

## Article

# Brain morphometry and cognitive features in prediction of irritable bowel syndrome

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**Abstract:** *Background:* Irritable bowel syndrome (IBS) is a common condition within the spectrum of gut-brain disorders, characterized by abdominal pain, bloating, altered bowel habits, and different patterns of psychological distress. While brain-gut interactions are increasingly recognized in IBS pathophysiology, the relationship between brain morphometry, cognitive function, and clinical presentation remains poorly understood. *Objectives:* To investigate whether multivariate analysis of brain morphometric measures and cognitive test performance can distinguish patients with IBS from healthy controls (HCs), and to evaluate the relative importance of structural and cognitive features in this discrimination. *Methods:* In this cross-sectional study, 49 patients with IBS and 29 HCs underwent structural magnetic resonance imaging (MRI) brain examination and completed the Repeatable Battery for the Assessment of Neuropsychological Status (RBANS). Brain morphometry was analyzed using two versions of FreeSurfer software (v6.0.1 and v7.4.1). IBS severity was assessed using the IBS-Severity Scoring System (IBS-SSS). We employed both univariate and multivariate statistical and machine learning approaches, including cross-validation, to analyze morphometric and cognitive measures. *Results:* Univariate and multivariate analyses showed limited discrimination between IBS and HC groups using morphometric measures alone. However, when combining morphometric and cognitive measures in a machine learning framework, the model achieved 93% sensitivity in identifying IBS patients, albeit with 78% specificity. Feature importance analysis highlighted the significance of subcortical structures (particularly hippocampus, caudate, and putamen) and two cognitive domains (recall and verbal skills) in group discrimination. Software version comparison revealed substantial impact on morphometric measurements. *Conclusions:* Our findings suggest that the combination of brain morphometry and cognitive measures provides better discrimination between IBS and HC groups than either measure alone. The identified importance of subcortical structures and specific cognitive domains supports a complex brain-gut interaction in IBS. These results emphasize the need for multimodal approaches in IBS research and careful consideration of methodological factors in brain morphometry studies.

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**Keywords:** Irritable bowel syndrome; structural MRI; brain morphometry, cognition; supervised classification; machine learning

## Introduction

Irritable bowel syndrome (IBS) represents a prevalent and complex gastrointestinal (GI) disorder, affecting approximately 10% of the global population [1]. The syndrome is clinically defined by a characteristic symptom pattern: recurrent abdominal pain associated with defecation, accompanied by alterations in bowel habits [2], and can be divided into clinical phenotypes based on predominant bowel patterns [3] and overall symptom severity [4]. The clinical presentation is heterogeneous, with experiences ranging from mild discomfort to severe symptoms that substantially impair quality of life and daily functioning [4]. Notably, women are disproportionately affected, a difference that appears to arise from a complex interplay of biological factors (including hormonal influences), healthcare-seeking behaviors, and sociocultural determinants [5–8]. Such epidemiological patterns highlight the multifactorial nature of IBS and underscore the importance of considering both biological and psychosocial factors in its study and treatment.

A bidirectional relationship between gastrointestinal symptoms of IBS and psychological functioning is well-documented [9]. While gastrointestinal symptoms can trigger or exacerbate psychological distress, anxiety and depression may in turn amplify the intensity and frequency of abdominal pain [10]. Recent research has expanded this psychobiological framework to include cognitive function, revealing a more nuanced picture of brain-gut interactions in IBS. Although cognitive impairments have been demonstrated at the group level [11,12], these deficits seem to characterize specific subgroups rather than being a universal feature of IBS [9,13]. This heterogeneity in psychological and cognitive presentations aligns with contemporary models of the gut-brain axis [14,15], which conceptualize IBS as a disorder of disrupted neural-enteric communication. In these models, the brain serves as the central integration hub for processing and interpreting the complex array of visceral signals, emotional responses, and cognitive processes that may be involved in IBS.

The relationship between brain structure and cognitive function has evolved from simple localization models to more sophisticated network-based frameworks [16,17]. This network perspective gained particular relevance for understanding IBS through Mayer et al.'s [18] seminal paper in 2015, which proposed that alterations in brain networks could directly influence multiple cognitive domains in IBS patients. Recent empirical support for this systems-level approach comes from Li et al. [19], who identified several associations between symptom severity and regional brain volumes, including positive correlations with subcortical structures (globus pallidus, caudate, and putamen) and negative correlations with cortical regions (anterior cingulate, dorsolateral prefrontal cortex, anterior and mid-cingulate cortices) and subcortical areas (anterior insula, hippocampus, parahippocampal cortex, thalamus). Of special interest to the present study, they also showed that these brain regions were linked to cognitive performance on tests of language skills and memory function.

Studies of abdominal pain and visceral stimulation have consistently demonstrated involvement of distributed brain networks, encompassing both cortical and subcortical structures [20,21]. Building on this network perspective, Skrobisz et al. [22] conducted a comprehensive morphometric analysis in patients with non-specific digestive disorders, including IBS. Using FreeSurfer software (version 6.0.1), they analyzed 36 brain regions, including subcortical, cortical, and global measures derived from structural magnetic resonance imaging (MRI). Their univariate analyses revealed reduced thalamic volume in IBS patients compared to healthy controls, though volumes remained larger than in patients with inflammatory bowel diseases. While these findings suggest structural brain differences in IBS, univariate approaches may not capture the full complexity of brain-gut interactions. Therefore, our study builds upon Skrobisz et al.'s work in two key ways. First, we examine the robustness of their findings by comparing analyses using both FreeSurfer v6.0.1 and a more recent version, allowing us to differentiate between software-dependent and

true biological effects. Second, we extend beyond univariate analyses by implementing multivariate approaches, including supervised machine learning techniques, to capture complex patterns in brain morphometry that might better characterize IBS. This dual approach - methodological validation and advanced pattern analysis - aims to provide a more comprehensive understanding of the structural brain differences associated with IBS. Finally, responding to Skrobisz et al.'s [22] call for integrating clinical measures, we investigated whether combining cognitive performance data with morphometric features would enhance the accuracy of IBS versus HC classification.

We will address these aims in the following way:

- A We aim to replicate the morphometric differences between IBS patients and HC reported in [22] by using the same FreeSurfer software version (FS 6.0.1) and a similar univariate analysis approach as in the original study.
- B We will compare outcomes of the morphometric segmentation of the T1-w recordings in the current dataset ( $n = 78$ ) for the FreeSurfer software version used in [22] (FS 6.0.1) with a more recent version (FS 7.4.1) by calculating pairwise correlations.
- C We will investigate if morphometric features (derived from the FS 7.4.1 brain segmentation, both cross-sectional and longitudinal-stream-based analyses) can be used to distinguish between IBS individuals and HC. To this end, we will employ four different strategies: (i) we will conduct univariate comparisons between IBS individuals and HC; (ii) we will conduct multivariate comparisons between the two groups (by incorporating covariance structures of the morphometric features); (iii) we will apply a machine learning framework (ML) to predict group membership based on morphometric features; and (iv) we will identify feature importance of respective morphometric features should the classification into IBS individuals and HC be successful.
- D We will investigate if the prediction of group membership (IBS vs. HC) is improved when cognitive performance is added as a predictor to the morphometric features ML model. If prediction is improved, we will identify those features that have the strongest influence on distinguishing IBS individuals from HC.

## Materials and Methods

### Participants

This investigation is part of the Bergen Brain-Gut project, a comprehensive clinical study conducted at Haukeland University Hospital, Norway, between 2020 and 2022 (detailed protocol in Berentsen et al. [23]). Our sample comprised 49 patients with IBS and 29 healthy controls (HCs), all aged 18 years or older. Participant recruitment employed multiple strategies, including media advertisements, informational flyers, and direct referrals from the hospital's outpatient clinic. All potential participants underwent systematic screening by a trained nurse using standardized inclusion and exclusion criteria (Table 1), followed by a comprehensive assessment battery including gastrointestinal measures, psychometric testing, and multiparametric magnetic resonance imaging (MRI).

The final sample size was determined by several factors. While no formal a priori power analysis was conducted as effect sizes for brain morphometric differences in IBS were not well established at the study's inception, our sample size aligns with or exceeds those of similar neuroimaging studies in functional gastrointestinal disorders [22]. For the current analyses, we included only participants with complete data on all key measures and artifact-free MRI scans suitable for automated brain segmentation, ensuring data quality while maintaining the largest possible sample size. This approach balanced statistical power requirements with practical constraints and methodological rigor.

### Measures

Age and sex (not genetically verified) were self-reported by the participants at baseline.

Inclusion criteria	Exclusion criteria
Rome-IV criteria: Recurrent abdominal pain average at least 1 day/week during the last 3 months, and associated alterations in bowel habits at least 6 months before diagnosis. Other causes are excluded.  Normal diet at least 3 weeks before inclusion IBS score equal to or above 175	Pharmacological treatment affecting GI tract, including medication for anxiety and depression, diabetes, coeliac disease, IBS, Polycystic ovary syndrome, active Helicobacter pylori infection, Parkinson's disease, amyotrophic lateral sclerosis, or Psychiatric disorders.  Treated with antibiotics for the last 3 months Diets such as vegetarian or vegan Use of probiotics or low-FODMAP diet within the last 3 weeks Previous intestinal surgery, except appendectomy Metallic implants, claustrophobia, incompatible with MRI Travel outside Europe last 3 weeks Plan to travel in the near future Pregnancy

**Table 1.** Exclusion and inclusion criteria for the IBS patients. Source: Retrieved from [23]

#### The IBS-Severity Scoring system (IBS-SSS)

The IBS-Severity Scoring system is a questionnaire used to assess the severity and frequency of GI-related IBS symptoms [24]. The questionnaire includes five items related to (i) abdominal *pain intensity*, (ii) abdominal *pain frequency*, (iii) abdominal *distention/bloating*, (iv) dissatisfaction with *bowel habits*, and (v) interference with *quality of life* – over the past 10 days. The maximum score for each question is 100. A sum of scores < 75 is used to define "no or minimal problems", and the scores in the ranges [75, 175), [175, 300], and > 300 as "mild", "moderate", and "severe" IBS symptoms, respectively [24]. In the present study, an IBS-SSS score ≥ 175 was used as the inclusion criteria for the IBS group. Almost all HCs obtained an IBS-SSS score at the lowest level (< 75), with some reporting a score within the mild level ([75, 175)).

#### Repeatable Battery for the Assessment of Neuropsychological Status (RBANS)

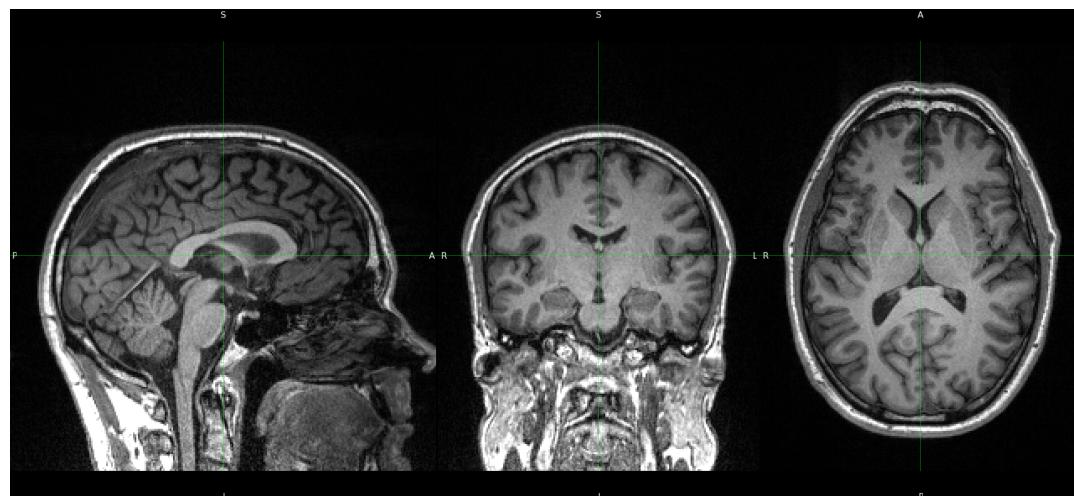
RBANS provides a quick and comprehensive assessment of five key cognitive domains, allowing the identification of specific areas of cognitive impairment, and takes about 30 minutes to administer. RBANS is sensitive to mild cognitive impairment, has good reliability and validity, can track changes over time, and is useful for both screening and detailed assessment. The five cognitive domains are (i) *immediate memory* (e.g., story memory and list learning tasks), (ii) *visuospatial/constructional skills* (e.g., copying geometric designs and identifying line orientation), (iii) *language* (e.g., picture naming and semantic fluency tasks), (iv) *attention* (e.g., digit span and coding tasks), and (v) *delayed memory* (e.g., recall of previously learned stories or lists). All participants performed the Norwegian A version of RBANS, administrated by a nurse trained by a clinical neuropsychologist, following the test manual's instructions [25]. The test battery comprises ten subtests, which are combined into five index scores and a total score. These scores are expressed both as raw and as age-corrected scaled scores. The scaled scores have a mean value of 100 and a standard deviation of 15 and are based on performance in a normative group matched to population statistics of 2012 in Norway, Sweden, and Denmark. We used these scaled scores on each of the five RBANS indices for a pairwise correlation analysis between brain morphometric measures and cognitive performance.

#### MRI Data Acquisition

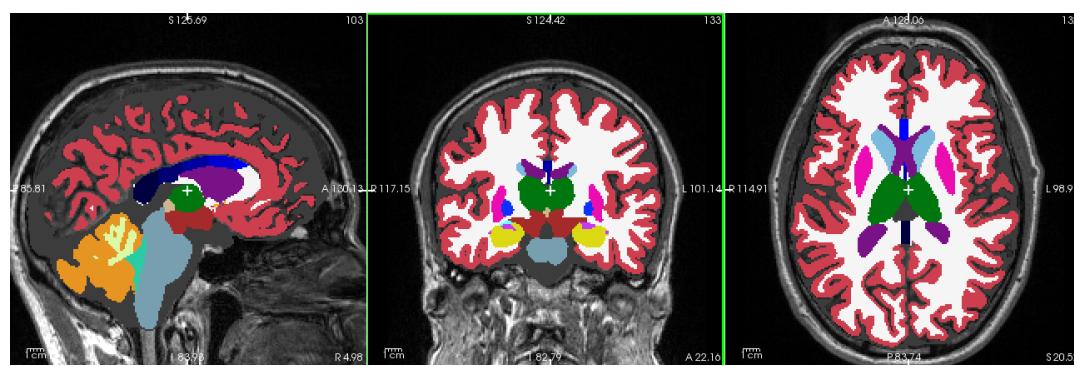
All neuroimaging data were acquired using a 3 Tesla Siemens Biograph mMR PET/MRI scanner equipped with a standard 12-channel head coil. The comprehensive multiparametric imaging protocol consisted of five sequences: a 3D T1-weighted MPRAGE (TA =

5:35), T2-weighted structural imaging (TA = 5:12), gradient echo (GRE) field mapping (TA = 0:54), resting-state functional MRI using echo-planar imaging (EPI) with integrated motion correction (TA = 9:48), and diffusion-weighted imaging with 30 gradient directions and three b-values (TA = 8:34). The total examination time was approximately 45 minutes. For the current morphometric analyses, we utilized only the high-resolution T1-weighted images, acquired using a 3D MPRAGE (Magnetization Prepared Rapid Gradient Echo) sequence. The acquisition parameters included a spatial resolution of 1.0 mm isotropic ( $1 \times 1 \times 1 \text{ mm}^3$ ) across 192 sagittal slices, with repetition time (TR) of 2500 ms, echo time (TE) of 2.26 ms, and inversion time (TI) of 900 ms. The field of view (FOV) was set to  $256 \times 256 \text{ mm}^2$  with a corresponding matrix size of  $256 \times 256$ , and parallel imaging was employed using GRAPPA with an acceleration factor of 2.

Figure 1 shows a representative T1-weighted image from our dataset, demonstrating the high tissue contrast necessary for accurate morphometric analysis. The corresponding FreeSurfer-generated segmentation mask, which forms the basis for our morphometric measurements, is illustrated in Figure 2. These images exemplify the quality standards maintained throughout our dataset.



**Figure 1.** 3D T1-weighted MPRAGE recording from BGA\_046. Panels left to right: Sagittal, Coronal, and Axial section, respectively. (01-freesurfer-freeview-t1-aseg-bga-046.ipynb)



**Figure 2.** The color-coded ASEG segmentation mask by FreeSurfer 7.4.1 overlaid on 3D T1-w MPRAGE from BGA\_046. Panels left to right: Sagittal, Coronal, Axial section, respectively. The white cross is located in the medial part of Left-Thalamus. Thalamus: green, Hippocampus: yellow, Caudate: light blue, Putamen: pink, Pallidum: purple, Cortex: red, White-Matter: white.

#### Brain Morphometry Analysis using FreeSurfer

Image processing and morphometric analyses were performed using FreeSurfer (<https://freesurfer.net>), a widely-validated open-source software suite for analyzing brain MRI

data [26]. To address both methodological and biological questions, we conducted parallel analyses using two FreeSurfer versions: version 6.0.1, which was employed in the reference study by Skrobisz et al. [22], and the current version 7.4.1.

The evolution of FreeSurfer's capabilities is particularly relevant to our investigation of brain structure in IBS. Version 7.0 (July 2020) introduced significant improvements in subcortical segmentation accuracy, while version 7.4.1 (June 2023) further enhanced the precision of limbic system structures, notably the hippocampus and amygdala. Additionally, version 7.4.1 provides superior compatibility with multi-modal imaging data and implements refined longitudinal processing algorithms. Since our multimodal MRI examinations were part of a longitudinal intervention study of IBS (Berentsen et al. [23]), we also used the longitudinal stream capability of FreeSurfer 7.4.1 to compare baseline longitudinal analysis with a cross-sectional analysis of the first MRI examination.

For both versions, we focused on the automated segmentation of subcortical structures using FreeSurfer's aseg pipeline, which identifies and quantifies the volume of distinct brain regions (detailed in Table A1). This dual-version approach serves two purposes: first, it enables direct comparison with Skrobisz et al.'s [22] findings, and second, it allows us to assess the impact of software evolution on morphometric measurements on a fixed dataset, and differences in cross-sectional and longitudinal stream analysis to discriminate HC and IBS from brain morphometric features. This methodological consideration is crucial, as previous studies have demonstrated that version-dependent variations in automated segmentation can significantly influence morphometric results [27–32].

The enhanced accuracy of version 7.4.1 is particularly relevant for our investigation of IBS, as it provides more reliable quantification of brain regions implicated in visceral sensation, pain processing, emotional regulation, and cognitive function. However, by analyzing our data with both versions, we can distinguish between genuine biological differences and methodologically-induced variations in brain morphometry.

#### *Statistical and Machine Learning Analysis*

All analyses were implemented in Python (version 3.10), with complete computational workflows and reproducibility materials available in our public GitHub repository (<https://arvidl.github.com/ibs-brain>). Our analytical approach combined traditional statistical methods with advanced machine learning techniques, employing both parametric and non-parametric approaches as appropriate for the data distributions.

For group comparisons, statistical significance was assessed using a threshold of  $p < 0.05$ , with Bonferroni correction applied to control for multiple comparisons. Effect sizes were quantified using Cliff's Delta, a robust non-parametric measure particularly suitable for non-normally distributed data. Following established conventions, we interpreted Cliff's Delta values as negligible (0.00-0.14), small (0.15-0.33), medium (0.34-0.47), or large (0.48-1.00).

Relationships between variables were evaluated using Spearman's rank correlation coefficient ( $\rho$ ), chosen for its robustness to non-normality and ability to capture monotonic relationships. Correlation strengths were classified as weak (0.20-0.39), moderate (0.40-0.59), strong (0.60-0.79), or very strong (0.80-1.00). Values below 0.20 were considered negligible to minimize the risk of over-interpreting weak associations.

To ensure reproducibility and transparency, all analysis scripts, including data preprocessing steps, statistical analyses, and visualization code, are documented in Jupyter notebooks accessible through our GitHub repository. These notebooks provide detailed documentation of parameter choices, statistical assumptions, and analytical decisions.

Our analysis strategy addressed four interconnected research objectives, progressing from replication to more advanced multivariate approaches:

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<b>A - Replication Analysis :</b>	239
Is it possible to replicate the morphometric findings in Skrobisz et al. [22] regarding IBS vs. HC, using the same FreSurfer-derived features and the same FreeSurfer version?	240
(i) By employing a feature-by-feature (univariate) comparison?	241
(ii) Employing a multivariate comparison, incorporating covariance structures in the morphometric features?	242
<b>B - Software Version Comparison :</b>	244
Are there IBS vs. HC disparities in morphometric feature values between FreeSurfer 6.0.1 and FreeSurfer 7.4.1 applied to the same set ( $n = 78$ ) of T1-weighted recordings in our Bergen cohort?	245
What about FreeSurfer 7.4.1 cross-sectional analysis versus FS 7.4.1 longitudinal stream?	246
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(iii) By predicting IBS vs. HC from the morphometric features using a machine learning framework (ML)?	257
(iv) identifying feature importance of the morphometric measures in the model with the best prediction?	259
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Would adding cognitive performance as a predictor improve the accuracy of separating IBS from HC?	263
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(iii) By predicting IBS vs. HC from the morphometric and cognitive features using a machine learning framework (ML)?	267
(iv) identifying feature importance of the morphometric and cognitive measures included in the model with the best prediction?	269
This hierarchical analytical framework progresses from basic replication to more advanced multivariate approaches, enabling both methodological validation and novel insights into IBS-related brain structure and function.	272
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Given the complexity of our research questions and the combination of traditional and advanced analytical methods, we implemented a comprehensive statistical framework encompassing both univariate and multivariate approaches. Here we detail our analytical strategy and its methodological justification.	277
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### Exploratory and Univariate Analyses

Initial analyses followed established protocols [22], beginning with exploratory data analysis of numerical features and cross-tabulation of categorical variables (Group: HC/IBS; Sex: F/M). For univariate comparisons (Objectives A-D), we employed both parametric (independent t-tests) and non-parametric (Mann-Whitney U) tests, depending on normality assessments. Multiple comparison correction used the Bonferroni method, and effect sizes were quantified using both Cohen's d (for parametric tests) and Cliff's delta (for non-parametric analyses).

### Permutation Testing

To address the challenges of small sample sizes and potential non-normal distributions, we implemented permutation testing with 1,000 iterations. This approach calculates an observed test statistic (sum of squared differences between group means) and generates a null distribution by randomly reassigning group labels and recalculating the statistic. The resulting empirical p-value represents the proportion of permuted statistics exceeding the observed value. This method offers several advantages: robustness to non-normality, natural handling of multiple comparisons, and suitability for small samples in multivariate analyses.

### Multivariate Approaches

For multivariate analyses (Objectives B-D), we first assessed multivariate normality using two complementary methods: Mardia's test (examining skewness and kurtosis) and the more comprehensive Henze-Zirkler's test, both implemented in SciPy's stats module. Effect sizes were quantified using Partial  $\eta^2$  (Eta-squared), interpreted following established guidelines [33]: small (0.01 – 0.06), medium (0.06 – 0.14), and large ( $\geq 0.14$ ) effects.

### Advanced Distance Metrics

To capture the complexity of high-dimensional relationships, we employed the Generalized Mahalanobis distance [34]. This sophisticated metric accounts for the covariance structure of the data while maintaining scale invariance across features. The method employs Moore-Penrose pseudoinversion to handle near-singular covariance matrices, enabling robust analysis even with highly correlated features. Significance can be assessed either parametrically through chi-square distribution or non-parametrically via permutation testing, providing flexibility in statistical inference. While this metric assumes similar covariance structures between groups, we supplemented it with permutation testing when this assumption was potentially violated. The combination of these approaches ensures robust statistical inference while acknowledging the inherent complexity of brain morphometry data.

### Prediction of class belonging using machine learning

In tasks C(iii) - D(iii) we applied a comprehensive machine learning framework, utilizing morphometric features derived from FreeSurfer (aseg) to develop predictive models for two distinct classification tasks. We employed PyCaret (<https://pycaret.org>), an open-source, low-code machine learning library in Python, to develop and evaluate our classification models.

### Machine Learning Model Development

Our machine learning approach followed a systematic protocol designed to ensure robust classification while addressing the challenges of limited sample size and potential overfitting. The analysis pipeline consisted of several carefully constructed stages optimized for neuroimaging data classification.

Initial data preparation employed a stratified sampling approach, partitioning the dataset into training (70%) and testing (30%) sets while preserving the distribution of key variables (IBS/HC status, sex, and cognitive function levels) across both partitions. This stratification

was crucial for maintaining representative samples and ensuring valid model evaluation, particularly given our modest sample size and the inherent complexity of neuroimaging data.

Model development utilized PyCaret's comprehensive machine learning framework to evaluate multiple classification algorithms, ranging from traditional approaches to advanced ensemble methods. The classifier suite included linear models (Logistic Regression with L1 and L2 regularization), non-linear algorithms (Support Vector Machines (SVM) with various kernels), tree-based methods (Random Forests, Gradient Boosting Machines including XGBoost and LightGBM), and instance-based learners (K-Nearest Neighbors). This diverse algorithm selection enabled exploration of different decision boundaries and feature interaction patterns.

To ensure robust model assessment and mitigate overfitting risks, we implemented a nested cross-validation strategy. The outer loop employed 10-fold cross-validation for model selection, while the inner loop optimized hyperparameters through random search with internal cross-validation. This approach provided unbiased performance estimates while preventing data leakage between model selection and evaluation phases. Hyperparameter optimization focused on algorithm-specific parameters crucial for neuroimaging data: regularization strengths for linear models, kernel parameters for SVMs, tree structure parameters for ensemble methods, and neighborhood configurations for instance-based learners. The final model selection prioritized both predictive performance and model interpretability, considering the clinical relevance of our findings.

### *Model Performance Assessment*

Model evaluation employed a multi-faceted approach to ensure comprehensive performance assessment, particularly important given our class imbalance between IBS and HC groups. While overall classification accuracy provided an initial performance indicator, we implemented additional metrics to capture nuanced aspects of model behavior and clinical relevance.

Our primary evaluation framework combined complementary performance measures. The F1 score, computed as the harmonic mean of precision and recall, offered a balanced assessment of model performance by considering both false positives and false negatives. We supplemented this with the Receiver Operating Characteristic Area Under Curve (ROC-AUC), which quantifies discrimination ability across different classification thresholds and is particularly robust to class imbalance. For detailed error analysis, we generated confusion matrices showing the distribution of correct and incorrect predictions across classes, enabling identification of specific classification patterns and potential biases. For the cognitive function analysis, which involved multiple classes, we employed macro-averaged versions of all metrics. This approach, calculating metrics independently for each class before averaging, ensures equal weighting of all classes regardless of their relative frequencies in the dataset. This consideration was particularly crucial given the uneven distribution of cognitive performance levels in our sample.

Performance assessment followed a dual-track strategy, evaluating models on both cross-validated training data and the held-out test set. This approach enabled us to assess both learning capacity and generalization ability, crucial considerations for clinical applications. To quantify uncertainty in our performance estimates, we calculated 95% confidence intervals for all metrics using bootstrapping methods. These intervals provide important context for interpreting model reliability and potential clinical utility, particularly given our moderate sample size.

The comprehensive evaluation strategy, combining multiple performance metrics with detailed error analysis and uncertainty quantification, ensures a thorough understanding of model behavior and reliability. This approach is particularly valuable for potential clinical applications, where understanding both model capabilities and limitations is crucial for

responsible implementation.

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### *Feature Importance and Model Interpretability Analysis*

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Understanding the relative contribution of different morphometric features to classification performance is crucial for both methodological validation and biological insight. We implemented two complementary approaches to feature importance analysis, combining model-agnostic methods with sophisticated game-theoretic interpretability techniques.

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The first approach employed permutation importance analysis, which quantifies feature relevance by measuring the degradation in model performance when individual features are randomly permuted. For each morphometric feature, we performed multiple permutation iterations, calculating the mean decrease in model performance. This approach provides an intuitive measure of feature importance while remaining independent of the underlying model architecture, enabling consistent interpretation across different classification algorithms.

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Our second approach utilized SHAP (SHapley Additive exPlanations) values, grounded in cooperative game theory [35]. This method offers a unified framework for model interpretation at both global and local levels. Global analysis aggregates SHAP values across all cases to identify consistently important features, while local analysis examines feature contributions to individual predictions. This dual-scale interpretation provides crucial insights into how different brain regions contribute to classification decisions, both at the population level and for individual cases.

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We applied both analytical approaches to our highest-performing models across all classification tasks. The results were synthesized through comprehensive visualization techniques, including SHAP summary plots depicting feature value distributions and their associated impact on model predictions. These visualizations integrate both the magnitude and directionality of feature effects, providing insight into how specific morphometric characteristics influence classification decisions.

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The integration of permutation importance and SHAP analysis provides complementary perspectives on feature relevance. While permutation importance identifies features whose disruption most impacts model performance, SHAP analysis reveals the complex interactions between features and their contributions to specific predictions. This comprehensive approach to model interpretation enhances our understanding of the neurobiological features distinguishing IBS patients from healthy controls, while also illuminating potential relationships between brain structure, sex/gender differences, and cognitive function. The results not only validate our modeling approach but also suggest specific brain regions and networks that may play key roles in IBS pathophysiology.

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## Results

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### *Sample Demographics and Clinical Characteristics*

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Our study included 78 participants (49 IBS patients and 29 healthy controls), with demographic and clinical characteristics summarized in Table 2. The age distributions were comparable between groups, with median ages of 34 years in the IBS group and 33 in the healthy control group. Both groups showed a predominance of female participants, with slightly higher representation in the IBS group (38 of 49) compared to the control group (20 of 29).

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IBS symptom severity, assessed using the IBS-SSS, demonstrated clear group separation. The IBS group reported substantially higher symptom severity compared to healthy controls, consistent with our inclusion criteria. Missing IBS-SSS data for three participants in each group (total n = 6) were handled through multiple imputation. The observed severity scores in the IBS group indicate predominantly moderate to severe symptomatology, while

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healthy controls showed minimal gastrointestinal symptoms, as expected.

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**Table 2.** Demographic and Clinical Characteristics of the Study Sample

Group	Age Median (IQR)	IBS_SSS Median (IQR)	Sex F/M (%)	N	Missing IBS_SSS
HC (N=29)	33.0 (23.0)	21.0 (30.0)	69.0/31.0	29	3
IBS (N=49)	34.0 (14.0)	264.0 (95.0)	77.6/22.4	49	3

Age is reported in years; IBS-SSS scores range from 0-500, with higher scores indicating greater symptom severity. IQR = Interquartile Range; F/M = Female/Male ratio expressed as percentages.

Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/02-demographics-and-clinical-characteristics.ipynb>

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Replication analysis of Skrobisz (2022) using the Bergen cohort (with FS 6.0.1)

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Table 3 addresses the attempt to replicate, in our Bergen cohort, the morphometric findings in Skrobisz et al. [22] regarding IBS vs. HC, using the same metrics (i.e., 35 eTIV-normalized volumes) from FreeSurfer-derived brain regions and the same FreeSurfer version (FS 6.0) as reported in Skrobisz et al. (2022).

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**Table 3.** Comparison of eTIV-normalized regional brain volumes between the two cohorts.

Brain Region	Skrobisz Cohort (FS 6.0)				Bergen Cohort (FS 6.0.1)			
	HC (N=19)		IBS (N=20)		HC (N=29)		IBS (N=49)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Left-Cerebellum-WM	0.00992	0.00113	0.00971	0.00107	0.01050	0.00092	0.01048	0.00092
Left-Cerebellum-Cortex	0.03628	0.00302	0.03553	0.00256	0.03894	0.00344	0.03931	0.00373
Left-Thalamus	0.00511	0.00037	0.00500	0.00024	0.00523	0.00046	0.00514	0.00039
Left-Caudate	0.00239	0.00025	0.00228	0.00021	0.00236	0.00026	0.00236	0.00031
Left-Putamen	0.00336	0.00033	0.00324	0.00028	0.00348	0.00038	0.00344	0.00039
Left-Pallidum	0.00140	0.00012	0.00135	0.00010	0.00140	0.00015	0.00137	0.00011
Left-Hippocampus	0.00270	0.00021	0.00272	0.00020	0.00291	0.00027	0.00290	0.00024
Left-Amygdala	0.00118	0.00013	0.00113	0.00015	0.00122	0.00010	0.00120	0.00010
Left-Accumbens-area	0.00031	0.00005	0.00034	0.00006	0.00043	0.00007	0.00042	0.00006
CSF	0.00061	0.00009	0.00060	0.00012	0.00067	0.00012	0.00070	0.00014
Right-Cerebellum-WM	0.00908	0.00106	0.00922	0.00100	0.00997	0.00089	0.00998	0.00085
Right-Cerebellum-Cortex	0.03652	0.00321	0.03616	0.00264	0.03972	0.00344	0.03998	0.00376
Right-Thalamus	0.00488	0.00030	0.00475	0.00024	0.00512	0.00044	0.00507	0.00036
Right-Caudate	0.00244	0.00024	0.00236	0.00024	0.00244	0.00024	0.00244	0.00030
Right-Putamen	0.00336	0.00030	0.00330	0.00028	0.00351	0.00037	0.00349	0.00035
Right-Pallidum	0.00136	0.00012	0.00133	0.00010	0.00132	0.00013	0.00130	0.00011
Right-Hippocampus	0.00282	0.00022	0.00285	0.00021	0.00301	0.00024	0.00298	0.00023
Right-Amygdala	0.00125	0.00012	0.00120	0.00012	0.00128	0.00009	0.00127	0.00010
Right-Accumbens-area	0.00034	0.00004	0.00036	0.00005	0.00043	0.00005	0.00043	0.00006
WM-hypointensities	0.00047	0.00015	0.00048	0.00013	0.00079	0.00031	0.00069	0.00025
CC_Posterior	0.00065	0.00013	0.00065	0.00010	0.00065	0.00010	0.00070	0.00011
CC_Mid_Posterior	0.00038	0.00007	0.00036	0.00007	0.00037	0.00007	0.00040	0.00007
CC_Central	0.00039	0.00009	0.00043	0.00008	0.00039	0.00009	0.00039	0.00010
CC_Mid_Anterior	0.00041	0.00009	0.00044	0.00013	0.00038	0.00008	0.00041	0.00011
CC_Anterior	0.00062	0.00010	0.00061	0.00008	0.00062	0.00010	0.00065	0.00010
BrainSegVol	0.75340	0.01784	0.74913	0.01647	0.80464	0.02487	0.80558	0.02397
BrainSegVolNotVent	0.74137	0.01880	0.73857	0.01836	0.79224	0.02511	0.79132	0.02490
lhCortexVol	0.15339	0.00620	0.15313	0.00871	0.16670	0.00800	0.16693	0.00951
rhCortexVol	0.15490	0.00690	0.15467	0.00859	0.16614	0.00828	0.16646	0.00939
CortexVol	0.30829	0.01298	0.30780	0.01715	0.33283	0.01611	0.33339	0.01880
lhCerebralWhiteMatterVol	0.15101	0.00748	0.15058	0.00742	0.15990	0.00858	0.15915	0.00876
rhCerebralWhiteMatterVol	0.15103	0.00757	0.15075	0.00727	0.15925	0.00829	0.15827	0.00938
CerebralWhiteMatterVol	0.30205	0.01500	0.30133	0.01461	0.31915	0.01678	0.31742	0.01808
SubCortGrayVol	0.03930	0.00194	0.03855	0.00162	0.04092	0.00258	0.04063	0.00236
TotalGrayVol	0.42105	0.01376	0.41884	0.01868	0.45307	0.02208	0.45396	0.02432

Note: All volumes are normalized to estimated total intracranial volume (eTIV)

HC = Healthy Controls; IBS = Irritable Bowel Syndrome; SD = Standard Deviation; WM = White Matter

Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/03-replication-analysis-fs6.ipynb>

The volumetric analysis comparing brain structures between IBS patients and healthy controls across the two cohorts reveals several noteworthy patterns. The Bergen cohort shows consistently larger volumes (6–8% for global measures, up to 35% for specific structures like the *accumbens*). However, global brain eTIV-normalized volumes show remarkable consistency between IBS and HC groups within each cohort, with BrainSegVol values being nearly identical (Skrobisz: HC  $0.753 \pm 0.018$ , IBS  $0.749 \pm 0.016$ ; Bergen: HC  $0.805 \pm 0.025$ , IBS  $0.806 \pm 0.024$ ). Cortical volumes demonstrate similar stability, with total cortical volume (CortexVol) showing minimal between-group differences in both cohorts. Subcortical structures exhibit subtle variations, with a minor trend toward volume reduction in IBS patients' subcortical gray matter (SubCortGrayVol), though these differences remain within standard deviation ranges. The analysis of white matter reveals consistent volumes between groups within cohorts, while white matter hypointensities show intriguing variation in the Bergen cohort. Corpus callosum segments maintain relatively uniform volumes across all groups. Methodologically, it is important to note the different cohort sizes (Skrobisz: HC=19, IBS=20; Bergen: HC=29, IBS=49) and the possible subtle differences in FreeSurfer versions (6.0 vs 6.0.1) and operating systems, which might contribute to the systematic differences observed between cohorts. The normalization to estimated total intracranial volume (eTIV) enables direct comparisons while controlling for head size variation within cohorts but not between cohorts.

A more penetrating reproducibility analysis is illustrated in Figure 3 showing the differences in HC versus IBS of eTIV-normalized brain regions volumes between the Skrobisz (2022) and the Bergen cohorts. The plot displays effect sizes from the Skrobisz cohort on the x-axis against the Bergen cohort on the y-axis, with a diagonal reference line indicating perfect agreement. We are here using Cohen's d values for region-wise effect size which are a standardized measure of the difference between means. More specifically, for each eTIV-normalized brain region volume and cohort we are calculating the pooled standard deviation as:

$$s_p = \sqrt{\frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2}{n_1 + n_2 - 2}}$$

where  $n_1$  and  $n_2$  are the sample sizes, and  $s_1$  and  $s_2$  are the standard deviations of the two groups, IBS and HC, respectively. Cohen's d effect size was then computed as:

$$d = \frac{\bar{x}_1 - \bar{x}_2}{s_p}$$

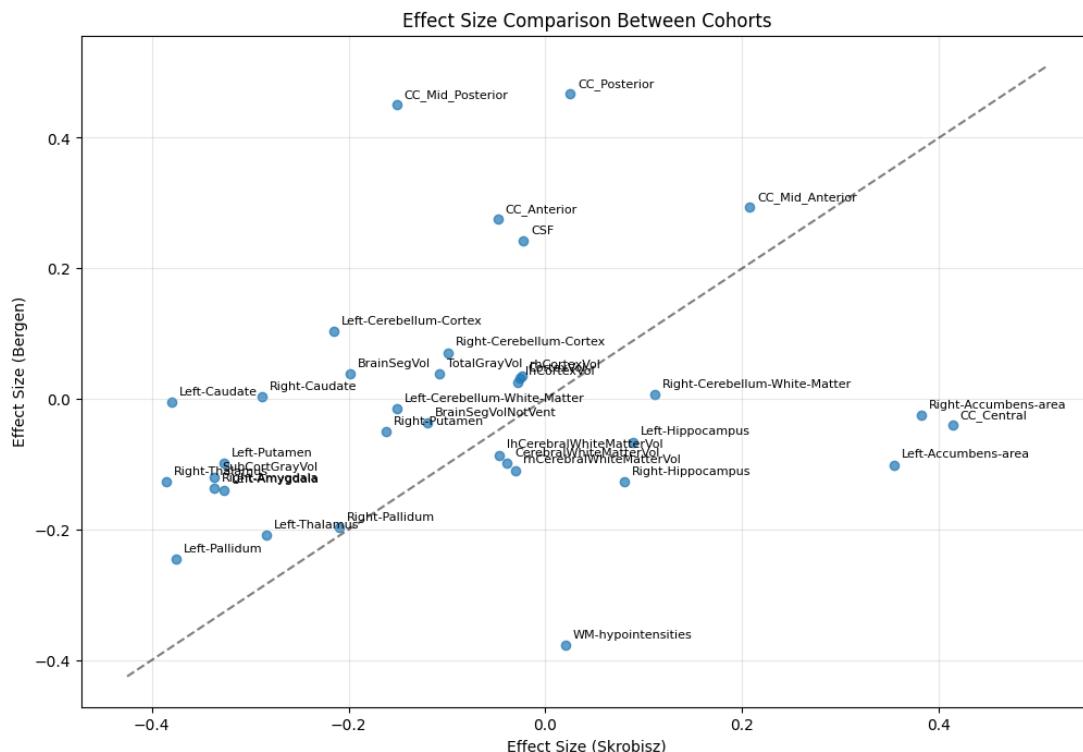
where  $\bar{x}_1$  and  $\bar{x}_2$  are the means of the two groups. The 95% confidence interval for d was calculated using:

$$CI_{95\%} = d \pm 1.96 \sqrt{\frac{n_1 + n_2}{n_1 n_2} + \frac{d^2}{2(n_1 + n_2)}}$$

where the standard error term accounts for both sampling variance and uncertainty in the effect size estimate.

An overall score ( $S$ ) for each brain region was computed as a composite measure combining three key components: binary sign consistency ( $\sigma$ ), confidence interval overlap ( $\omega$ ), and effect magnitude ( $\epsilon$ ). Specifically,  $S = \sigma + \omega + \epsilon$ , where  $\sigma$  equals 1 if the direction of effect is consistent between cohorts and 0 otherwise,  $\omega$  equals 1 if the 95% confidence intervals overlap and 0 otherwise, and  $\epsilon$  represents the minimum absolute effect size observed across cohorts. This scoring system yields higher values for brain regions demonstrating consistent effects across cohorts, with the magnitude component ( $\epsilon$ ) providing additional weight to regions showing stronger effects. A higher overall score thus indicates greater reliability and robustness of the observed effects across different study populations

and analysis pipelines, offering a quantitative basis for identifying the most consistently replicated findings in brain morphometry.



**Figure 3.** The large disparity of region-wise effect size of IBS vs. HC in comparison between the Skrobisz (2022) cohort and the Bergen cohort. Scatterplot of calculated Cohen's d effect sizes for each region in both cohorts (see text for details).

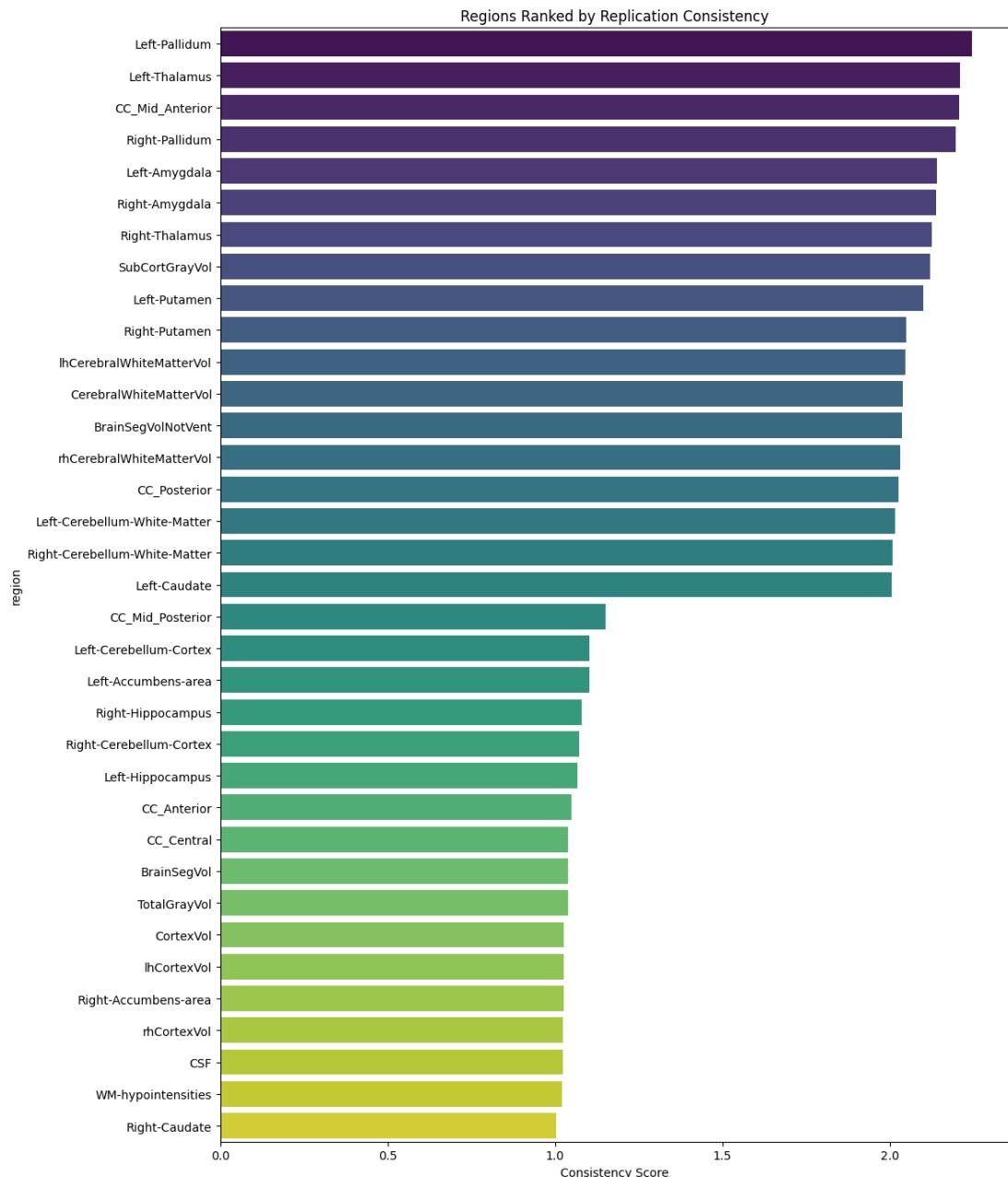
Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/03-replication-analysis-fs6.ipynb>

The scatter plot, presenting a comparison of effect sizes between the two cohorts, reveals a moderate effect size correlation of 0.203 ( $p=0.243$ ) between the cohorts. We found a sign consistency of 51.4%, meaning that in approximately half of the brain regions (51.4%), the direction of the difference between IBS and HC groups was the same in both cohorts. We also found a complete (100%) CI overlap rate, i.e., for all brain regions the 95% confidence intervals of the effect sizes from the two cohorts overlapped (data not shown). The relatively low sign consistency suggests poor replication of the directional effects between cohorts. The complete overlap suggests that while the point estimates of effects might differ between cohorts, we cannot conclude that these differences are statistically significant, given the uncertainty in our measurements.

Several brain regions show notable consistency between cohorts, particularly the CC\_Mid\_Anterior, Left-Pallidum, Left-Thalamus, Right-Pallidum, and Left-Amygdala, which are identified as the most consistent regions with overall scores ( $S$ ) ranging from 2.14 to 2.26. Some regions demonstrate substantial divergence from the diagonal, suggesting differential effects between cohorts. For instance, WM-hypointensities show particularly discordant effects between cohorts, as do several corpus callosum regions (CC\_Posterior and CC\_Mid\_Posterior) which exhibit stronger effects in the Bergen cohort. The cerebellar regions cluster near the center, indicating relatively modest effects in both cohorts. To conclude our analysis, the pattern of data points suggests modest overall agreement between cohorts, with considerable variability in effect sizes across brain regions. The widespread distribution of points around the diagonal line, combined with the moderate correlation coefficient, indicates that while there is some consistency in findings between cohorts, there are also substantial differences in how IBS-related brain alterations manifest in these two

independent samples.

Figure 4 plots a ranking of brain regions on how consistently they show similar patterns between the cohorts.



**Figure 4.** Barplot of regions ranked by their replication consistency . The consistency score used for ranking combines several factors: sign match, which assesses whether the direction of difference is the same in both cohorts; the CI overlap, which checks whether the confidence intervals for the effects overlap between cohorts; and effect magnitude, which evaluates how similar the size of the effect is between cohorts (see text for details).

Generated by: <https://github.com/arvid1/ibs-brain/blob/main/notebooks/03-replication-anaylsis-fs6.ipynb>

The replication consistency scores and their rankings reveal that several brain regions exhibit high levels of replication consistency across cohorts. The top-ranked regions, such as the Left-Pallidum, Left-Thalamus, and CC\_Mid\_Anterior, all have an overall score above 2.0, indicating strong consistency. These regions show both a consistent

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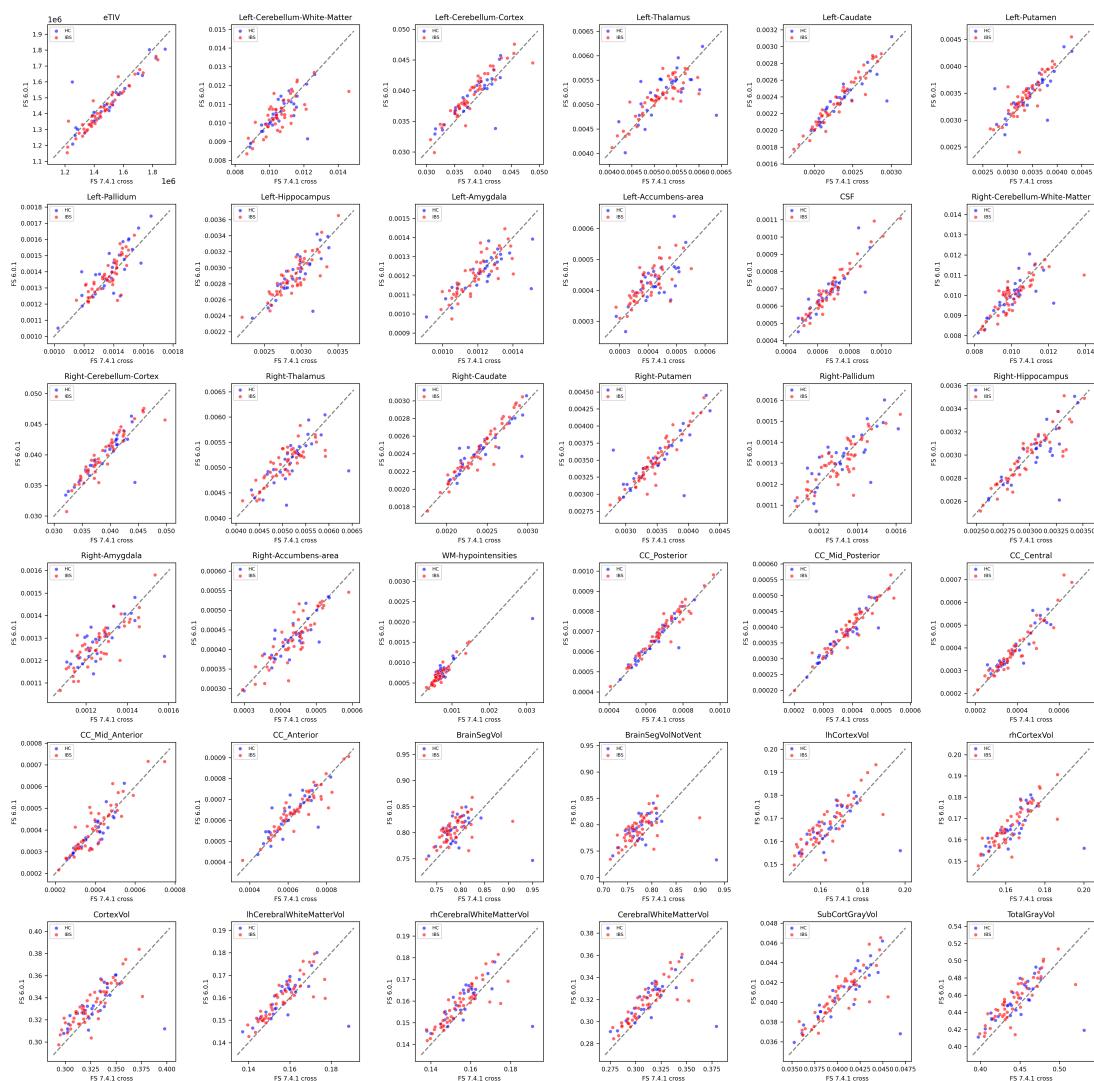
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direction of effect (sign match) and complete confidence interval overlap, along with relatively high effect magnitudes. This suggests that these regions are reliably replicated across studies, making them robust findings. In contrast, regions like the Right-Caudate, Right-Cerebellum-Cortex, Left- and Right-Hippocampus, CC\_Mid\_Posterior and Left-Cerebellum-Cortex, which have overall scores around 1.1, lack sign consistency despite having complete confidence interval overlap. This indicates that while the magnitude of effects is similar, the direction of effects is not consistent across cohorts, suggesting variability in these findings. Overall, the analysis highlights that while some regions demonstrate strong replication consistency, others show variability, particularly in the direction of effects, which may reflect differences in cohort characteristics or measurement variability. To conclude this consistency analysis, the rankings suggest that while eTIV-normalized volumes differ systematically between cohorts, some regions show more consistent relative patterns than others. This information is valuable for understanding which brain measurements might be more reliable across different studies and analysis pipelines. An interim conclusion is that the attempt to replicate the specific HC vs IBS morphometric differences reported in Skrobisz et al. (2022) was largely unsuccessful, suggesting that reported brain structural differences in IBS might be less robust or more heterogeneous than previously thought.

To evaluate the reproducibility and reliability of brain morphometry measurements in IBS research, we conducted a comparative analysis using multiple FreeSurfer processing pipelines on neuroimaging data from the Bergen cohort. Specifically, we examined how morphometric measurements and IBS versus healthy control (HC) group differences were affected by different FreeSurfer versions (6.0.1 versus 7.4.1) and different processing streams within FreeSurfer 7.4.1 (cross-sectional versus longitudinal). The inclusion of longitudinal stream analysis was enabled by our interventional study design, allowing us to assess the consistency of morphometric findings across different analytical approaches. Moreover, in these comparisons, we had access to all measurements from all subjects and not only summary statistics as in the replication analysis of the Skrobisz (2022) cohort.

#### *Univariate comparisons of morphometric measures using different FreeSurfer versions*

In this analysis, we aimed to compare brain region volumes between two FreeSurfer versions, FS 6.0.1 and FS 7.4.1 cross-sectional, to assess their consistency in measuring volumetric differences between IBS patients and healthy controls (HC). The scatter plot matrix in Figure 5 visualizes the relationship between corresponding brain regions across the two versions. Each subplot represents a specific brain region, with volumes from FS 6.0.1 plotted on the y-axis and those from FS 7.4.1 on the x-axis. Data points are color-coded by group, with blue representing HC and red representing IBS. An identity line is included in each plot to facilitate visual assessment of agreement between the two versions.



**Figure 5.** Scatter plot matrix comparing brain region volumes between FreeSurfer versions, FS 6.0.1 and FS 7.4.1 cross-sectional. Blue dots represent HC, and red dots represent IBS (see text for details).

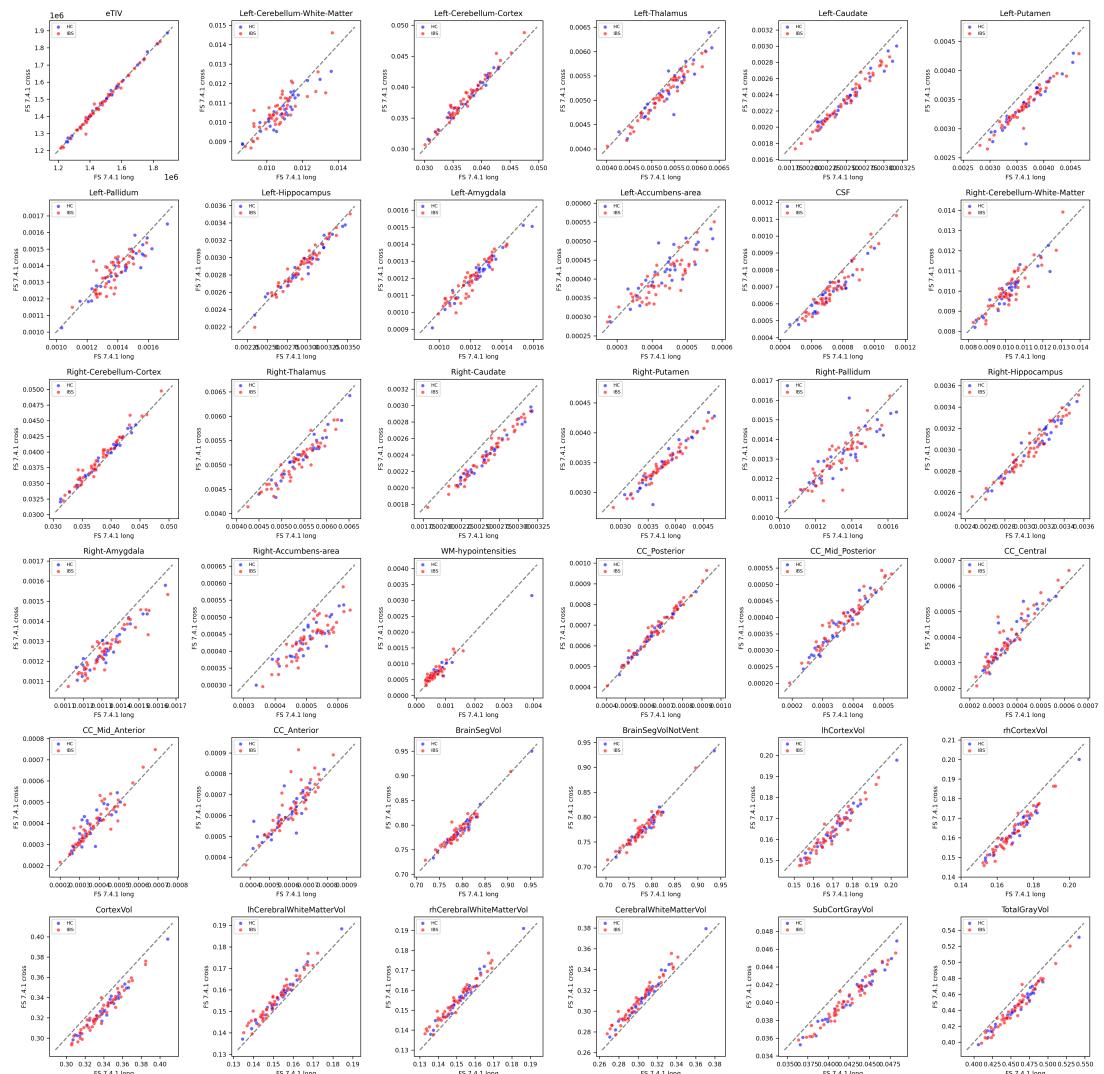
Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/04-comparing-FS-versions-on-same-dataset.ipynb>

The scatter plot matrix reveals a generally strong agreement between FS 6.0.1 and FS 7.4.1 cross-sectional measurements, as indicated by the clustering of data points around the identity line in most brain regions. This suggests that both versions provide consistent volumetric estimates. However, some variability is observed, particularly in regions such as the corpus callosum and subcortical structures, where deviations from the identity line are more pronounced. These discrepancies may reflect differences in segmentation algorithms or version-specific sensitivity to regional signal intensity variations. Overall, the figure supports the reliability of both FreeSurfer versions in capturing brain volume differences, though minor inconsistencies warrant further investigation regarding morphometry-based group prediction.

The summary statistics by the mean and standard deviation in HC and IBS patients on each of the 35 included brain regions (also reported by Skrobisz et al. [22]) derived from the `aseg.stats` files using cross-sectional Freesurfer 6.0.1 and Freesurfer 7.4.1, respectively, is shown in the Appendix as Table A1.

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Figure 6 depicts a scatter plot matrix comparing brain region volumes between two FreeSurfer pipelines: FS 7.4.1 cross-sectional and FS 7.4.1 longitudinal. Each subplot represents a specific brain region, with FS 7.4.1 cross-sectional volumes on the y-axis and FS 7.4.1 longitudinal on the x-axis. Data points are color-coded by group, with blue for HC and red for IBS. An identity line is included to facilitate visual assessment of agreement between the two processing streams. This visualization allows for a comprehensive comparison of how each stream measures brain volumes, highlighting potential discrepancies.



**Figure 6.** Scatter plot matrix comparing brain region volumes between FS 7.4.1 cross-sectional and FS 7.4.1 longitudinal stream. Blue dots represent HC, and red dots represent IBS (see text for details).

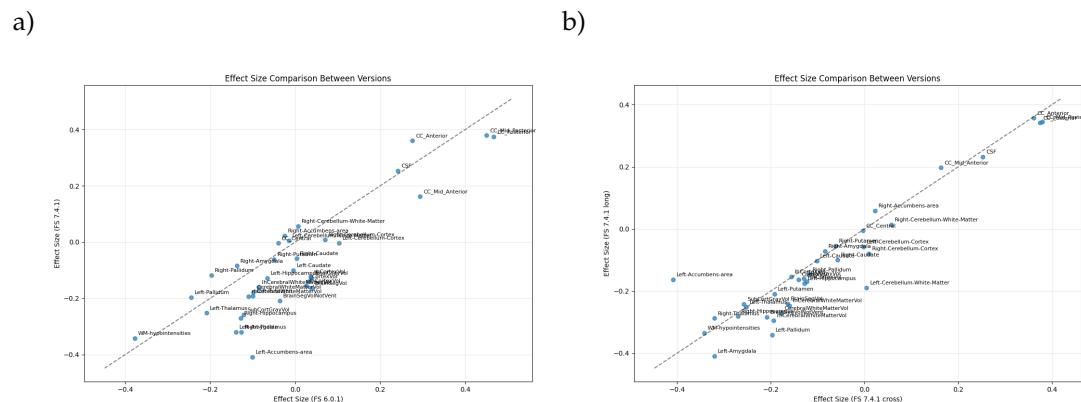
Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/04-comparing-FS-versions-on-same-dataset.ipynb>

The scatter plot matrix reveals a strong overall agreement between FS 7.4.1 cross-sectional and FS 7.4.1 longitudinal measurements, as indicated by the clustering of data points around the identity line in most brain regions. This suggests that both processing streams provide consistent volumetric estimates. However, some variability is observed, particularly in regions such as the corpus callosum and subcortical structures, where deviations from the identity line are more pronounced. The scatter plot matrix also reveals notable systematic biases between cross-sectional and longitudinal processing streams in several key regions. Particularly striking are the systematic differences in CortexVol, where the longitudinal stream consistently produces lower volume estimates compared to the

cross-sectional approach. Similar systematic biases are observed in subcortical structures, with Right-Putamen and Caudate showing consistent offsets from the identity line. CSF volumes demonstrate an opposite trend, with longitudinal processing typically yielding higher estimates than cross-sectional processing. These systematic differences suggest that the choice of processing stream can significantly impact volume estimates, even within the same FreeSurfer version (7.4.1). Interestingly, these biases appear consistent across both IBS and HC groups, maintaining the relative differences between groups regardless of the processing stream. This systematic variation between cross-sectional and longitudinal processing was not observed in the previous comparison between FS 6.0.1 and FS 7.4.1 cross-sectional analyses, indicating that these differences are specifically related to the processing stream rather than version updates. These findings highlight the importance of maintaining consistent processing approaches when comparing across studies or conducting longitudinal analyses.

The summary statistics by the mean and standard deviation for Freesurfer v. 7.4.1 cross-sectional and v. 7.4.1 longitudinal stream, respectively, is shown in the Appendix as Table A2.

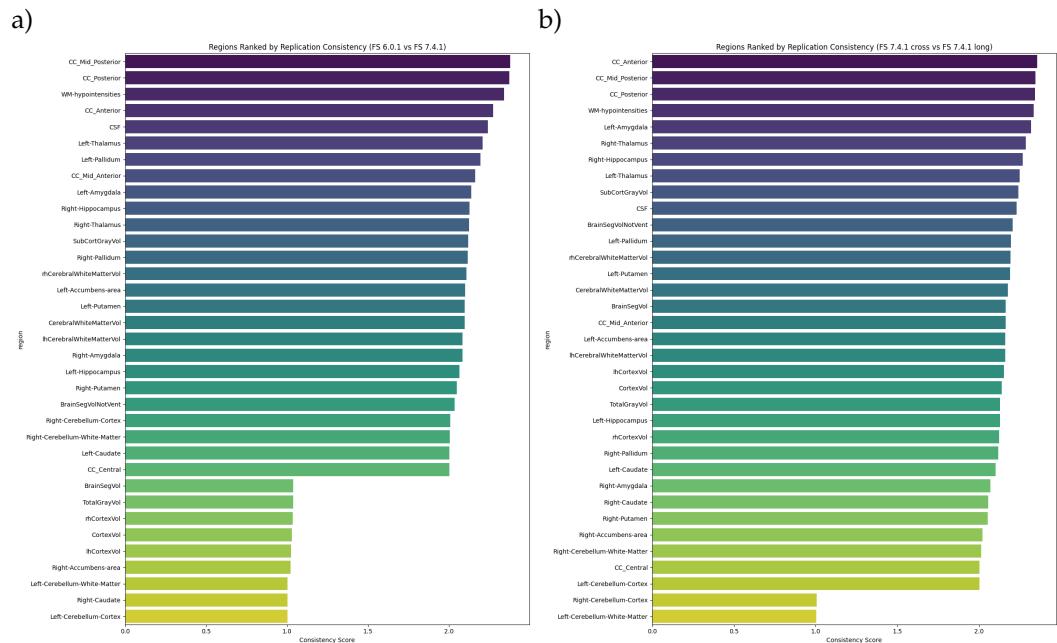
Figure 7 shows the region-wise effect size of IBS vs. HC dependent on FreeSurfer version (a) and FS 7.4.1 cross-sectional versus longitudinal stream (b).



**Figure 7.** Region-wise effect size of IBS vs. HC in comparison of FreeSurfer version and analysis pipeline. a) FS 6.0.1 cross-sectional versus F 7.4.1 cross-sectional. b) F 7.4.1 cross-sectional analysis versus FS 7.4.1 longitudinal stream (see text for details).

Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/04-comparing-FS-versions-on-same-dataset.ipynb>

Figure 8 shows regions ranked by their consistency of IBS vs. HC discrimination dependent on FreeSurfer version (a) and FS 7.4.1 cross-sectional versus longitudinal stream (b).

**Figure 8.** Barplot of regions ranked by their IBS vs HC separation consistency across FS versions.

a) FS 6.0.1 cross-sectional versus F 7.4.1 cross-sectional. b) F 7.4.1 cross-sectional analysis versus FS 7.4.1 longitudinal stream (see text for details).

Generated by: <https://github.com/arvind1/ibs-brain/blob/main/notebooks/04-computing-FS-versions-on-same-dataset.ipynb>

### Multivariate analyses: IBS versus HC

A generalized Mahalanobis distance, a partial  $\eta^2$  (Eta-squared), and a permutation test were computed to reveal how distinct the two groups (IBS and HC) are in a multidimensional space. Analog to the univariate analyses, no significant differences were observed: general Mahalanobis distance (= 2.0427), Partial  $\eta^2$  (= 0.013), the permutation test ( $p = 0.45$  (4515)). From a clinical perspective, these results indicate that the brain morphometry of IBS patients, as measured by these variables, does not differ substantially from that of healthy controls.

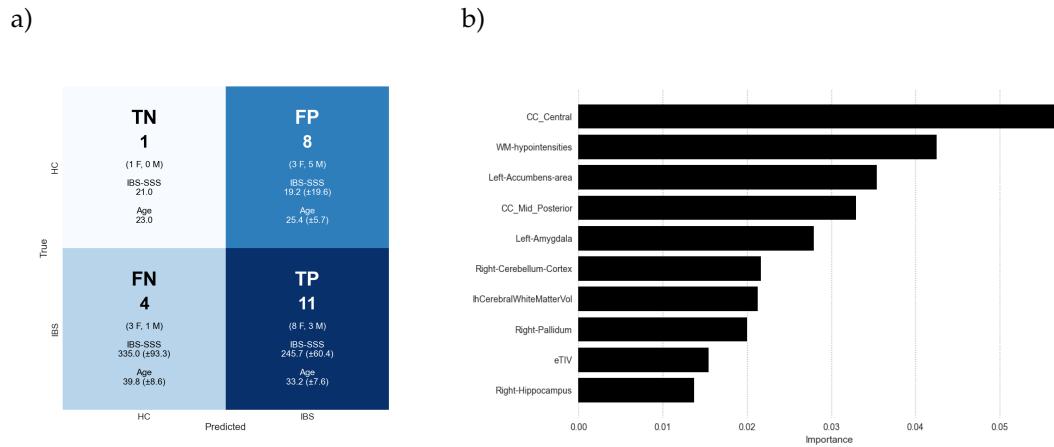
To evaluate the discriminative power of brain volumetric differences between IBS patients and healthy controls (HC) across FreeSurfer versions (6.0.1, 7.4.1 cross-sectional, and 7.4.1 longitudinal), we employed a machine learning approach combining feature selection and logistic regression. The analysis pipeline included standardization of features, selection of the most informative brain regions using F-scores, and L1-regularized logistic regression with cross-validation to assess classification performance. This approach was chosen to handle the high-dimensional nature of brain volume data while identifying the most relevant anatomical features. Results showed that none of the FreeSurfer versions achieved above-chance classification performance (ROC-AUC: FS 6.0.1 =  $0.423 \pm 0.132$ ; FS 7.4.1 cross =  $0.410 \pm 0.094$ ; FS 7.4.1 long =  $0.451 \pm 0.073$ ), indicating that brain volume differences alone are insufficient to reliably distinguish between IBS and HC groups. While the FS 7.4.1 longitudinal pipeline demonstrated slightly better and more stable performance, all versions consistently identified corpus callosum regions, white matter hypointensities, and certain subcortical structures (amygdala, thalamus) as potentially relevant features, suggesting these regions might warrant targeted investigation in further studies using additional imaging or clinical metrics beyond volumetric measures.

### Prediction of IBS versus HC from morphometric measures

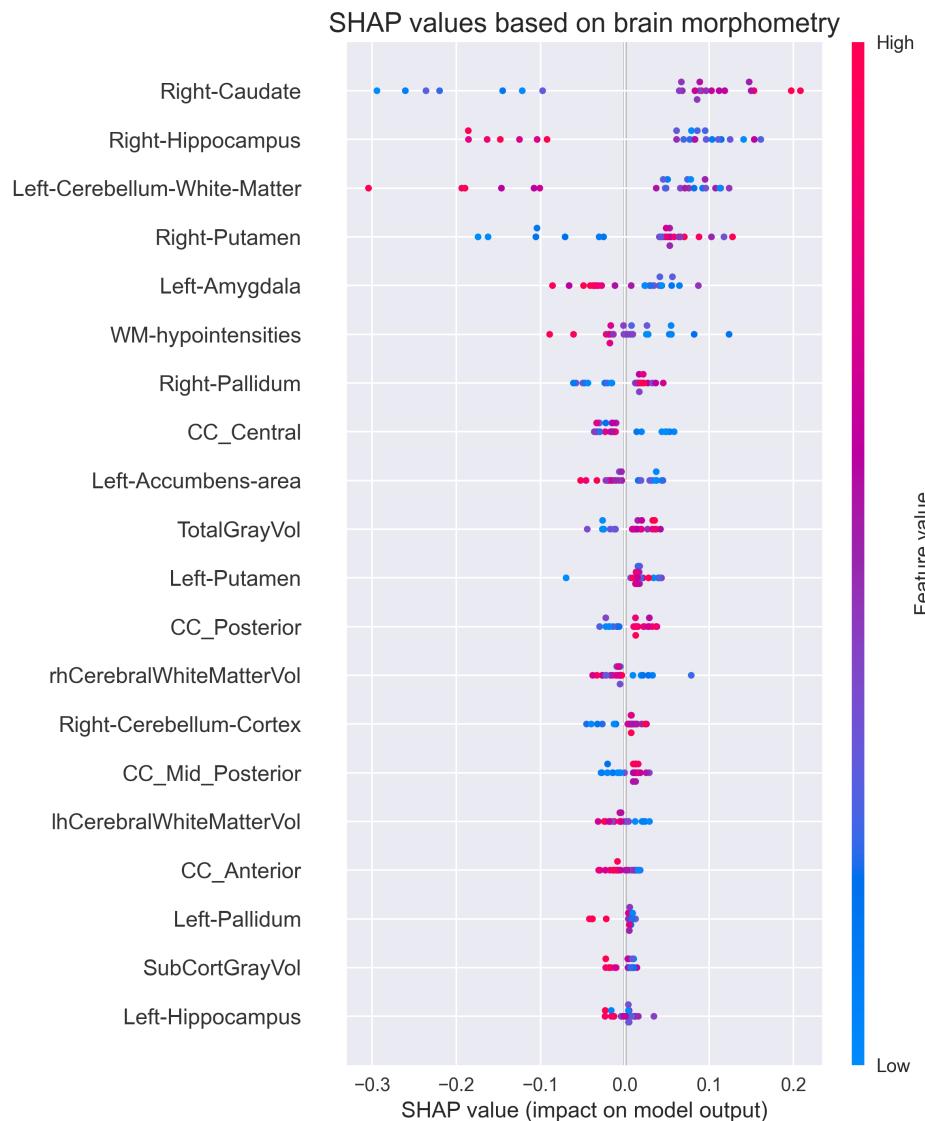
A large set of machine learning models were trained on a data set including 54 participants (43 women), of which 34 belonged to the IBS group. Among the 24 participants allocated to the test data set (15 women), 15 participants were from the IBS group.

Extreme Gradient Boosting (xgboost) was selected as the best model, with an accuracy of 0.7200 in the train-set, with a reduction (accuracy = 0.5000 ) in the test set. The confusion matrix, Figure 3 (a) shows that the majority of the IBS patients in the test set (73%) were correctly classified (11/15), while all but one HC was misclassified as IBS. The misclassified IBS patients were somewhat older than the correctly classified patients, with a higher and more distributed IBS-SSS score.

Figure 3 (b) shows the features with the strongest importance. Among the top ten features we find different parts of the corpus callosum and subcortical structures like Accumbens, Amygdala, Pallidum and Hippocampus, but also more global measures like the eTIV.



**Figure 9.** (a) Confusion matrix (XGBOOST) predicting IBS versus HC from brain morphometry. TN = true negative, FP = false positive, FN = false negative, TP = true positive. (b) Top 10 feature importance with Group (IBS vs HC) as the outcome variable and morphometric features as predictors.



**Figure 10.** SHAP values with morphometry as predictors. The SHAP values (x-axis) indicate the impact each feature (y-axis) has on the model's output, which is the probability of classifying someone as IBS patient or HC. Values to the right of 0 indicate a positive contribution (towards one class, likely HCs in this case). Values to the left indicate a negative contribution (likely towards IBS). Color gradient (Feature Value): This represents the actual feature value: Red/pink indicates a high feature value. Blue/purple indicates a low feature value.

Figure 10 shows SHAP values, which measure how different brain regions impact a predictive model's output. The right-hemispheric structures (caudate, hippocampus) appear most important for whatever outcome the model is predicting, suggesting these areas might deserve particular attention in patient assessment. There is also a pattern suggesting the basal ganglia network as a whole as important for the model's predictions, with right-hemispheric structures (especially caudate) showing stronger effects than left-hemispheric ones. Other subcortical structures like the amygdala and hippocampus also show notable effects, supporting a broader subcortical involvement in predicting IBS versus HC.

#### Univariate analysis of the cognitive features

Table 4 shows that the HC group obtained significantly higher scores on the full-scale RBANS measure, with a medium effect size. Non-parametric comparisons were used to account for non-normality distributions. Scores for the Visuospatial Index and the Attention

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Index were non-significant, with negligible effect sizes. For the two memory indexes, the Immediate Memory and Recall Indexes, the HC group scored significantly higher than the IBS group. Statistical significance was observed after Bonferroni correction solely for the Recall index, where Cliff's delta indicated a small effect size.

Variable	HC	IBS	p-value	Cliff's delta
Fullscale_RBANS	103.0 (93.0-108.0)	91.0 (85.0-100.0)	0.002	0.213
Memory_Index	100.0 (86.0-109.0)	86.0 (81.0-105.0)	0.031	0.147
Visuoaspatial_Index	97.0 (90.0-107.0)	96.0 (90.0-105.0)	0.763	0.021
Verbal skills Index	105.0 (95.0-113.0)	95.0 (89.0-111.0)	0.087	0.116
Attention_Index	98.0 (89.0-108.0)	97.0 (83.0-101.0)	0.118	0.107
Recall_Index	107.0 (92.0-113.0)	95.0 (85.0-100.0)	0.006	0.186

**Table 4.** A non-parametric analysis comparing cognitive features in the IBS and HC groups. Cliff's delta is used to estimate effect sizes.

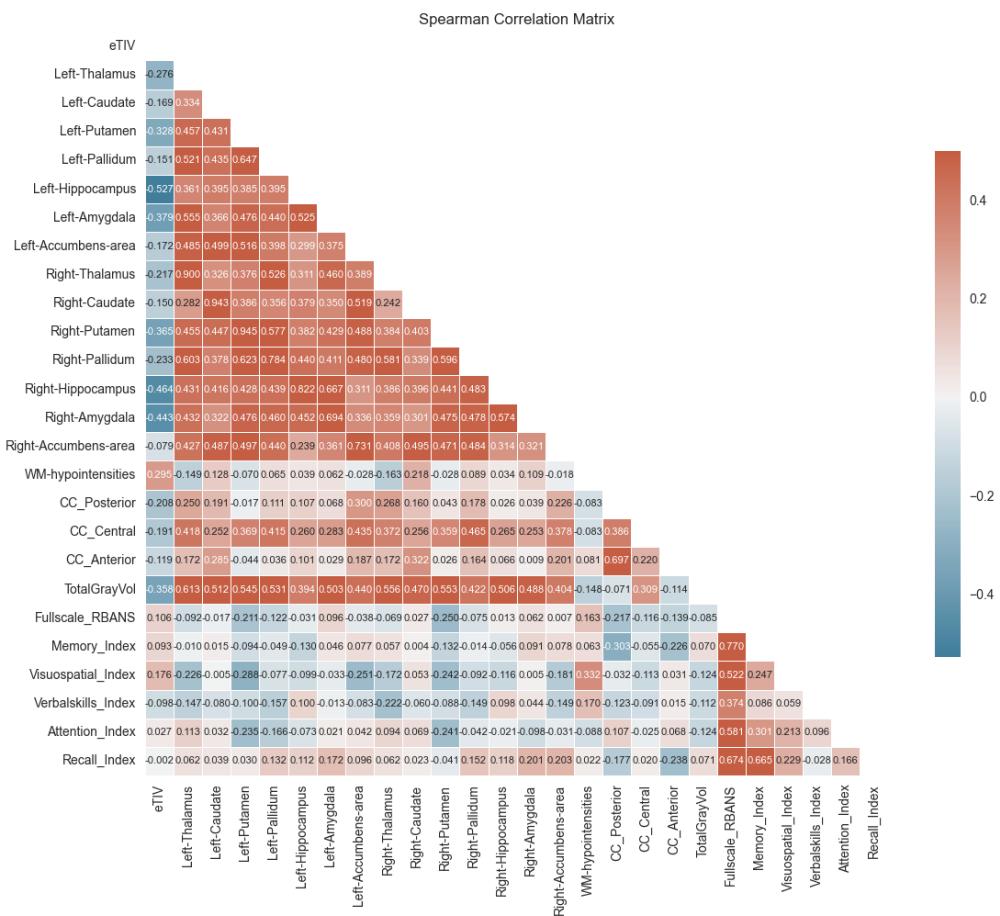
#### *Correlations between morphometric and cognitive features*

Figure 11 shows a heatmap to illustrate the pairwise correlations between the included morphometric volume measures, the RBANS indexes, and the Fullscale RBANS score. The correlations between the Fullscale score and each of the five indexes are moderate to strong for all expect the Verbal skills index. Furthermore, there is a strong correlation between the two memory scores (Immediate memory and Recall). The strongest correlations between RBANS and the morphometric measures were found between the Recall index and subcortical structures like hippocampus, amygdala and accumbens, but these correlations should still be defined as weak. The strongest correlation ( $r = .33$ ) was found between white matter hypointensities and the Visuospatial index. Most of the correlations between subcortical structures were moderate, with some very strong correlations between structures on the left and right hemisphere (e.g., Putamen  $r_s = .95$ ; Caudate  $r_s = .94$ ; Thalamus  $r_s = .90$ ; Hippocampus  $r_s = .82$  and Pallidum  $r_s = .78$ ).

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**Figure 11.** Pairwise correlations between the morphometric and cognitive variables.

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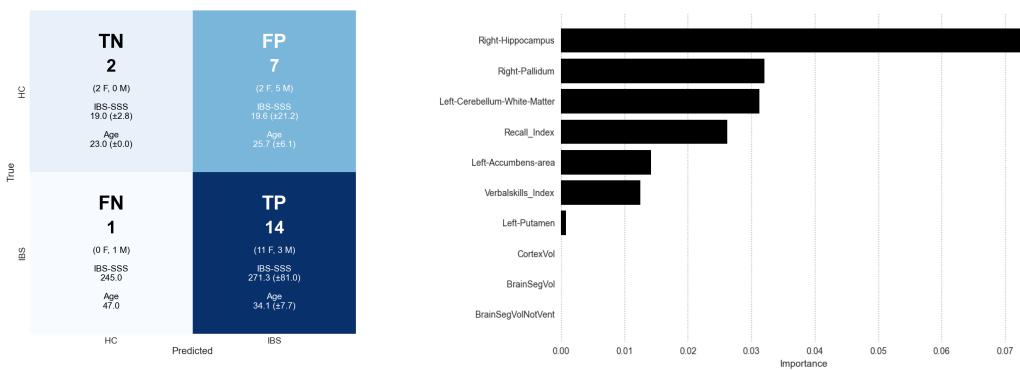
#### *Prediction of IBS versus HC from morphometric and cognitive measures*

The Extreme Gradient Boosting (xgboost) was also selected as the best model when cognitive features (the five indexes) were included as predictors together with the morphometric volume measures. An accuracy of 0.6500 in the training set improved to 0.6667 in the test set, and the confusion matrix shown in Figure 6 (a) indicates an improved generalization when cognitive features were included. The Figure shows that 93% of the IBS patients (14/15) were correctly classified (one misclassified man in the older age range). However, the specificity was lower, in that 78% in the HC group, all men, were misclassified as IBS patients.

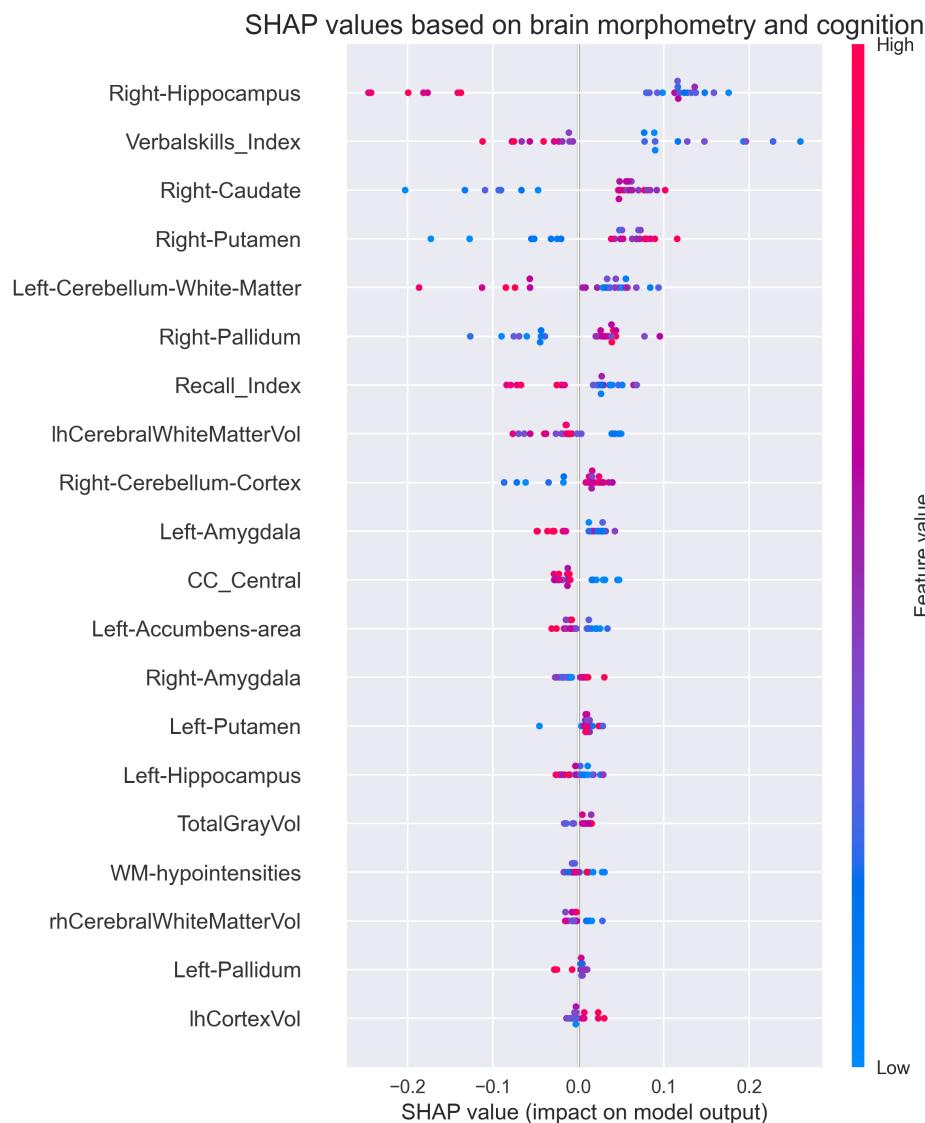
Figure 6 illustrates how different brain measurements and cognitive abilities influence our model's predictions. Each row shows a different measurement, with the most influential features at the top. The dots represent individual participants, with red dots indicating higher values and blue dots indicating lower values for each measurement. The position of each dot shows whether that particular value pushes the model's prediction higher (towards the right) or lower (towards the left). The Right Hippocampus, shown at the top, has the strongest overall impact: when some people have high values (red dots), this strongly pushes predictions in one direction, while low values (blue dots) push predictions in the other direction. Verbal skills also shown important patterns, with a wide spread

across different participants, suggesting that both high and low verbal skills can be important indicators, depending on the individual case. Among subcortical regions, Right Caudate and Right Putamen demonstrate particularly notable influences. These patterns reveal meaningful differences between our two groups in both brain structure and cognitive abilities. Particularly important are the variations we see in verbal abilities and in memory-related brain regions like the hippocampus, which could inform how we approach patient care and communication strategies.

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**Figure 12.** (a) Confusion matrix (XGBOOST) predicting IBS versus HC from brain morphometry and cognition. (b) Top 10 feature importance (permutation importance, 100 repeats) with Group (IBS vs HC) as the outcome variable and morphometric and cognitive features as predictors



**Figure 13.** SHAP values with morphometry and cognition as predictors.

Note:X-axis (SHAP values): The SHAP values indicate the impact each feature has on the model's output, which is the probability of classifying someone as IBS patient or HC.

## Discussion

Our study yields two key methodological insights and one substantial findings regarding brain structure and function in IBS. First, we were unable to replicate the morphometric differences between the IBS and healthy control group reported by Skrobisz et al. [22], regardless of whether we used FreeSurfer version 6.0.1 or 7.4.1. Second, we observed substantial discrepancies in morphometric measurements between these software versions, highlighting the critical importance of considering methodological factors in neuroimaging research. Application of advanced multivariate and machine learning techniques to investigate brain-behavior relationships in IBS is another primary contribution. While morphometric features alone proved insufficient for reliable group discrimination, the integration of cognitive performance measures with brain morphometry substantially improved classification accuracy. Specifically, our analyses revealed that two cognitive domain indices, combined with volumetric measures of subcortical structures—particularly the hippocampus and basal ganglia—provided robust discrimination between IBS patients and healthy controls. The consistency of these findings across different approaches to feature importance analyses strengthens their validity and suggests a fundamental relationship

between brain structure, cognitive function, and IBS symptomatology. This observation aligns with views of IBS as a disorder involving complex interactions between central nervous system function and gastrointestinal symptoms, rather than purely peripheral manifestations.

#### *Brain Structures involved in discriminating between IBS and HC*

Our results showed that subcortical structures, particularly within the basal ganglia, played a key role in distinguishing IBS patients from healthy controls. While traditionally associated with motor control, the basal ganglia also critically influence reward processing, habit formation, and pain modulation - functions directly relevant to IBS symptomatology, and the impact on patients' experience of gastrointestinal symptoms. These findings align with recent results from a UK Biobank study [19], which also highlighted the importance of hippocampal and basal ganglia structures, including the Pallidum and Caudate, in IBS. Beyond the basal ganglia, several other subcortical structures relevant to IBS symptomatology emerged as discriminators. The nucleus accumbens, fundamental to reward processing and motivation, may mediate the emotional and motivational aspects of chronic pain in IBS. Dysfunction in this structure could explain the intensified emotional distress and pain sensitivity commonly reported by IBS patients [9]. Similarly, the amygdala appears significant, particularly given its connection to pain-modulation and emotion-processing networks, including the prefrontal cortex and insula. This aligns with previous research [36] demonstrating enhanced amygdala-insula connectivity in IBS patients. Although our results differ from Skrobisz et al.'s [22] findings regarding thalamic involvement, other studies have supported its role in IBS. Diffusion tensor imaging has revealed altered thalamic organization in IBS patients, with reduced fractional anisotropy and increased mean diffusivity [37]. These alterations suggest compromised structural integrity of thalamic circuits, potentially affecting pain processing and sensory integration. The involvement of corpus callosum should also be mentioned, as interhemispheric integration is crucial for visceral sensation processing, pain modulation [38] as well as in mental disorders [39]. Taken together, our findings support that integrated neural signatures are involved in predicting IBS [40].

#### *Integration of Cognitive Performance and Brain Structure in IBS*

The enhanced diagnostic accuracy by including cognitive measures strongly support that IBS should be understood as a disorder of the gut-brain interaction [14,41]. The brain's integral role in cognitive, emotional, and autonomic regulation suggests that these manifestations are fundamentally interconnected rather than merely coincidental. The prominent role of hippocampal volume was a principal finding. The fundamental role of Hippocampus in cognitive processing is well known [42], and was supported by the Recall index being identified as another feature with strong importance. The role of Verbal skills was more surprising. Although research has established connections between memory systems and language processing, particularly in semantic memory organization [43], a negligible correlation between the two indices suggests that IBS affects multiple cognitive domains through independent mechanisms.

Our findings may also have implications for other somatic and psychiatric disorders, like Alzheimer's disease, Parkinson's disease, and major depression. The gut-brain axis are involved in all these diseases, which also are characterized by cognitive impairment. Recent research has identified potential pathways linking gut microbiota alterations to neurological function, particularly through inflammatory responses and tryptophan metabolism [44,45]. The emergence of the microbiota-gut-brain axis as a key framework [46] offers new perspectives on how peripheral inflammation might influence both brain structure and cognitive function in IBS. This integrated view suggests that cognitive assessment, combined with brain morphometry, might provide valuable insights not only for IBS but

for a broader spectrum of gut-brain disorders.

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#### *Brain-Gut Axis: Implications for Understanding and Treating IBS*

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Our findings should have important implications for clinical practice and treatment development. The observed relationship between brain structure, cognitive function, and IBS symptomatology suggests that effective interventions should target multiple domains simultaneously. Such a multifaceted approach recognizes IBS as a complex disorder requiring coordinated intervention across multiple domains.

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Future research directions should expand upon these findings through multimodal investigation. Integration of functional neuroimaging, gut microbiome analysis, and broader clinical assessment [19] could provide a more comprehensive understanding of IBS pathophysiology. Particularly crucial will be longitudinal studies to determine the temporal relationship between brain changes and symptom development. Such studies would allow us to track the evolution of cognitive and structural alterations over time, identify early markers of disease progression, and evaluate the impact of various therapeutic interventions. This temporal perspective is essential for understanding whether observed brain changes represent cause or consequence of IBS symptoms.

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This comprehensive approach to understanding IBS aligns with the emerging paradigm of precision medicine. By considering the full spectrum of biological, cognitive, and behavioral manifestations, we may better identify patient subgroups and develop more personalized treatment strategies. The integration of brain structure, cognitive function, and clinical symptoms represents a promising framework for advancing both our understanding and treatment of this complex disorder. Ultimately, this integrated perspective may lead to more effective, personalized interventions that address the full range of IBS manifestations.

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#### *Strengths and Limitations: Critical Evaluation and Future Directions*

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Although contributing through its multimodal analytical approach, several limitations warrant discussion. The primary limitation concerns sample size, which particularly constrained our ability to conduct robust sex/gender-based analyses. This limitation is especially noteworthy given the evidence for substantial sex/gender differences in IBS presentation, progression, and treatment response [47]. The importance of sex/gender considerations in IBS research has become increasingly apparent. Clinical presentations show clear sex-based patterns, with IBS-C predominating in women and IBS-D in men [48]. These differences reflect complex interactions between biological and environmental factors. Sex hormones, particularly estrogen and progesterone, influence both gastrointestinal function and pain processing in the central nervous system [49]. Recent research has revealed sex-based differences extending to gut microbiota composition [50] and sensory processing. Notably, Labus et al. [20] demonstrated enhanced sensory sensitivity in women with IBS, potentially related to sex-specific morphometric variations in brain structure.

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An inability to fully account for IBS symptom severity in our analyses was another limitation. Recent work by Li et al. [19] has demonstrated that symptom severity correlates significantly with both cognitive performance and brain volumetric measures, particularly in regions associated with emotional processing and cognitive control. This finding suggests that future studies should incorporate detailed severity measures to better understand the relationship between symptom intensity and brain-behavior patterns.

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The present study's methodological strengths lie in its comprehensive multivariate approach to analyzing brain-behavior relationships. This approach better captures the complex interactions between multiple brain regions and cognitive measures, providing a more nuanced understanding than traditional univariate analyses. However, we acknowledge important limitations. While our sample exceeds that of many comparable studies, multivariate analyses and machine learning approaches typically benefit from larger datasets.

To address this limitation, we implemented robust cross-validation procedures, including 10-fold validation and hold-out test sets, thereby reducing the risk of overfitting and enhancing the generalizability of our findings. Moreover, the results point to several important directions for future research. First, larger-scale studies are needed to validate and extend our multivariate findings. Such studies should maintain rigorous methodological standards while increasing statistical power. Second, standardization of neuroimaging analysis protocols, including careful documentation of software versions and processing parameters, is crucial for reproducibility. Third, the field would benefit from systematic investigation of how different analysis approaches might influence morphometric findings in IBS research. Overall, future studies should consider implementing standardized protocols for both imaging and cognitive assessment, facilitating meta-analyses and enabling more direct comparisons across studies. This standardization, combined with transparent reporting of methodological details, would strengthen the field's ability to build cumulative knowledge about brain-gut interactions in IBS. Longitudinal studies represent a particularly important future direction. Such studies could address crucial questions about the temporal dynamics of brain-gut interactions in IBS, including whether observed structural and cognitive changes precede or follow symptom development. Longitudinal data would also enable better prediction of disease trajectories and treatment responses, potentially informing personalized interventions such as dietary modifications (e.g., Low FODMAP diet) or targeted cognitive interventions. The combination of longitudinal design with multimodal assessment (including brain structure, cognitive function, and clinical symptoms) could provide unprecedented insights into the development and progression of IBS.

## Conclusions and Future Directions

The present study advances our understanding of brain-gut interactions in IBS through several key contributions. First, our comprehensive multivariate analyses reveal the inherent complexity of IBS pathophysiology, demonstrating that single-modality approaches may be insufficient for characterizing this multifaceted disorder. While we did not replicate previously reported volumetric differences in thalamic structure, our machine learning analyses uncovered more subtle and complex patterns of brain-behavior relationships. Particularly noteworthy was the finding that morphometric features gain discriminative power when integrated with cognitive measures, especially in subcortical regions including the hippocampus and basal ganglia. These results strongly support a systems-level conceptualization of IBS, where the condition emerges from complex interactions between neural structure, cognitive function, and gastrointestinal symptoms. This perspective suggests that effective characterization and treatment of IBS requires consideration of multiple biological and cognitive markers rather than focusing on isolated symptoms or structures. The successful integration of structural and functional measures in our analyses points toward more sophisticated approaches for both diagnosis and treatment planning. Moving forward, several research priorities emerge from our findings. Large-scale validation studies are needed to confirm the reliability and generalizability of our brain-cognition relationships across diverse patient populations. Such studies should incorporate standardized protocols for both imaging and cognitive assessment to facilitate cross-study comparisons. Longitudinal investigations are particularly crucial for understanding how these markers evolve over time and relate to treatment response. Additionally, future research should explore how individual differences in brain structure and cognitive function might predict treatment outcomes, potentially enabling more personalized therapeutic approaches. Ultimately, our findings suggest that advancing IBS treatment may require a fundamental shift toward integrated, multimodal assessment approaches that capture both structural and functional aspects of brain-gut interactions. This more comprehensive understanding of IBS pathophysiology could lead to more effective, personalized interventions that address the full spectrum of patient symptoms and experiences.

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**Institutional Review Board Statement:** The B-BGM project was approved by the Southeast Regional Ethical Committees (REC) for medical and health research ethics in Norway (REK2015-01621). All participants provided written consent to participate, and the project was conducted following the ethical requirements from the Declaration of Helsinki. The project is registered at [www.clinicaltrials.gov](http://www.clinicaltrials.gov) (#NCT04296552). 1039  
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**Informed Consent Statement:** Informed consent was obtained from all subjects involved in the study. 1044  
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**Data Availability Statement:** The implementation of the complete workflow, the setup of the corresponding conda environment, the cleaned input dataset in .csv format, and code for all tables and figures in the Results section are available as *Jupyter notebooks* at <https://arvidl.github.com/ibs-brain>. 1046  
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## Abbreviations

The following abbreviations are used in the manuscript:

AUC	Area Under Curve	1051
CM	Confusion matrix	1052
Cohen's d	effect size	1053
Cliff's delta	effect size	
DGBI	Disorders of the gut-brain interaction	
FS	Freesurfer	
GI	Gastrointestinal	
GitHub	Meeting platform for collaboration	
HC	Healthy Control	
IBS	Irritable bowel syndrome	
IBS-SSS	IBS Severity Scoring System	1057
IQR	Inter Quartile Range	
ML	Machine-learning	
MRI	Magnetic Resonance Imaging	
RBANS	Repeatable Battery for the Assessment of Neuropsychological Status	
RF	Random Forest	
ROC	Operating Characteristic	
SHAP	SHapley Additive exPlanations	
SD	Standard deviation	
SHAP	SHapley Additive exPlanations	
XGBoost	eXtreme Gradient Boosting	

## Appendix A Supplementary tables and figures

### Appendix A.1 Comparing Freesurfer 6.0.1 and FreeSurferr 7.4.1 cross-sectional

Table A1 gives the summary statistics, mean and standard deviation from HC and IBS patients in the Bergen cohort on each of the 35 included brain regions (also reported by Skrobisz et al. [22]) derived from the `aseg.stats` files using cross-sectional Freesurfer 6.0.1 and Freesurfer 7.4.1, respectively.

**Table A1.** Comparison of Brain Region Volumes in IBS Patients and Healthy Controls.

Brain Region	Bergen Cohort FS 6.0.1				Bergen cohort FS 7.4.1			
	HC (N=29)		IBS (N=49)		HC (N=29)		IBS (N=49)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Left-Cerebellum-WM	0.010496	0.000915	0.010483	0.000924	0.010603	0.000932	0.010607	0.001027
Left-Cerebellum-Cortex	0.038939	0.003435	0.039314	0.003733	0.038066	0.003526	0.038056	0.003684
Left-Thalamus	0.005232	0.000464	0.005144	0.000393	0.005236	0.000522	0.005114	0.000458
Left-Caudate	0.002356	0.000259	0.002355	0.000306	0.002346	0.000283	0.002317	0.000294
Left-Putamen	0.003479	0.000385	0.003441	0.000386	0.003438	0.000396	0.003370	0.000329
Left-Pallidum	0.001405	0.000154	0.001374	0.000107	0.001380	0.000136	0.001358	0.000095
Left-Hippocampus	0.002913	0.000272	0.002896	0.000242	0.002926	0.000251	0.002895	0.000243
Left-Amygdala	0.001218	0.000097	0.001203	0.000105	0.001228	0.000133	0.001190	0.000111
Left-Accumbens-area	0.000427	0.000069	0.000421	0.000057	0.000424	0.000061	0.000400	0.000057
CSF	0.000670	0.000120	0.000702	0.000141	0.000658	0.000114	0.000689	0.000130
Right-Cerebellum-WM	0.009973	0.000891	0.009979	0.000851	0.010052	0.000934	0.010108	0.001015
Right-Cerebellum-Cortex	0.039719	0.003445	0.039978	0.003760	0.038881	0.003534	0.038912	0.003673
Right-Thalamus	0.005120	0.000438	0.005071	0.000358	0.005190	0.000455	0.005053	0.000413
Right-Caudate	0.002438	0.000240	0.002439	0.000301	0.002418	0.000286	0.002402	0.000285
Right-Putamen	0.003506	0.000366	0.003489	0.000351	0.003487	0.000402	0.003466	0.000322
Right-Pallidum	0.001323	0.000126	0.001301	0.000107	0.001321	0.000137	0.001306	0.000118
Right-Hippocampus	0.003013	0.000240	0.002983	0.000229	0.003049	0.000230	0.002986	0.000235
Right-Amygdala	0.001284	0.000087	0.001271	0.000098	0.001269	0.000107	0.001260	0.000106
Right-Accumbens-area	0.000428	0.000053	0.000427	0.000061	0.000434	0.000054	0.000435	0.000057
WM-hypointensities	0.000791	0.000306	0.000688	0.000253	0.000787	0.000481	0.000667	0.000244
CC_Posterior	0.000652	0.000096	0.000702	0.000113	0.000645	0.000096	0.000685	0.000113
CC_Mid_Posterior	0.000369	0.000067	0.000401	0.000071	0.000366	0.000069	0.000394	0.000073
CC_Central	0.000395	0.000089	0.000391	0.000105	0.000390	0.000091	0.000390	0.000101
CC_Mid_Anterior	0.000379	0.000081	0.000409	0.000113	0.000384	0.000078	0.000400	0.000105
CC_Anterior	0.000623	0.000096	0.000650	0.000101	0.000608	0.000098	0.000646	0.000112
BrainSegVol	0.804644	0.024872	0.805581	0.023967	0.792112	0.037690	0.786845	0.028349
BrainSegVolNotVent	0.792235	0.025106	0.791323	0.024898	0.779857	0.037538	0.772948	0.030305
lhCortexVol	0.166698	0.008003	0.166929	0.009510	0.164771	0.010207	0.163181	0.010196
rhCortexVol	0.166137	0.008276	0.166462	0.009388	0.164149	0.010295	0.162912	0.009834
CortexVol	0.332835	0.016110	0.333391	0.018798	0.328920	0.020369	0.326092	0.019888
lhCerebralWhiteMatterVol	0.159895	0.008578	0.159148	0.008757	0.157377	0.010114	0.155820	0.009472
rhCerebralWhiteMatterVol	0.159252	0.008291	0.158267	0.009384	0.156808	0.010522	0.154840	0.009950
CerebralWhiteMatterVol	0.319147	0.016780	0.317415	0.018079	0.314184	0.020552	0.310659	0.019351
SubCortGrayVol	0.040924	0.002583	0.040629	0.002364	0.040864	0.002871	0.040194	0.002433
TotalGrayVol	0.453068	0.022076	0.453961	0.024324	0.446625	0.027101	0.443252	0.025556
eTIV [mm <sup>3</sup> ]	1468820.2	155501.4	1426237.4	136412.8	1494273.2	171472.3	1462310.8	144145.1

Note: All volumes are normalized to estimated total intracranial volume (eTIV)

HC = Healthy Controls; IBS = Irritable Bowel Syndrome; SD = Standard Deviation; WM = White Matter

Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/04-computing-FS-versions-on-same-dataset.ipynb>

### Appendix A.2 Comparing FreeSurfer 7.4.1 cross-sectional and longitudinal stream

Table A2 gives the summary statistics, mean and standard deviation from HC and IBS patients in the Bergen cohort on each of the 35 included brain regions derived from the `aseg.stats` files using Freesurfer 7.4.1 cross-sectional analysis and Freesurfer 7.4.1 longitudinal stream, respectively.

**Table A2.** Comparison of Brain Region Volumes in Bergen cohort, FS 7.4.1 cross-sectional vs. FS 7.4.1 longitudinal stream

Brain Region	FS 7.4.1 cross-sectional				FS 7.4.1 longitudinal stream			
	HC (N=29)		IBS (N=49)		HC (N=29)		IBS (N=49)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Left-Cerebellum-White-Matter	0.010603	0.000932	0.010607	0.001027	0.010888	0.001076	0.010688	0.001044
Left-Cerebellum-Cortex	0.038066	0.003526	0.038056	0.003684	0.037438	0.003633	0.037232	0.003575
Left-Thalamus	0.005236	0.000522	0.005114	0.000458	0.005385	0.000517	0.005262	0.000473
Left-Caudate	0.002346	0.000283	0.002317	0.000294	0.002506	0.000313	0.002474	0.000315
Left-Putamen	0.003438	0.000396	0.003370	0.000329	0.003702	0.000418	0.003621	0.000369
Left-Pallidum	0.001380	0.000136	0.001358	0.000095	0.001415	0.000156	0.001373	0.000098
Left-Hippocampus	0.002926	0.000251	0.002895	0.000243	0.002970	0.000273	0.002925	0.000244
Left-Amygdala	0.001228	0.000133	0.001190	0.000111	0.001253	0.000138	0.001203	0.000111
Left-Accumbens-area	0.000424	0.000061	0.000400	0.000057	0.000440	0.000072	0.000429	0.000070
CSF	0.000658	0.000114	0.000689	0.000130	0.000712	0.000127	0.000742	0.000130
Right-Cerebellum-White-Matter	0.010052	0.000934	0.010108	0.001015	0.010218	0.001008	0.010231	0.000955
Right-Cerebellum-Cortex	0.038881	0.003534	0.038912	0.003673	0.038471	0.003722	0.038176	0.003654
Right-Thalamus	0.005190	0.000455	0.005053	0.000413	0.005475	0.000474	0.005341	0.000456
Right-Caudate	0.002418	0.000286	0.0002402	0.000285	0.002608	0.000310	0.002577	0.000303
Right-Putamen	0.003487	0.000402	0.003466	0.000322	0.003788	0.000418	0.003766	0.000375
Right-Pallidum	0.001321	0.000137	0.001306	0.000118	0.001350	0.000156	0.001330	0.000120
Right-Hippocampus	0.003049	0.000230	0.002986	0.000235	0.003102	0.000245	0.003034	0.000241
Right-Amygdala	0.001269	0.000107	0.001260	0.000106	0.001332	0.000115	0.001323	0.000115
Right-Accumbens-area	0.000434	0.000054	0.000435	0.000057	0.000503	0.000065	0.000507	0.000063
WM-hypointensities	0.000787	0.000481	0.000667	0.000244	0.000757	0.000644	0.000607	0.000274
CC_Posterior	0.000645	0.000096	0.000685	0.000113	0.000632	0.000097	0.000669	0.000112
CC_Mid_Posterior	0.000366	0.000069	0.000394	0.000073	0.000350	0.000066	0.000375	0.000075
CC_Central	0.000390	0.000091	0.000390	0.000101	0.000364	0.000082	0.000363	0.000091
CC_Mid_Anterior	0.000384	0.000078	0.000400	0.000105	0.000361	0.000071	0.000379	0.000101
CC_Anterior	0.000608	0.000098	0.000646	0.000112	0.000587	0.000096	0.000620	0.000093
BrainSegVol	0.792112	0.037690	0.786845	0.028349	0.798892	0.038555	0.790867	0.029736
BrainSegVolNotVent	0.779857	0.037538	0.772948	0.030305	0.785834	0.038695	0.776074	0.031471
lhCortexVol	0.164771	0.010207	0.163181	0.010196	0.170575	0.010496	0.168994	0.010110
rhCortexVol	0.164149	0.010295	0.162912	0.009834	0.170608	0.010675	0.168900	0.009646
CortexVol	0.328920	0.020369	0.326092	0.019888	0.341183	0.021084	0.337894	0.019676
lhCerebralWhiteMatterVol	0.157377	0.010114	0.155820	0.009472	0.153875	0.009612	0.151551	0.009309
rhCerebralWhiteMatterVol	0.156808	0.010522	0.154840	0.009950	0.152915	0.009898	0.150025	0.009752
CerebralWhiteMatterVol	0.314184	0.020552	0.310659	0.019351	0.306790	0.019437	0.301576	0.018991
SubCortGrayVol	0.040864	0.002871	0.040194	0.002433	0.042919	0.003186	0.042213	0.002739
TotalGrayVol	0.446625	0.027101	0.443252	0.025556	0.459994	0.028590	0.455681	0.025742

Note: All volumes are normalized to estimated total intracranial volume (eTIV)

HC = Healthy Controls; IBS = Irritable Bowel Syndrome; SD = Standard Deviation; WM = White Matter

Generated by: <https://github.com/arvidl/ibs-brain/blob/main/notebooks/04-computing-FS-versions-on-same-dataset.ipynb>