that FW and ISOVF changes may reflect distinct, localized processes, potentially influenced by acute or chronic inflammation associated with depression. Given the reward network's sensitivity to neuroinflammatory processes and its

central role in dopamine signaling (9–11), these regions may exhibit differential responses to such changes, with prolonged depressive episodes possibly impacting the VTA's structural integrity in a manner distinct from acute depressive symptoms.

Our results indicate that the VTA is also susceptible to microstructural aberrations possibly linked to inflammation in depression. While ICVF and ODI differences did not survive Bonferroni correction in group comparisons, our regression analysis identified both metrics as independent predictors of depression severity. Elevated ICVF, which is typically interpreted as increased neurite density and intracellular diffusion, has also been associated with glial swelling and microglial reactivity, key features of depression (20,45). Similar increases in ICVF have been reported in the striatum following interferon alpha-induced inflammation (46). Although fatigue (a symptom of depression) increased, no significant rise in depression severity was observed, suggesting that inflammation affects reward circuitry even in the absence of amplifying mood symptoms (46). This may explain the vulnerability of the VTA and striatum to inflammation in depression, which is consistent with our findings of increased ICVF in the VTA, a structure closely linked to the brain's reward system, reinforcing its vulnerability to inflammatory processes and depression. Higher ODI levels in the VTA may reflect microstructural processes in the extracellular space, where microglia modulate immune defense. Yi *et al.* (39) demonstrated that increased ODI correlated with microglial density, further suggesting that microglial activation could be driving these microstructural changes. Similarly, Ota *et al.* (47) reported reduced ODI in the thalamus of individuals with MDD, and postmortem studies found decreased astrocytic markers in the thalamus of deceased individuals with MDD, lending additional support for the involvement of glial cells in depression (48).

Our study found no significant differences in VTA volume between individuals with depression and healthy individuals. This contrasts with findings by Morris *et al.* (49), who reported increased VTA volume linked to alterations in neuromelanin and dopamine activity in individuals with mood disorders. The absence of volumetric differences in our study suggests that the observed microstructural changes, such as higher ICVF and ODI, may occur independently of gross volumetric changes.

Even though QSM is sensitive to the effects of prolonged inflammation, our study found no significant differences in magnetic susceptibility between individuals with major depression and HCs in the VTA; it was also not a predictor of acute depressive severity. This suggests that processes such as iron deposition, demyelination, or cell death, typically associated with chronic inflammation, may not significantly impact VTA structure as hypothesized. However, Yao *et al.* (20) reported higher susceptibility values correlating with current depression severity in the putamen and thalamus. Duan *et al.* (19) found that magnetic susceptibility in the thalamus, hippocampus, and putamen correlated with depression duration rather than current symptom severity, suggesting that microstructural changes accumulate over time. The discrepancy between our VTA findings and previous work in other regions may reflect the distinct cellular architecture and inflammatory vulnerability of different brain areas.

The VTA, primarily composed of dopaminergic neurons and their supporting glial cells, may respond differently to inflammatory processes compared with regions such as the putamen and thalamus, which have different cellular compositions and metabolic demands. While Yao *et al.* (20) and Duan *et al.* (19) demonstrated susceptibility changes in striatal and limbic regions, our findings suggest that the VTA may be more resilient to chronic inflammatory processes that lead to iron accumulation or cellular damage. Instead, the VTA appears to show more dynamic changes in the extracellular space (as indicated by FW measures) that may reflect acute rather than chronic inflammatory states.

While statistically significant, our findings had small effect sizes, reflecting the multifactorial nature of depression. Depression is influenced by numerous factors, including sex, lifestyle factors, genetic predisposition, and metabolic variables such as BMI (50), which explained a substantial portion of the variance in our models. Our findings showed that females were significantly more likely to have higher depressive symptoms, consistent with previous literature indicating a higher prevalence of depression in females, potentially due to factors such as reporting tendencies and hormonal influences (51). BMI was also a positive predictor of depression severity, consistent with studies that have shown obesity to be associated with elevated cytokine levels, reduced physical activity, and heightened depression risk (52,53).

Additionally, depression history and chronicity, as measured by the number and duration of previous depressive episodes, were significant predictors of current symptom severity. This is consistent with previous research showing that recurrent and prolonged episodes are associated with greater functional impairment and an increased risk of future depressive episodes (54). However, we acknowledge that these measures were based on retrospective self-report and did not include precise timing information (e.g., when episodes occurred), which may limit the reliability and interpretability of these findings. These results underscore the complex interplay between neuroinflammation, lifestyle factors, and depressive symptoms, suggesting that future studies should consider these factors in more detail to fully understand the multifactorial nature of depression.

One important limitation of our study is the cross-sectional design, which limits our ability to establish causal relationships between neuroinflammation, VTA microstructural changes, and depression severity. Additionally, while a significant portion of our participants experienced multiple major depressive episodes between their initial ICD-10 depression diagnosis and the time of scanning, the average time since their first diagnosis was approximately  $27 \pm 11.15$  years prior to the scan. Future longitudinal studies should explore the relationship between recurrent depressive episodes and the interplay of chronic and acute depression on brain structure. Furthermore, while DWI and QSM provide valuable insights into brain structure and inflammation, they are indirect measures. Integrating these methods with more direct biomarkers, such as cerebrospinal fluid cytokine levels or PET imaging, would provide stronger evidence of neuroinflammation's role in depression. Lastly, while the atlas-based registration captures the VTA well, it encompasses small portions of neighboring structures such as the red nucleus and substantia

nigra, which is a common limitation when studying such small, anatomically adjacent nuclei. As a result, the diffusion metrics attributed to the VTA may partially reflect signal from these adjacent regions.

## Conclusions

Our findings reveal distinct patterns of microstructural changes in the VTA associated with both major depression history and current depressive symptom severity, suggesting different underlying pathophysiological mechanisms. The elevated FW markers observed in individuals with a major depression history, coupled with the relationship between NODDI metrics and current symptom severity, point to dynamic inflammatory processes that may differ between acute and chronic states of depression. While our results support the role of neuroinflammation in depression, the absence of QSM changes suggests that the VTA may be more resilient to chronic inflammatory processes compared with other brain regions. These findings could ultimately inform more targeted therapeutic approaches that consider the distinct biological processes underlying depression.

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All the neuroimaging preprocessing and analyses conducted in this study involved the use of publicly available toolboxes and resources. Specifically, the following software tools were used: FreeSurfer (version 7.4.1) (55,56), FSL (version 6.0.7.4) (57,58), FSLeves (version 1.10.4) (59), MRtrix3 (version 3.0.3) (60), and ANTs (version 2.3.5) (33). For statistical analysis, both ANOVAs and regression analyses were run through R and RStudio (61), using *dplyr*, *ggplot2*, *ggpubr*, *skmr*, *plotly*, *htmlwidgets*, *MatchIt*, *gghalves*, *psych*, *GGally*, *tidyverse*, *car*, *kableExtra*, *officer*, *tidtk*, and *scales* packages and functions (62–77). The FW imaging code, developed by Pasternak *et al.* (28), is not publicly available but can be obtained from the authors upon request. All other code and materials used in this study are available at [https://github.com/Psychedelics-Clinical-Neuroimaging-Lab/UKB\\_VTA\\_Inflammation\\_Data\\_Prep\\_and\\_Stat/tree/main](https://github.com/Psychedelics-Clinical-Neuroimaging-Lab/UKB_VTA_Inflammation_Data_Prep_and_Stat/tree/main). Due to ethical considerations, raw data from the UK Biobank cannot be shared directly.

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## ARTICLE INFORMATION

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