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# Brain imaging derived phenotypes: a biomarker for the onset of inflammatory bowel disease and a potential mediator of mental complications

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**Background and aims:** Inflammatory bowel disease (IBD), mainly categorized into Crohn's disease (CD) and ulcerative colitis (UC), is a chronic relapsing gastrointestinal disorder that significantly impairs patients' quality of life. IBD patients often experience comorbidities such as anxiety and depression, and the underlying mechanisms and treatment strategies remain areas of investigation.

**Methods:** We conducted a Mendelian randomization(MR) analysis utilizing brain image derived phenotypes (IDP) from the UK Biobank database to investigate the causal relationships between IBD and alterations in brain structural morphology and connectivity of neural tracts. This study aimed to identify biological evidence linking IBD to psychiatric disorders such as anxiety and depression.

**Results:** Specifically, the volume of grey matter in the Left Frontal Orbital Cortex exhibited a negative association with the onset of Crohn's disease (odds ratio (OR) [95% confidence interval (CI)]: 0.315[0.180~0.551], adjusted P=0.001), while the volume of the superior frontal cortex in the right hemisphere showed a positive correlation with the development of Ulcerative colitis (OR [95% CI]: 2.285 [1.793~2.911], adjusted P<0.001), and the volume of lateral occipital cortex in the left hemisphere demonstrated a positive relationship with Crohn's disease onset (OR [95% CI]: 1.709[1.671~1.747], adjusted P<0.001). In the context of reverse causality, the onset of UC or CD has led to alterations in imaging derived phenotypes associated with five disorders (anxiety, depression, schizophrenia, bipolar disorder, pain) and three functions (memory, emotion, language).

**Conclusion:** Our study has demonstrated a causal relationship between IBD and IDPs. IDPs may serve as potential biomarkers for the progression of IBD and as predictive intermediaries for the development of neurological diseases in IBD patients.

#### KEYWORDS

Mendelian randomization, imaging derived phenotypes, ulcerative colitis, Crohn's disease, inflammatory bowel disease

## **1** Introduction

Inflammatory Bowel Disease (IBD), encompassing ulcerative colitis (UC) and Crohn's disease (CD), represents a chronic, relapsing autoimmune gastrointestinal disorder that has now emerged as a global health concern. Over the past 30 years, there has been a global rise in the incidence, prevalence, and mortality rates of IBD. In the next two decades, the burden of IBD is projected to persistently increase due to population growth and aging, particularly in developing countries (1).. The primary clinical manifestations of the disease include fever, abdominal pain, bloody stools, fatigue, malnutrition, as well as extraintestinal manifestations such as arthritis and uveitis, significantly impacting the quality of life. Furthermore, due to the prolonged suffering from the disease, IBD patients often have a higher susceptibility to mental disorders, including anxiety and depression (2). Current research on the gutbrain axis involves the study of IBD's clinical presentations, such as pain, stress states, and emotional changes, inducing alterations in brain structure (3). Some voxel-based morphometry studies have demonstrated differences in certain brain structures, such as the frontal and temporal lobes and the amygdala, between patients with CD and UC and healthy individuals (4). However, due to variations in the delineation of brain regions and the different stages of disease chosen in previous research, it remains challenging to precisely determine how the disease affects brain structure. Therefore, we employed a Mendelian randomization approach to investigate the causal relationships between CD and UC and brain imaging derived phenotypes (IDPs) at the genetic level.

Mendelian randomization (MR) is an epidemiological approach that uses summary data from existing genome-wide association studies (GWAS) to identify eligible genetic variants as instrumental variables for inferring causal relationships between exposures and outcomes (5). Randomized controlled trials (RCTs) are often considered the gold standard for establishing causal relationships in observational study conclusions. However, due to the complexity of experimental design, financial constraints, ethical considerations, and other factors, RCTs are not always feasible to conduct (6). In such cases, MR serves as a valuable method for simulating RCT studies. This method is based on Mendel's law, in which each trait is randomly allocated to independent individuals with allele variants, similar to the random allocation of individuals into control and exposure groups in an RCT. As genetic factors determine traits, the selection of appropriate instrumental variables can help mitigate the influence of common confounders on study outcomes and establish causal relationships (7).

In this study, we employed publicly available GWAS data to conduct two-sample bidirectional MR analyses to investigate the impact of UC and CD on brain IDPs. This study investigates the causal relationship between IBD and alterations in brain IDP, aiming to explore potential factors contributing to the occurrence of psychiatric symptoms in IBD patients. It emphasizes the significance of MRI in clinical practice, providing new evidence for the diagnosis of IBD and the prevention of neuropsychiatric complications.

# 2 Methods

## 2.1 Data sources

The GWAS summary data for UC and CD, stratified after inflammatory bowel disease, were obtained from the International IBD Genetics Consortium (IIBDGC). This study analyzed data from 86,640 European participants and 9,846 individuals of non-European ancestry (8), employing quality control, genotype data imputation, meta-analysis, and trans-ethnic analysis to investigate the genetic basis and heterogeneity of Crohn's disease and ulcerative colitis in-depth. The diagnosis of IBD was based on recognized radiological, endoscopic, and histopathological evaluations. All included cases met the clinical criteria for IBD. Recruitment of study subjects was approved by the ethics committees or institutional review boards of all individual participating centers or countries. All study participants provided written informed consent. In our study, we exclusively utilized datasets of European ancestry for UC and CD. Both datasets included both male and female participants, with 6,968 UC cases and 20,464 controls in the UC dataset, and 5,956 CD cases and 14,927 controls in the CD dataset.

The GWAS summary data for brain imaging-derived phenotypes were obtained from a meta-analysis of 3,144 IDPs in 8,428 participants from the UK Biobank, including both male and female individuals (9). High-throughput sequencing technologies were employed, with filters applied to remove variants with a minor allele frequency (MAF) below 0.1% and imputation information scores below 0.3, excluding variants with Hardy-Weinberg equilibrium P-values <10<sup>-7</sup> and MAF <0.1%. They identified 148 significant association clusters between single nucleotide polymorphisms (SNPs) and IDPs, with particularly notable and interpretable clusters encompassing genes related to iron transport and storage, white matter microstructure and pathology, brain development, and plasticity. For the definition of each IDP, segmentation was conducted using FreeSurfer v6.0.0 on the Desikan-Killiany-Tourville Atlas (referred to as DKT) and the Destrieux Atlas (referred to as a2009s). The UK Biobank has received approval from the Northwest Multi-center Research Ethics Committee (MREC) to acquire and disseminate participant data and samples (http://www.ukbiobank.ac.uk/ethics/), regulations that encompass the work in this study. All participants provided written informed consent. We included IDP phenotypes encompassing regional and tissue volumes (such as Cortex, thalamus, hypothalamus, basal ganglia, hippocampus and other structures), tfMRI activation, and white matter hyperintensity volume (see Supplementary Table 1). Our study solely extracted GWAS statistical data from previously published research, which is publicly accessible and does not require additional ethical approvals or informed consent. Based on the information from the two data set sources, there is no sample overlap in the MR analysis of this study.

## 2.2 Study approach

In this study, we utilized a European population GWAS summary datasets, conducted stratified analyses for two subtypes of inflammatory bowel disease, and performed reverse analyses by altering exposures and outcomes. We applied three MR hypotheses to select eligible instrumental variables and employed various analytical methods, including inverse variance weighting (IVW), MR-Egger regression, weighted median, weighted mode, and MR-RAPS. Sensitivity analyses, such as testing for heterogeneity, horizontal pleiotropy, and leave-one-out analysis, were performed to ensure the robustness of our results. We defined the causal direction from imaging derived phenotypes to inflammatory bowel disease as forward, and the opposite as reverse.

## 2.3 Genetic instrument selection

We selected single nucleotide polymorphisms as instrumental variables and randomized exposure factors indicated by genetic variations into groups to validate the association between exposure and outcomes.

Initially, we extracted genetic variations significantly associated with exposures at a genome-wide significance level  $(5 \times 10^{-8})$ . To ensure the retrieval of an adequate number of SNPs for further investigation, we lowered the significance level to  $1 \times 10^{-6}$  in forward batches. To ensure the independence of SNPs and reduce the risk of multicollinearity in the model, we filtered out SNPs with linkage disequilibrium (LD) by setting the parameters to  $r^2 < 0.01$  and a physical distance of 10,000 base pairs. We estimated the strength of the SNP-exposure association using the formula  $F = beta^2/se^2$  and defined SNPs with F > 10 as non-weak instrumental variables, which were retained (10). When harmonizing data, we excluded unmatched ambiguous SNPs and palindromic SNPs. We queried each SNP for its phenotypic associations using PhenoScanner at a significance level of  $P < 5 \times 10^{-8}$  and removed SNPs related to confounding factors (11). We applied MR-PRESSO to test and exclude the horizontal pleiotropy outliers.

We used MR-Steiger to verify the causal direction represented by all SNPs, and SNPs indicating the incorrect direction were removed. Finally, we applied Bonferroni multiple testing correction (P < 0.05/n, where n refers to the number of remaining SNPs) to eliminate SNPs directly associated with the outcome.

Through the instrumental-variable selection process described above, the eligible SNPs included in the study met the following criteria: 1) The SNP should exhibit a strong correlation with the exposure factor; 2) The SNP should only influence the outcome through the exposure factor; 3) The SNP should not be associated with confounding factors (Figure 1).

## 2.4 Mendelian randomization analysis

We conducted analyses using five different MR methods, and for each batch of studies, we determined a primary analytical method conclusion based on the results of sensitivity analyses, which served as the final conclusion. Inverse variance weighted is a weighted linear regression without an intercept term. It derives the final causal estimate by analyzing the Wald ratio results for each SNP (12, 13). The weights in IVW are the inverse of the genetic association variance and the outcome. Using first-order weights, IVW estimation is equivalent to the two-stage least squares (2SLS) method. According to guidelines (14), in the absence of horizontal pleiotropy and heterogeneity, we employ the IVW method with a multiplicative random-effects model as the primary analysis approach. Weighted median and weighted mode methods calculate causal effects based on the majority valid assumption and plurality valid assumption, respectively (14, 15). As long as over 50% of genetic variants are valid IVs, the median of the weighted median ratio estimate will tend towards the true causal effect. When less than 50% of variants are valid instrumental variables, weighted mode can determine the true causal effect as long as there is no larger group of invalid instrumental variables with the same ratio estimate. When there is heterogeneity among SNPs, both the weighted median method and the IVW method are required to jointly support a significant conclusion. The MR-Egger method inherently assumes the presence of horizontal pleiotropy and estimates this effect using the intercept term in the model (16). When the intercept term in the MR-Egger regression exhibits statistical significance, we consider the existence of horizontal pleiotropy, and employ MR-Egger as the primary analytical approach. The MR-RAPS method is a common modeling approach that estimates causal effects using the likelihood-based probability profile under the assumption of pleiotropy following a normal distribution centered at zero with position variance (17). The null hypothesis (H<sub>0</sub>) for all the aforementioned MR methods is that there is no causal relationship between exposure and outcome, whereas the alternative hypothesis  $(H_1)$  posits a causal relationship between exposure and outcome. We applied FDR multiple testing correction to the significance P-values of MR results, using a threshold of corrected P < 0.05 to infer the likelihood of causal relationships.

## 2.5 Sensitivity analysis

To ensure the robustness of Mendelian randomization analysis results, we conducted various sensitivity analyses. We assessed heterogeneity among instrumental variables by calculating the weighted sum of squared differences between each variant estimate and the overall estimate using Cochran's Q test. SNP with a Q test P-value < 0.05 were considered heterogeneous. We estimated the overall causal direction using MR-Steiger (18). To determine if any strongly influential SNPs were present, we conducted a leave-one-out sensitivity test. The MR-Egger regression assumes adherence to the InSIDE (Instrument Strength Independent of Direct Effect) assumption and the NOME (No Measurement Error) hypothesis. We generated a funnel plot and calculated the I2 statistic to validate the adherence to these assumptions. When I2 is less than 90% and the primary analytical method is MR-Egger, a correction for causal estimates is necessary (16, 19). Finally, we computed the statistical power for each batch of



studies and excluded positive conclusions with power less than 80% to reduce the occurrence of statistical Type II errors (20).

## 2.6 Two-step mediation analysis

Following recommendations, we employed the neuropsychiatric GWAS dataset from FinnGen to conduct a twostep mediation analysis, which investigated the mediating effects of various IDPs on six neuropsychiatric disorders caused by UC or CD, and calculated the mediation effect ratio.

## 2.7 Visualization and statistical software

For each batch of MR analyses in this study, scatterplots, regression curve graphs, and forest plots of SNP effects were generated, which will be presented in the results. Circular heatmaps and forest plots for the overall results were also created. Some of the figures were created using Servier Medical Art (smart.servier.com), provided by Servier, and are licensed under a Creative Commons Attribution 3.0 unported license. Statistical analysis and visualization in this study were performed using R (version 4.1.2), with the application of R packages such as "TwoSampleMR," "MR-PRESSO," "mr.raps," "forestploter," and several foundational R packages. Flowcharts were created using Microsoft Office Powerpoint (version 2312).

# **3** Results

### 3.1 Selection of instrumental variables

In the forward MR analysis, we initially screened a total of 916 SNPs related to the IDPs. No weak instrument variables were identified. 8 SNPs were removed due to missing data in the outcome datasets, 115 SNPs were excluded as ambiguous or palindromic SNPs when harmonizing two datasets, and 7 SNPs, identified through PhenoScanner, were found to be related to confounding factors such as serum vitamin D levels (21) and depression (2), and were subsequently excluded (see Supplementary Table 4). MR-PRESSO test revealed 8 SNPs with horizontal pleiotropy. MR-Steiger test did not identify any SNPs with erroneous causal direction. After Bonferroni correction, 26 SNPs directly related to the outcome were removed. Ultimately, 752 eligible SNPs were included in the study (for detailed SNP data, see Supplementary Table 3).

In the reverse MR analysis, we initially screened a total of 23,000 SNPs related to the IDPs. No weak instrument variables were identified. 252 SNPs were removed due to missing data in the outcome database, 3,499 SNPs were excluded as ambiguous or palindromic SNPs when harmonizing two datasets, and 1,250 SNPs were excluded after a PhenoScanner search, as they were associated with confounding factors. MR-PRESSO test identified 10 SNPs with horizontal pleiotropy. MR-Steiger test found no SNPs with erroneous causal directions. After Bonferroni correction, 27 SNPs directly related to the outcome were removed. Ultimately, 17,962 eligible SNPs were included in the study.

## 3.2 IDPs as the cause of IBD onset

In this study, 252 IDPs were included in the analysis, including regional and tissue volume, tfMRI activation, and white matter hyperintensity volume. The number of SNPs for each IDP ranged from 1 to 9, and the detailed information for the IVs of the 252 IDPs is provided in the Supplementary Table. Through MR analysis, we identified 3 types of IDPs associated with the causality of CD or UC subtypes. Analyses for these batches could not calculate heterogeneity and pleiotropy due to fewer than 3 SNPs. Specifically, for each 1 standard deviation (SD) increase in the Volume of grey matter (VGM) in the Left Frontal Orbital Cortex,

the risk of CD decreased by 68% (odds ratio (OR) [95% confidence interval (CI)]: 0.315[0.180~0.551], adjusted P=0.001). For each 1 SD increase in the Volume of superior frontal in the right hemisphere, the risk of UC increased by 129% (OR [95% CI]: 2.285[1.793~2.911], adjusted P<0.001). Moreover, for each 1 SD increase in the Volume of lateral occipital in the left hemisphere, the risk of CD increased by 71% (OR [95% CI]: 1.709[1.671~1.747], adjusted P<0.001). The results of the MR analysis are depicted in Figure 2. Detailed information on the other batches is shown in the Supplementary Materials.

# 3.3 IBD promotes changes in mental complications and function-related IDPs

We conducted a functional query in the PubMed database for IDPs significantly altered due to the onset of IBD and found that the onset of UC or CD led to changes in IDPs related to five disorders (anxiety, depression, schizophrenia, bipolar disorder, pain) and three functions (memory, emotion, language)(See in Figure 3).

### 3.3.1 IDPs classified by mental complications

The pallidum is associated with emotion regulation and contributes to the pathogenesis of anxiety disorders (22), with its volume changing in relation to the onset risk of UC. We identified 10 IDPs located within 3 brain regions (temporal lobe, cerebellum, anterior cingulate cortex) associated with CD and UC. Prior studies have shown alterations in low-frequency oscillatory amplitudes of the cerebellum and inferior temporal lobe, an index reflecting spontaneous brain activity, in patients with depression (23). We observed that 8 IDPs located in the brain regions of the parietal lobe, frontal lobe, paracentral lobule, olfactory-insularhippocampus, and hippocampal parahippocampus were affected by the onset of CD and UC. The parietal lobe is associated with visual information processing, the frontal lobe is linked to decisionmaking and emotion processing, the olfactory-insularhippocampus region is involved in memory, and the paracentral lobule is implicated in sensory and motor control. Previous research suggests their potential involvement in the onset of schizophrenia (24, 25). We also observed that UC onset altered 1 IDP associated with Bipolar Disorder, and CD onset altered 1 IDP related to pain.



#### FIGURE 2

Forward Mendelian Randomization Analysis Results and Brain Region Schematic. Notes: Top left panel, Brain Region Schematic; Top right panel, Manhattan plot of positive IDP-associated GWAS data; Bottom panel, Forest plot of positive MR analysis. VGM, Volume of Gray Matter; VWM, Volume of White Matter; L, Left; R, Right; CD, Crohn's Disease; UC, Ulcerative Colitis; IVW, Inverse Variance Weighted; MRE, Multiplicative Random Effects Model; OR, Odds Ratio; CI, Confidence Interval; NSNP, Number of SNPs included in the study.

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CD to VGM R cerebellum X     11     IVW(MRE)     0.025 (0.003 - 0.047)     0.028       CD to VGM L cerebellum X     41     IVW(MRE)     0.031 (0.008 - 0.047)     0.037       UC to VGM L cerebellum X     31     IVW(MRE)     0.036 (0.011 - 0.061)     0.008       UC to VGM L cerebellum X     31     IVW(MRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM L infermo cof.usif cortex     31     IVW(MRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM L infermo grups post     31     IVW(MRE)     0.038 (0.007 - 0.068)     0.017       UC to VGM L infermo grups post     31     IVW(MRE)     0.037 (0.022 - 0.027)     -0.001       Schizophrenia     CD to VVM L paracentral area     41     IVW(MRE)     0.037 (0.023 - 0.026)     0.046       CD to VGM L hippocampus     41     IVW(MRE)     -0.053 (0.011 - 0.051)     0.006       CD to VGM L parachipp grups post     31     IVW(MRE)     -0.037 (0.014 - 0.059)     0.031       UC to VGM R parahipp grups post     31     IVW(MRE)     -0.037 (0.014 - 0.059)     0.008       UC to VGM R parahipp grups post     31     IVW(MRE)	CD to VGM R cerebellum X CD to VGM R cerebellum X CD to VGM L cerebellum X UC to VGM L cerebellum X UC to VGM L tempeblum X UC to VGM L inft temp gyrus post UC to VGM L inft temp gyrus tempocc UC to VGM L inft temp gyrus tempocc CD to VGM L cerebellum X hrenia CD to VGM L caracentral area CD to VGM L entoRinal area CD to VGM R hippocampus UC to VGM R hippocampus	area	41 41 31 31 31 31 31	IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE)		III	0.025 (0.003 ~ 0.047) 0.026 (0.004 ~ 0.047) 0.031 (0.008 ~ 0.054)	0.028 0.037 0.008	86.9 91.1
CD to VWM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVV(IMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVVIMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVVIMRE)     Image: CD to VMM R rostralanteriorcingulate area     41     VVVVIMRE	CD to WMR R rostralanteriorcingulate a CD to VGM L cerebellum X UC to VGM L cerebellum X UC to VGM L temp occ fusif cortex UC to VGM R int femp gyrus post UC to VGM R int femp gyrus tempocc UC to VGM R cerebellum X hrenia CD to VWM L paracentral area CD to VWM L entoRinal area CD to VGM R hippocampus LIC to VGM R hippocampus	area	41 41 31 31 31 31	IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE)		II	0.026 (0.004 ~ 0.047) 0.031 (0.008 ~ 0.054)	0.037	91.1
CD to VGM L cerebellum X     41     VVV(IMRE)     Image: CD to VGM L cerebellum X     0.031 (0.008 - 0.054)     0.008       UC to VGM L temp occ fusif cortex     31     IVVV(IMRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM L temp occ fusif cortex     31     IVVV(IMRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM L temp ogrus person     31     IVVV(IMRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM L temp ogrus person     31     IVVV(IMRE)     0.038 (0.007 - 0.065)     0.030       UC to VGM C erebellum X     31     IVVV(IMRE)     0.038 (0.007 - 0.065)     0.017       UC to VGM R carebellum X     31     IVVV(IMRE)     0.037 (0.014 - 0.056)     0.019       CD to VMM L entoRinal area     41     IVVV(IMRE)     -0.032 (-0.0560.066)     0.045       CD to VGM R parahipp grups post     31     IVVV(IMRE)     -0.037 (0.014 - 0.051)     0.006       UC to VGM R parahipp grups post     31     IVVV(IMRE)     -0.037 (0.014 - 0.053)     0.003       UC to VGM R parahipp grups post     31     IVVV(IMRE)     -0.032 (-0.018 - 0.146)     0.024       Bipolar Disorder     UC	CD to VGM L cerebellum X UC to VGM L cerebellum X UC to VGM L cerebellum X UC to VGM L temp occ fusif cortex UC to VGM R inf temp gyrus tempocc UC to VGM R cerebellum X trenia CD to VGM L paracentral area CD to VGM L entoRinal area CD to VGM R hippocampus UC to VGM R hippocampus		41 31 31 31 31	IVW(MRE) IVW(MRE) IVW(MRE) IVW(MRE)			0.031 (0.008 ~ 0.054)	0.008	
UC to VGM L cerebellum X     31     VVV(INRE)	UC to VGM L cerebellum X UC to VGM L temp occ fusif cortex UC to VGM L inf temp gyrus post UC to VGM L inf temp gyrus tempocc UC to VGM L erebellum X hrenia CD to VWM L paracentral area CD to VWM L entoRinal area CD to VGM L entoRinal area CD to VGM L entoRinal area UC to VGM R hippocampus UC to VGM R hippocampus		31 31 31 31	IVW(MRE) IVW(MRE) IVW(MRE)			. ,		97.2
UC to VGM L temp occ fusif cortex 31 IVW(MRE) III 0.036 (0.007 - 0.065) 0.030 UC to VGM L inft temp gyrus prost 31 IVW(MRE) III 0.038 (0.009 - 0.067) 0.022 UC to VGM L inft temp gyrus tempcoc 31 IVW(MRE) III 0.038 (0.009 - 0.067) 0.022 UC to VGM L inft temp gyrus tempcoc 31 IVW(MRE) III 0.038 (0.010 - 0.068) 0.017 UC to VGM L inft temp gyrus tempcoc 31 IVW(MRE) III 0.047 (0.023 - 0.072) - 0.0011 CD to VMM L entoRinal area 41 IVW(MRE) III 0.032 (-0.056 - 0.008) 0.020 CD to VGM L inft pocampus 41 IVW(MRE) III 0.033 (0.011 - 0.051) 0.006 UC to VGM R pippocampus 41 IVW(MRE) III 0.037 (0.014 - 0.053) 0.003 UC to VGM R parahipp gyrus post 31 IVW(MRE) III 0.037 (0.014 - 0.053) 0.003 UC to VGM R parahipp gyrus post 31 IVW(MRE) III 0.037 (0.014 - 0.059) 0.003 UC to VGM R caneal cortex 31 IVW(MRE) III 0.037 (0.014 - 0.059) 0.003 UC to VGM R caneal cortex 31 IVW(MRE) III 0.036 (0.009 - 0.063) 0.018 UC to VGM R caneal cortex 31 IVW(MRE) III 0.037 (0.014 - 0.059) 0.003 UC to VGM R caneal cortex 31 IVW(MRE) III 0.037 (0.026 - 0.048) 0.024 Bipolar Disorder UC to VGM R caneal cortex 31 IVW(MRE) III 0.037 (0.014 - 0.059) 0.003 UC to VGM R caneal cortex 31 IVW(MRE) III 0.037 (0.026 - 0.048) 0.024 VGM P value adjusted by fabe discovery rate (FCP) method: * in Method' column refer to existance of phetograms/ * in Method' column refer to existance of phetograms/ * in Method' column refer to existance of phetograms/ CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.023 (0.003 - 0.045) 0.034 42 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.023 (0.003 - 0.045) 0.034 42 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.023 (0.003 - 0.045) 0.034 42 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.025 (0.003 - 0.045) 0.034 42 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.025 (0.003 - 0.045) 0.034 42 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.035 (0.001 - 0.051) 0.006 52 CD to VGM L parahipp gyrus post 41 IVW(MRE) III 0.035 (0.001 - 0.051) 0.006 52 CD to VGM L parahipp gyrus post 41 IVW(MRE) III	UC to VGM L temp occ fusif cortex UC to VGM L inf temp gyrus post UC to VGM L inf temp gyrus tempoce UC to VGM R cerebellum X CD to VMM L parcentral area CD to VMM L entoRinal area CD to VGM L entoRinal area CD to VGM L entoRinal procempus UC to VGM R hippocampus		31 31 31	IVW(MRE) IVW(MRE)			0.036 (0.011 ~ 0.061)	0.008	96.5
UC to VGM R inf temp gyrus person     31     IVW(MRE)     IVW(MRE)     0.033 (0.000 - 0.067)     0.022       UC to VGM I. inf temp gyrus tempoco     31     IVW(MRE)     IVW(MRE)     0.047 (0.023 - 0.077)     0.0010     0.0020     0.0010     0.0010     0.0020     0.0010     0.0020     0.0010     0.0010     0.0020     0.0010     0.0020     0.0010     0.0020     0.0010     0.0020     0.0010     0.0020     0.0010     0.0020     0.0010     0.0050     0.0010     0.0050     0.0010     0.0020     0.0011     0.0010     0.0020     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011     0.0011 <td< td=""><td>UC to VGM R inf temp gyrus post UC to VGM L inf temp gyrus tempoco UC to VGM R cerebellum X trenia CD to VWM L paracentral area CD to VWM L entoRinal area CD to VGM R hippocampus UC to VGM R hippocampus</td><td></td><td>31 31</td><td>IVW(MRE)</td><td></td><td><b>—</b></td><td>0.036 (0.007 ~ 0.065)</td><td>0.030</td><td>96.8</td></td<>	UC to VGM R inf temp gyrus post UC to VGM L inf temp gyrus tempoco UC to VGM R cerebellum X trenia CD to VWM L paracentral area CD to VWM L entoRinal area CD to VGM R hippocampus UC to VGM R hippocampus		31 31	IVW(MRE)		<b>—</b>	0.036 (0.007 ~ 0.065)	0.030	96.8
UC to VGM L inf temp grous tempooc     31     IVW(MRE)     Image: Constraint of the second se	UC to VGM L inf temp gruss tempooc UC to VGM R cerebellum X C to VWM L paracentral area CC to VWM L entoRinal area CC to VWM L entoRinal area CC to VGM L hippocampus CC to VGM R hippocampus UC to VGM R hippocampus		31			<b></b>	0.038 (0.009 ~ 0.067)	0.022	98.0
Schizophrenia     UC to VGM R cerebellum X     31     IVW(MRE)     IVV     0.047 (0.023 - 0.072)     < 0.001       Schizophrenia     CD to VWM L paracentral area     41     MR Egger#	UC to VGM R cerebellum X CD to VVM L paracentral area CD to VVM L entoRinal area CD to VVM L entoRinal area CD to VGM L hippocampus CD to VGM R hippocampus UC to VGM R parateino gruis post		~ .	IVW(MRE)		<b>—</b>	0.039 (0.010 ~ 0.068)	0.017	98.2
Schizophrenia     CD to VVML L paracentral area     41     NR Egger <sup>4</sup> 0.063 (-0.1100.016) 0.019       CD to VGML LentoRinal area     41     IVW(IMRE)     -0.032 (-0.0560.008) 0.020       CD to VGM R hippocampus     41     IVW(IMRE)     -0.031 (0.011 - 0.51) 0.006       UC to VGM R hippocampus     41     IVW(IMRE)     -0.032 (-0.056 - 0.008) 0.021       UC to VGM R hippocampus     41     IVW(IMRE)     -0.031 (0.011 - 0.051) 0.006       UC to VGM R panipp gvrus post     31     IVW(IMRE)     -0.032 (0.009 - 0.063) 0.018       UC to VGM R frontal pole     31     IVW(IMRE)	hrenia CD to VWM L paracentral area CD to VWM L entoRinal area CD to VCM L hippocampus CD to VCM R hippocampus UC to VCM R parabing ours post		31	IVW(MRE)		<b>H</b>	0.047 (0.023 ~ 0.072)	< 0.001	99.9
CD to VWM LendRinal area     41     VVV(MRE)    0.032 (-0.0560.008) 0.020       CD to VGM L hippocampus     41     VVV(MRE)     0.025 (0.003 - 0.046)     0.045       CD to VGM R parahipp gyrus post     31     IVV(MRE)     0.031 (0.011 - 0.051)     0.006       UC to VGM R parahipp gyrus post     31     IVV(MRE)     0.035 (0.003 - 0.046)     0.045       UC to VGM R parahipp gyrus post     31     IVV(MRE)     0.037 (0.014 - 0.051)     0.006       UC to VGM R frontal pole     31     MR Eggert     0.032 (0.018 - 0.146)     0.024       Bipolar Disorder     UC to VGM R curael cortex     31     NVV(MRE)     0.044 (0.011 - 0.077)     0.017       Pain     CD to VMM R rotratanterioringulate area     1     IVV(MRE)     0.044 (0.011 - 0.077)     0.017       Pain     CD to VMM R rotratanterioringulate area     1     IVV(MRE)     0.026 (0.004 - 0.047)     0.037       View Prake adjutate dy friate discovery rote FR3P method;     edits	CD to VWM L entoRinal area CD to VGM L hippocampus CD to VGM R hippocampus LIC to VGM R parabino gvus post		41	MR Egger#	<b>—</b> —	1	-0.063 (-0.110 ~ -0.016	0.019	100.0
CD to VGM L hippocampus     41     IVW(MRE)     Image: Constraint of the state of the	CD to VGM L hippocampus CD to VGM R hippocampus UC to VGM R parahing arrus post		41	IVW(MRE)		-	-0.032 (-0.056 ~ -0.008	0.020	97.1
CD to VGM R hippocampus     41     VVV(MRE)     Image: Constraint of the co	CD to VGM R hippocampus		41	IVW(MRE)			0.025 (0.003 ~ 0.046)	0.045	91.1
UC to VGM R parahipp gyrus post     31     IVW(MRE)     Image: Constraint of the state	LIC to VGM R parabipp gyrus post		41	IVW(MRE)		<b>H</b>	$0.031(0.011 \sim 0.051)$	0.006	98.2
UC to VGM L parahipp gyrus post     31     IVW(MRE)     Image: Figure Fi			31	IVW(MRE)			0.036 (0.009 ~ 0.063)	0.018	96.1
UC to VGM R frontal pole     31     MR Egger#     0.082 (0.018 ~ 0.146)     0.028       UC to VGM R frontal pole     31     MR Egger#     0.082 (0.018 ~ 0.146)     0.028       Bipolar Disorder     UC to VGM R uncel cortex     31     MR Egger#     0.07 (0.026 ~ 0.188)     0.024       Pain     CD to VMM R rostralanterioroingulate area     41     IVW(MRE)     0.04 (0.011 ~ 0.077)     0.017       Note, P value adjusted by folles dicovery rate (PDR) immod?	UC to VGM L parahipp gyrus post		31	IVW(MRE)			$0.037(0.014 \sim 0.059)$	0.003	96.8
Bipolar Disorder     UC to VGM L particular opero cortex     31     MR Egget*     0.107 (0.026 - 0.188)     0.024       Pain     CD to VGM R cuneal cortex     31     IVW(MRE)     0.044 (0.011 - 0.077)     0.017       Note: Pvalue adjutted by false discovery rate (CR) method: ** n Method' column refer to existence of pleatory of SMR: ** n Method' column refer to existence of pleatory of SMR:	UC to VGM R frontal pole		31	MR Egger#			■ 0.082 (0.018 ~ 0.146)	0.028	100.0
Bipolar Disorder     UC to VGM R cuneal cortex     31     IVW(MRE)     Image: Disorder     0.044 (0.011 ~ 0.077)     0.017       Pain     CD to VVMR rostralanterioringulate area     41     IVW(MRE)     Image: Disorder     0.044 (0.011 ~ 0.077)     0.017       Note P-value adjusted by false downyr rate (FD) method:     Image: Disorder     0.044 (0.011 ~ 0.077)     0.017       Note P-value adjusted by false downyr rate (FD) method:     Image: Disorder     Image: Disorder     0.026 (0.004 ~ 0.047)     0.037       Mote P-value adjusted by false downyr rate (FD) method:     Image: Disorder     Image: Diso	UC to VGM L parietal operc cortex		31	MR Egger#			0 107 (0 026 ~ 0 188)	0.024	100.0
Pain     CD to VWM R rostralanteriorcingulate area     1     IVW(MRE)     0.026 (0.004 ~ 0.047)     0.037       Note: P value adjusted by false discovery rate (PDI) method: * In Method" column refer to existence of pleiotoxy of SNPs;     -dis     -dis<	Disorder UC to VGM B cuneal cortex		31	IVW(MRE)		i i i i i i i i i i i i i i i i i i i	$0.044(0.011 \sim 0.077)$	0.017	99.6
Note:     Palue adjusted by false discovery rate (PDF) method;	CD to VWM R rostralanteriorcingulate a	area	41	IV/W(MRE)		<u> </u>	$0.026(0.004 \sim 0.047)$	0.037	91.1
CD to VGM L parahipp gyrus post     1     IVV(IMRE)     IVV(IMRE)     0.023 (0.002 - 0.045)     0.034 (0.012 - 0.045)     0.045 (0.002 - 0.045)     0.045 (0.045 - 0.045)     0.045 (0.045 - 0.045)	CD to VWML entoRinal area	41	IVA		( المحمل		-0.032 (-0.056 ~ -0.008)	0.020	07 1
CD to VGM L parahipp grus post     41     IV/W(MRE)     H     0.023 (0.002 - 0.045)     0.034     6       CD to VGM L popcampus     41     IV/W(MRE)     H     0.025 (0.003 - 0.046)     0.045     6       CD to VGM L hippocampus     41     IV/W(MRE)     H     0.021 (0.011 - 0.051)     0.006     5	CD to VWM L entoRinal area	41	IV	W(MRE)			-0.032 (-0.056 ~ -0.008)	0.020	97.1
CD to VGM L hippocampus 41 IVW(MRE) → 0.025 (0.003 ~ 0.046) 0.045 S CD to VGM R hippocampus 41 IVW(MRE) → 0.031 (0.011 ~ 0.051) 0.006 S	CD to VGM L parahipp gyrus post	41	IVI	W(MRE)			0.023 (0.002 ~ 0.045)	0.034	82.6
CD to VGM R hippocampus 41 IVW(MRE) - 0.031 (0.011 ~ 0.051) 0.006 9	CD to VGM L hippocampus	41	IVI	W(MRE)			0.025 (0.003 ~ 0.046)	0.045	91.1
	CD to VGM R hippocampus	41	IVI	W(MRE)			0.031 (0.011 ~ 0.051)	0.006	98.2
UC to VGM R precun cortex 31 IVW(MRE) - 0.032 (0.011 ~ 0.053) 0.005	UC to VGM R precun cortex	31	IV	W(MRE)			0.032 (0.011 ~ 0.053)	0.005	95.1
UC to VGM R parahipp gyrus post 31 IVW(MRE) 0.036 (0.009 ~ 0.063) 0.018	UC to VGM R parahipp gyrus post	31	IV	W(MRE)			0.036 (0.009 ~ 0.063)	0.018	96.1
UC to VGM L parahipp gyrus post 31 IVW(MRE) → 0.037 (0.014 ~ 0.059) 0.003	UC to VGM L parahipp gyrus post	31	IV	W(MRE)			0.037 (0.014 ~ 0.059)	0.003	96.8
UC to VGM L precun cortex 31 IVW(MRE) → 0.037 (0.013 ~ 0.061) 0.005 9	UC to VGM L precun cortex	31	IVI	W(MRE)		<b></b> -	0.037 (0.013 ~ 0.061)	0.005	98.7
UC to VGM R inf temp gyrus post 31 IVW(MRE) 0.038 (0.009 ~ 0.067) 0.022 9	UC to VGM R inf temp gyrus post	31	IVI	W(MRE)	•		0.038 (0.009 ~ 0.067)	0.022	98.0
UC to VGM L inf temp gyrus tempocc 31 IVW(MRE) 0.039 (0.010 ~ 0.068) 0.017	UC to VGM L inf temp gyrus tempocc	31	IV	W(MRE)	1		0.039 (0.010 ~ 0.068)	0.017	98.2
UC to VGM R frontal pole 31 MR Egger# 0.082 (0.018 ~ 0.146) 0.028	UC to VGM R frontal pole	31	MF	R Egger#			0.082 (0.018 ~ 0.146)	0.028	100.0
Language UC to VGM R frontal pole 31 MR Egger* 0.082 (0.018 ~ 0.146) 0.028	Je UC to VGM R frontal pole	31	MF	R Egger#		<b>—</b>	0.082 (0.018 ~ 0.146)	0.028	100.0
UC to VGM R supramarg gyrus post 31 MR Egger <sup>#</sup> ► 0.129 (0.039 ~ 0.218) 0.012	UC to VGM R supramarg gyrus post	31	MF	R Egger#			→ 0.129 (0.039 ~ 0.218)	0.012	100.0
Emotion UC to VGM L pallidum 31 IVW(MRE) -0.042 (-0.070 ~ -0.015) 0.004 9	UC to VGM L pallidum	31	IVI	W(MRE)			-0.042 (-0.070 ~ -0.015)	0.004	98.9
Note: Prusiue adjusted by false discovery rate (FDR) methodoits -oits -oits -oits oit oits oit "In Method" column refer to existence of pleichtopy of SNPs; "In Method" column refer to existence of heteropenently between SNPs.	ue adjusted by false discovery rate (FDR) method; d" column refer to existence of pleiotropy of SNPs; d" column refer to existence of heterogeneity between SNPs.			-0.15	-0.1 -0.05 0	0.05 0.1 0.15	0.2		

## 3.3.2 IDPs classified by function

Eleven IDPs located in 5 brain structures (wedge-shaped lobe, inferior temporal lobe, olfactory-hippocampal cortex, subhippocampal cortex, frontal lobe) are influenced by the risk of IBD onset. The olfactory-hippocampal cortex is a recognized vital component involved in memory formation (26). The frontal lobe, inferior temporal lobe, and wedge-shaped lobe have also been shown to play a role in visual memory formation (27). CD altered 4 IDPs, such as Volume of entorhinal in the left hemisphere, and UC modified 7 IDPs, such as VGM in Right Precuneous Cortex. Two IDPs located in 2 brain structures (supramarg gyrus post, frontal lobe) are influenced by UC onset, and these areas are considered as functional regions for language (28), specifically, VGM in the Right supramarg gyrus post and the right frontal pole. The pallidum, as a component of the limbic system, is involved in the regulation of emotions. Our results revealed that UC altered the VGM in the Left Pallidum.

According to Cochran's Q test and MR-Egger interception test, some batches exhibited heterogeneity, with no evidence of potential heterogeneity or multiple effects biasing our findings (see Table 1). For the functional origins of brain volume, refer to Supplementary Table 8.

## 3.4 Two-step mediation analysis

Through Mendelian randomization mediation analysis, we identified 34 IDPs as mediators for CD or UC in relation to six

neuropsychiatric disorders. Among these mediators, the majority were complete mediating effects, with only MO Superior frontooccipital fasciculus R and T1 FAST ROIsR parahippocampal gyrus post exhibiting partial mediating effects for UC-mediated pain (see Table 2).

## 4 Discussion

Inflammatory bowel disease is a chronic, relapsing autoimmune disorder. Previous observational studies suggest a relationship between IBD and changes in brain imaging derived phenotypes. However, whether due to confounding factors and the unclear direction of causality, the existence of a causal relationship between them remains inconclusive (29–31). In this study, we employed Mendelian randomization to investigate bidirectional causal relationships between IBD and IDPs, exploring the potential links between IBD onset and changes in brain structure. This provides new insights for the diagnosis and treatment of IBD.

Corticotropin-Releasing Hormone-Binding Protein (CRH-BP), a secreted glycoprotein, binds to CRH with very high affinity, modulating the signal transduction of CRH receptors. It suppresses adrenocorticotropic Hormone (ACTH) release mediated by CRH-Receptor 1 (CRHR1) while potentially facilitating it in CRHR2 (32). Additionally, it inhibits CRHinduced anterior pituitary cell ACTH release *in vitro* (33). Research by Ketchesin KD et al. confirms the specific expression of CRH-BP in GABAergic neurons in the prefrontal cortex,

Batch	Exposures and Outcomes	NSNP	Q from IVW	Pval_Q from IVW	Q from MR-Egger	Pval_Q from MR-Egger	I2 for MR-Egger	Pval of pleotropy	Dirction from MR-Steiger	LOO	
66	UC to VGM L pallidum	31	22.071	20.336	0.851	0.882	0.985	0.198	TRUE	TRUE	
777	CD to VWM R inferiortemporal area	41	34.649	34.366	0.709	0.681	0.980	0.598	TRUE	TRUE	
115	CD to VGM L cerebellum VIIb	40	30.007	30.007	0.849	0.819	0.982	0.977	TRUE	TRUE	
145	CD to VGM R cerebellum X	41	38.627	38.600	0.532	0.488	0.977	0.869	TRUE	TRUE	
815	CD to VWM R rostralanteriorcingulate area	41	41.580	40.376	0.402	0.409	0.980	0.288	TRUE	TRUE	
141	CD to VGM L cerebellum X	41	42.373	42.368	0.369	0.328	0.979	0.944	TRUE	TRUE	
142	UC to VGM L cerebellum X	31	22.805	21.520	0.823	0.839	0.984	0.266	TRUE	TRUE	
10	UC to VGM L temp occ fusif cortex	31	31.680	31.427	0.383	0.346	0.988	0.633	TRUE	TRUE	
1994	UC to VGM R inf temp gyrus post	31	31.875	31.179	0.373	0.357	0.988	0.428	TRUE	TRUE	
2016	UC to VGM L inf temp gyrus tempocc	31	29.705	28.759	0.481	0.478	0.989	0.339	TRUE	TRUE	
146	UC to VGM R cerebellum X	31	21.093	20.767	0.885	0.867	0.988	0.573	TRUE	TRUE	
559	CD to VWM L paracentral area	41	41.390	34.856	0.410	0.659	0.987	0.015	TRUE	TRUE	
535	CD to VWM L entoRinal area	41	42.154	39.029	0.378	0.469	0.990	0.085	TRUE	TRUE	
71	CD to VGM L hippocampus	41	47.030	46.911	0.207	0.180	0.990	0.755	TRUE	TRUE	
73	CD to VGM R hippocampus	41	40.871	40.680	0.432	0.396	0.990	0.671	TRUE	TRUE	
2724	UC to VGM R parahipp gyrus post	31	25.244	23.767	0.713	0.740	0.987	0.234	TRUE	TRUE	
2722	UC to VGM L parahipp gyrus post	31	17.564	17.445	0.965	0.955	0.976	0.733	TRUE	TRUE	
496	UC to VGM R frontal pole	31	25.818	20.749	0.684	0.868	0.988	0.032	TRUE	TRUE	
28	UC to VGM L parietal operc cortex	31	34.214	29.482	0.272	0.440	0.990	0.039	TRUE	TRUE	
2710	UC to VGM R cuneal cortex	31	39.940	34.912	0.106	0.207	0.992	0.050	TRUE	TRUE	
2721	CD to VGM L parahipp gyrus post	41	36.594	35.536	0.624	0.629	0.984	0.310	TRUE	TRUE	
2704	UC to VGM R precun cortex	31	21.482	21.163	0.872	0.853	0.982	0.577	TRUE	TRUE	
2682	UC to VGM L precun cortex	31	27.799	24.819	0.581	0.688	0.989	0.095	TRUE	TRUE	
2216	UC to VGM R supramarg gyrus post	31	39.002	30.882	0.126	0.371	0.990	0.010	TRUE	TRUE	
Batches with f Multiplicative	3atches with fewer than 3 included SNPs are not eligible for heterogeneity and pleiotropy analysis. VGM, Gray Matter Volume; VWM, White Matter Volume; L, Left; R, Right; CD, Crohn's Disease; UC, Ulcerative Colitis; IVW, Inverse Variance Weighted; MRE, Vultiplicative Random Effects Model; OR, Odds Ratio; CI, Confidence Interval; NSNP, Number of SNPs included in the study; LOO, Leave-One-Out test.										

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#### TABLE 2 Results of two-step Mendelian randomization mediation analysis.

			X-Y		X-M		M-Y			Effect
Exposure	Mediator	Outcome	OR 95%CI	Ρ	OR 95%Cl	Р	OR 95%CI	Р	Туре	rate
CD	T1 FAST ROIs L ventral striatum	Alzheimer's disease	0.992 (0.949~1.037)	0.717	1.021 (1.003~1.040)	0.023	2.162 (1.390~3.365)	0.027	ТМ	100%
CD	T1 FAST ROIs L cerebellum VIIb	Alzheimer's disease	0.992 (0.949~1.037)	0.717	1.025 (1.005~1.044)	0.012	1.194 (1.021~1.396)	0.027	ТМ	100%
CD	L2 Sagittal stratum L	Alzheimer's disease	0.992 (0.949~1.037)	0.717	0.972 (0.951~0.992)	0.007	0.862 (0.750~0.992)	0.038	ТМ	100%
UC	T1 SIENAX peripheral grey normalised volume	Alzheimer's disease	0.974 (0.940~1.010)	0.160	1.142 (1.069~1.221)	0.001	1.591 (1.062~2.383)	0.024	ТМ	100%
UC	T1 FAST ROIs L temp occ fusif cortex	Alzheimer's disease	0.974 (0.940~1.010)	0.160	1.037 (1.007~1.067)	0.015	1.612 (1.523~1.707)	<0.001	ТМ	100%
UC	T1 SIENAX peripheral grey unnormalised volume	Alzheimer's disease	0.974 (0.940~1.010)	0.160	1.096 (1.047~1.147)	<0.001	1.969 (1.089~3.558)	0.025	ТМ	100%
UC	ProbtrackX FA ml r	Alzheimer's disease	0.974 (0.940~1.010)	0.160	1.050 (1.018~1.083)	0.002	1.244 (1.040~1.489)	0.017	ТМ	100%
UC	ProbtrackX OD slf l	Alzheimer's disease	0.974 (0.940~1.010)	0.160	0.966 (0.938~0.995)	0.020	0.844 (0.718~0.991)	0.039	ТМ	100%
CD	L G&S paracentral area	Anxiety disorders	0.999 (0.984~1.014)	0.889	0.940 (0.894~0.987)	0.018	0.811 (0.748~0.879)	<0.001	ТМ	100%
CD	MO Corticospinal tract R	Anxiety disorders	0.999 (0.984~1.014)	0.889	0.968 (0.943~0.993)	0.014	0.963 (0.953~0.972)	< 0.001	ТМ	100%
UC	T1 FAST ROIs L temp occ fusif cortex	Anxiety disorders	1.002 (0.982~1.022)	0.847	1.037 (1.007~1.067)	0.015	1.196 (1.062~1.346)	0.003	ТМ	100%
CD	FA Sagittal stratum L	Bipolar disorders	1.087 (0.999~1.183)	0.060	1.032 (1.010~1.055)	0.005	1.329 (1.278~1.382)	< 0.001	ТМ	100%
CD	L2 Sagittal stratum L	Bipolar disorders	1.087 (0.999~1.183)	0.060	0.972 (0.951~0.992)	0.007	0.799 (0.723~0.883)	<0.001	ТМ	100%
CD	ProbtrackX FA mcp	Bipolar disorders	1.087 (0.999~1.183)	0.060	1.029 (1.004~1.054)	0.023	1.237 (1.107~1.382)	< 0.001	ТМ	100%
CD	R S collat transv ant area	Pain	1.005 (0.995~1.015)	0.327	0.936 (0.888~0.986)	0.017	0.922 (0.913~0.931)	< 0.001	ТМ	100%
CD	ProbtrackX MO unc r	Pain	1.005 (0.995~1.015)	0.327	0.924 (0.871~0.980)	0.012	0.963 (0.929~0.997)	0.034	ТМ	100%
UC	MO Superior fronto- occipital fasciculus R	Pain	1.019 (1.008~1.030)	0.001	1.142 (1.035~1.260)	0.013	1.062 (1.025~1.101)	0.001	РМ	42.43%
UC	T1 FAST ROIsR parahipp gyrus post	Pain	1.019 (1.008~1.030)	0.001	1.037 (1.009~1.065)	0.009	1.066 (1.053~1.079)	<0.001	РМ	12.33%
CD	T1 FAST ROIs L cerebellum VIIb	depression	1.007 (0.987~1.028)	0.493	1.025 (1.005~1.044)	0.012	1.090 (1.004~1.183)	0.041	ТМ	100%
CD	volume Right- Lateral-Ventricle	depression	1.007 (0.987~1.028)	0.493	0.971 (0.951~0.992)	0.006	0.805 (0.719~0.902)	<0.001	ТМ	100%
CD	L paracentral area	depression	1.007 (0.987~1.028)	0.493	0.939 (0.897~0.983)	0.010	0.958 (0.927~0.990)	0.010	ТМ	100%
CD	L G&S paracentral area	depression	1.007 (0.987~1.028)	0.493	0.940 (0.894~0.987)	0.018	0.901 (0.820~0.989)	0.028	ТМ	100%
CD	R S collat transv ant area	depression	1.007 (0.987~1.028)	0.493	0.936 (0.888~0.986)	0.017	0.827 (0.714~0.959)	0.012	ТМ	100%

(Continued)

	Mediator	Outcome	X-Y		X-M		M-Y			F#cot
Exposure			OR 95%CI	Ρ	OR 95%Cl	Р	OR 95%CI	Р	Туре	rate
CD	ProbtrackX FA mcp	depression	1.007 (0.987~1.028)	0.493	1.029 (1.004~1.054)	0.023	1.160 (1.016~1.325)	0.029	ТМ	100%
CD	ProbtrackX MO unc r	depression	1.007 (0.987~1.028)	0.493	0.924 (0.871~0.980)	0.012	0.894 (0.847~0.944)	<0.001	ТМ	100%
UC	T1 FAST ROIs L temp occ fusif cortex	depression	1.011 (0.979~1.044)	0.495	1.037 (1.007~1.067)	0.015	1.181 (1.062~1.313)	0.002	ТМ	100%
UC	L S front inf thickness	depression	1.011 (0.979~1.044)	0.495	1.033 (1.011~1.056)	0.004	1.187 (1.001~1.408)	0.049	ТМ	100%
CD	T1 FAST ROIs L ventral striatum	Schizophrenia	1.036 (0.991~1.084)	0.118	1.021 (1.003~1.040)	0.023	1.373 (1.037~1.819)	0.027	ТМ	100%
CD	L G&S paracentral area	Schizophrenia	1.036 (0.991~1.084)	0.118	0.940 (0.894~0.987)	0.018	0.802 (0.762~0.844)	<0.001	ТМ	100%
CD	R inferiortemporal area	Schizophrenia	1.036 (0.991~1.084)	0.118	0.979 (0.962~0.996)	0.015	0.676 (0.625~0.731)	<0.001	ТМ	100%
UC	L superiorparietal thickness	Schizophrenia	1.050 (0.990~1.113)	0.105	1.041 (1.006~1.077)	0.020	1.053 (1.040~1.066)	< 0.001	ТМ	100%
UC	L G temporal inf thickness	Schizophrenia	1.050 (0.990~1.113)	0.105	1.047 (1.019~1.076)	0.001	1.771 (1.497~2.094)	<0.001	ТМ	100%
UC	L S precentral-sup- part thickness	Schizophrenia	1.050 (0.990~1.113)	0.105	1.046 (1.011~1.082)	0.010	1.481 (1.442~1.520)	<0.001	ТМ	100%
UC	OD Middle cerebellar peduncle	Schizophrenia	1.050 (0.990~1.113)	0.105	0.964 (0.936~0.992)	0.014	0.728 (0.541~0.979)	0.036	ТМ	100%

#### TABLE 2 Continued

Mediator Type includes TM, Total Mediation; PM, Partial Mediation; L, Left; R, Right; CD, Crohn's Disease; UC, Ulcerative Colitis; OR, Odds Ratio; CI, Confidence Interval.

particularly in interneurons expressing somatostatin (32). Our research findings suggest an increase in VGM in the frontal lobe is associated with a reduced risk of Crohn's disease. This could be attributed to an upregulation of CRH-BP, which suppresses the activity of the neuroendocrine axis (i.e., the hypothalamuspituitary-adrenal axis) and lowers the levels of CRH. Simultaneously it participates in the regulation of CRHR, lowering the occurrence of intestinal inflammation through multiple pathways. Elevated levels of CRH can damage the intestinal mucosal barrier, increase intestinal permeability, enhance autophagy in the gut, induce dysbiosis in the gut microbiota, and promote an M1/M2 polarization, consequently facilitating intestinal inflammation (34, 35). Furthermore, Li et al.'s study suggests that CRHR1 mediates intestinal injury through the mentioned pathway. Conversely, CRHR2 activates intestinal stem cells, promoting intestinal repair (36).

Additionally, our study suggests that an increase in gray matter volume in the prefrontal and extra-frontal lobes may increase the risk of UC and CD, although research on whether these structural changes trigger intestinal inflammation is yet to be discovered, and the underlying mechanisms require further exploration.

In a reverse causal relationship, we found that an increased risk of IBD onset leads to alterations in certain brain structures. This may be associated with the complications of neurocognitive disorders in IBD. In addition to common gastrointestinal clinical manifestations such as diarrhea and abdominal pain, IBD patients also present neuropsychiatric complications, with central nervous system involvement including cerebrovascular diseases (37) and psychiatric disorders such as cognitive impairments, anxiety, and depression (2, 38). Peripheral nervous system manifestations include peripheral neuropathy and demyelinating diseases (39). These clinical symptoms may be associated with the observed changes in IDP in this study. In this study, an increased risk of UC or CD was found to be associated with changes in the volume of the pallidum, anterior cingulate gyrus, frontal lobe, temporal lobe, inferior temporal gyrus, temporal-occipital fusiform gyrus, hippocampus, parahippocampal gyrus, and cerebellum, which is consistent with the observations made by Bao, Chun Hui et al. (40).

Human perception, memory, and emotional regulation are associated with the prefrontal-limbic system (41), which comprises the cingulate cortex, insular cortex, parahippocampal gyrus, hippocampus, prefrontal cortex, amygdala, temporal lobe, hypothalamus, and thalamus. These regions are interconnected by structures such as the fornix, corpus callosum, and superior longitudinal fasciculus, forming various functional circuits (42). Research has shown that IBD patients are more susceptible to developing mental and psychological disorders, including anxiety, depression, schizophrenia, bipolar affective disorder, and cognitive impairment (43, 44). The olfactory-hippocampal circuit is considered to be associated with human episodic memory (26), and structural defects in the olfactory-hippocampal cortex can lead to cognitive inflexibility and pattern separation impairments, consequently resulting in depression (45). Experiments conducted by He, Xiao-Fei et al. and others suggest that in DSS-induced colitis mice, microglial cell activation, an increase in A1-like astrocytes, impairment of the lymphatic clearance pathway, and the induction of hippocampal cortical neuron loss ultimately lead to cognitive dysfunction and neuronal damage (46). This aligns with the findings of our study, providing anatomical evidence to some extent for the association of IBD with the development of psychiatric disorders.

The frontal, temporal, and occipital lobes are central to processes related to visual and auditory perception, attention, and language (47–49). The encoding and perception of facial features occur in the occipitotemporal region, while the matching of newly encoded facial features with previously stored ones takes place in the temporal and frontal lobes (50, 51). Observational studies have suggested that in comparison to healthy individuals, patients with schizophrenia exhibit significant reductions in the volume of certain brain regions, including the temporal, frontal, occipital, and parietal lobes (52, 53). Our findings may offer an explanation for the increased susceptibility of IBD patients to develop disorders such as schizophrenia.

Chronic abdominal pain is a primary clinical manifestation of IBD, with pain signals transmitted through the spinal thalamocortical pathway, spinal amygdala pathway, and spinal thalamocortical pathway, among others, to the cortical edge region (54). The central processing of visceral sensory/pain signals occurs in the orbitofrontal cortex, insula, and anterior cingulate cortex (54), with a notable increase in glutamate concentration, especially enhanced activity in the anterior cingulate cortex. Glutamate is involved in the energy metabolism of astrocytes and serves as an excitatory neurotransmitter in signal transduction. However, excessive glutamate leads to excitotoxicity, causing damage to neurons and axons (55). During acute inflammation in IBD, there is an increased release of inflammatory cytokines (such as TNF-a) and an elevated bloodbrain barrier permeability (56). Circulating cytokines, due to the compromised blood-brain barrier, impact neurotransmitter metabolism and induce apoptosis in astrocytes and oligodendrocytes, disrupting the conduction of brain structures and neural fiber bundles (57). Therefore, proactive management of intestinal inflammation, reducing the release of inflammatory factors, and alleviating chronic pain may potentially decrease the risk of IBD-associated mental health disorders.

Concurrently referencing previous studies, we also selected 6 neuropsychiatric disorder datasets for an additional mediation analysis of IDPs. Although the Mendelian randomization mediation effects differed from the mediating IDPs found in observational studies, both demonstrated that CD or UC promotes the onset of neuropsychiatric symptoms by mediating different brain IDPs.

Our research has unveiled a causal relationship between IBD and brain volume, offering anatomical and endocrine-related mechanisms for IBD-induced neurological disorders such as anxiety, depression, and schizophrenia. Therefore, medical doctors may consider early disease management by actively controlling intestinal inflammation, reducing systemic inflammatory factor release, preventing central nervous system damage, and reducing the incidence of long-term mental health disorders. Furthermore, certain changes in the volume of specific brain structures may serve as potential biomarkers for predicting the course of IBD. However, further research is required to substantiate this.

Prior neuroimaging studies on the impact of IBD on brain structure and function have been observational in nature. While they can establish an association between the disease and brain structure, they cannot confirm a causal relationship between the two. Our study innovatively applied Mendelian randomization, utilizing a selection of genetically associated variables significantly related to exposure factors, thereby mitigating confounding factors. This approach unveiled a bidirectional causal relationship between IBD and IDPs, with rigorous FDR correction, power calculations, and multiple sensitivity analyses effectively mitigating statistical Type I and Type II errors, as well as errors by abnormal SNP data, ensuring robust and reliable results. Finally, this paper provides an explanation of potential mechanisms, emphasizes the significance of neuroimaging applications, and offers a prospect for potential therapeutic approaches in the future.

Our study does have limitations. The population in this study consisted exclusively of individuals from European backgrounds, necessitating further exploration for populations from other regions. The brain imaging data used in this study were derived from a crosssectional study, which precluded our investigation into the impact of the duration of inflammatory bowel disease on brain IDPs. Additionally, due to the utilization of aggregated data rather than individual data, stratified analysis of variables such as gender was not possible.

Given that the method of Wald ratio estimation assumes a linear causal relationship, this study cannot exclude the possibility of a nonlinear relationship between IDPs and the onset of IBD. Lastly, it is inevitable that individual channelization exist in MR studies.

In conclusion, our study has unveiled a causal relationship between intestinal inflammation and changes in brain morphology through Mendelian randomization analysis, thereby emphasizing the role of distinct IDPs in patients with IBD. We believe that persistent inflammatory stimulation and continuous input of pain signals can lead to neuronal and neural pathway damage, consequently resulting in structural changes in the brain. Similarly, changes in brain structure can trigger the exacerbation of intestinal inflammation through regulatory mechanisms involving the nervous system and endocrine system. IDPs may serve as biomarkers for screening, monitoring, and identifying complications in clinical practice for IBD. This holds the potential to predict the progression of IBD disease by conducting and revisiting head MRI examinations, assessing changes in brain structure and neural fiber conductivity, and taking early therapeutic measures. Furthermore, proactive IBD treatment and inflammation control can reduce the risk of developing mental and psychological disorders associated with IBD.

## 5 Conclusion

Through Mendelian randomization analysis, we have revealed a causal relationship between inflammatory bowel disease and 252

Brain Imaging Derived Phenotypes. This underscores the role of brain IDPs in IBD patients. Three IDPs have been identified to have a close association with the onset of IBD, and we have also discovered some IDPs alterations related to anxiety, depression, and other 8 other disorders and functions that are attributable to IBD onset. These findings hold the promise of becoming valuable biomarkers in clinical practice for IBD screening and complication monitoring.

## Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

## Author contributions

FL: Conceptualization, Data curation, Methodology, Writing – original draft. QZ: Conceptualization, Investigation, Writing – original draft, Writing – review & editing, Methodology. TT: Conceptualization, Writing – review & editing. YYL: Conceptualization, Funding acquisition, Writing – review & editing. ZDW: Visualization, Writing – review & editing. YQL: Investigation, Project administration, Writing – review & editing. ZW: Visualization, Writing – review & editing. XH: Project administration, Writing – review & editing. ZX: Project administration, Writing – review & editing. YC: Visualization, Writing – review & editing. YC: Visualization, Writing – review & editing.

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# Conflict of interest

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

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## Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fimmu.2024.1359540/full#supplementary-material

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