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Article

Depressive Symptoms and Gut Microbiota after Bowel Preparation and Colonoscopy: A Pre-Post Intervention Study

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Abstract: Objective: Mechanical bowel preparation is essential for the visualisation of the colon during colonoscopy, and previous studies have identified changes in gut microbiota composition after bowel preparation and colonoscopy. Considering the gut microbiota is increasingly implicated in mood, we aimed to explore the potential impact of this intervention on mood and the microbiota-gut-brain axis. **Methods:** We conducted a pre-post intervention study, with timepoints of one week before and one month after bowel preparation and colonoscopy, in community-dwelling adults referred for colonoscopy. Our primary outcome was change in average Hospital Anxiety and Depression Scale depression sub-scores using generalised estimating equations. We also examined changes in average anxiety, stress, and quality of life scores, and faecal microbiota composition and potential function using 16S rRNA sequencing. We further explored associations between changes in depressive symptoms and the gut microbiota and conducted post-hoc analyses to explore potential effect modifiers. **Key Results:** 59 participants were included in the analyses. Depressive symptoms decreased one-month post-procedure compared to baseline (adjusted $\beta=-0.64$; 95%CI: $-1.18, -0.11$). Irritable bowel syndrome (IBS) appeared to moderate this relationship (1.78; 0.292, 3.26); depressive symptoms increased in those with, and decreased in those without, IBS. Changes in gut microbiome composition included reduced alpha diversity, modest effects on beta-diversity, and increases in genera associated with potential health benefits, one-month post-procedure compared to baseline. Increases in the CLR-transformed abundance of *Ruminococcaceae* UCG-009 were associated with improvements in depressive symptoms. **Conclusions:** There is preliminary evidence of a potential mental health effect of bowel preparation and colonoscopy, which may be associated with changes to the gut microbiota. There may also be a differential mental health impact of this procedure for those with and without IBS. Considering the limitations of our study, further research is required to confirm these findings and their clinical relevance.

Keywords: gut microbiota; colonoscopy; bowel preparation; depression; microbiota-gut-brain axis; mental health; psychiatry; gastroenterology

1. Introduction

The microbiome-gut-brain axis has emerged as a contributing factor to the aetiology and pathophysiology of depression^{1,2}. Preclinical evidence suggests that many physiological pathways influenced by the gut microbiota are also implicated in depression³. In humans, differences in gut microbiota composition have been associated with depression^{4,5}, and interventions that modulate the gut microbiota, such as diet⁶⁻⁸ and probiotic supplementation⁹, can alleviate depressive symptoms. Manipulation of the gut microbiota may also further our understanding of how the gut microbiota influences, and can be targeted to treat, depressive disorders. Reciprocally, understanding how common interventions affect the gut microbiota may have important clinical and pathophysiological implications.

Mechanical bowel preparations are osmotic laxatives consumed prior to colonoscopy. Bowel preparation is essential for visualisation of the colon and the detection of abnormalities during colonoscopy^{10,11}, and previous research has shown an impact of bowel preparation on the gut microbiome. Indeed, osmotic laxatives have strong associations with gut microbiota composition and function, with large effect sizes¹². Previous intervention studies that have examined the impact of bowel preparation and colonoscopy (hereafter referred to as the 'procedure') on the gut microbiota have reported mixed results. Whilst decreases in the number and distribution (i.e., alpha diversity) and relative abundances of bacterial taxa have been observed immediately after bowel preparation (before colonoscopy)^{13,14}, very minimal or no changes have also been described^{15,16}. Differences in gut microbiota composition have been reported one week^{14,15,17-19} to one month^{13,17,20,21} post-procedure compared to baseline, however in other studies there were no statistically significant differences at these follow up time points^{17,19,20}. Few studies have measured changes in gut microbiota composition beyond one month after bowel preparation and colonoscopy, and those that have provided limited evidence of ongoing changes six weeks and six months later^{18,22}.

Whilst studies have aimed to characterise changes in gut microbiota composition, no studies, to our knowledge, have investigated changes in depressive symptoms after bowel preparation and colonoscopy, nor explored the potential associations between changes in gut microbiota composition and changes in depressive symptoms. Considering the millions of colonoscopies conducted annually^{23,24}, there is a need to better understand the potential impact of this procedure on the microbiome-gut-brain axis.

Therefore, the aim of this study was to investigate changes in self-reported depressive symptoms one week before versus one month after bowel preparation and colonoscopy. We also aimed to explore changes in gut microbiota composition and functional potential post-procedure and identify any associations with changes in depressive symptoms.

2. Materials and Methods

2.1. Study Design

The data used for this manuscript were derived from the Micro-Scope study, which used a pre-post intervention study design to investigate changes in depressive symptoms and gut microbiota composition after bowel preparation and colonoscopy. Data are reported in accordance with the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) Statement²⁵, and the Strengthening the Organising and Reporting of Microbiome Studies (STORMS) checklist²⁶. This study had ethical approval from the Barwon Health (#15-129), Epworth Healthcare (#EH2016-146), and Deakin University (#2016-391) Human Research Ethics Committees. Microbial data were not uploaded to an online public repository as consent for the sharing of data was not obtained. This study was pre-registered on the Open Science Framework (OSF): <https://osf.io/fv7xd/>.

2.2. Participants and Setting

Participants were community-dwelling adults referred for colonoscopy between May 2017 and November 2018 at University Hospital Geelong, Australia. Colonoscopies were performed at

University Hospital or Epworth Hospital, Geelong, Australia. Recruitment of participants occurred at the time of their initial outpatient consultation with the General Surgery or Gastroenterology services. Any adults referred for colonoscopy during the study time frame were considered for recruitment. Exclusion criteria included those: highly dependent on medical care; unable to give informed consent (e.g., language barriers, significant intellectual or cognitive disability); or children under 18 years of age. There were no exclusions for antibiotic use. Those deemed eligible to participate by their clinician were directed to a member of the research team to discuss the study further and provide informed consent. Those with a diagnosis of cancer post-colonoscopy were withdrawn and did not provide follow up data.

2.3. Data Collection

Baseline data were collected one week pre-procedure. Participants completed paper-based questionnaires and collected a fresh faecal sample into a sterile collection jar with a scoop lid at home. The sample was stored in their freezer (-20°C) for approximately one week until transported on ice to the research team on the day of colonoscopy, when it was transferred to a -80°C freezer for storage until DNA extraction. Luminal aspirates were collected during colonoscopy to demonstrate the potential immediate changes in alpha diversity associated with the procedure. Faecal residue from within the colon was aspirated into a sterile collection jar (which had been flushed with saline for collection if necessary) and placed on ice, then transferred to a -80°C . One month post-procedure, participants completed another set of questionnaires, and collected a final faecal sample at home as previously.

2.4. Intervention

Bowel preparation was prescribed and carried out as per normal advice and practice for the colonoscopy service. Participants were instructed to commence a low-fibre (“white”) diet two days before their colonoscopy, and then fast for 12–24 hours prior to their procedure and consume a sodium picosulfate-based bowel preparation product in three separate doses. Adequacy of bowel preparation was reported during colonoscopy by the endoscopist using a modified overall Boston Bowel Preparation Scale score²⁷.

2.5. Outcomes

Depressive symptoms (primary outcome) were measured pre- and post-procedure using the depression sub-score of the Hospital Anxiety and Depression Scale (HADS)²⁸. All other outcomes were considered exploratory. The severity of depressive symptoms was measured using the Patient Health Questionnaire-9 (PHQ-9)²⁹; anxiety symptoms were measured using the anxiety sub-score of the HADS²⁸; total, psychosocial, and physical quality of life were measured using the Assessment of Quality of Life-8 (AQOL-8D)³⁰; and stress was measured using the Perceived Stress Scale (PSS)³¹. Gut microbiota alpha diversity was measured using the Shannon index — a within-sample index of both richness and evenness — and the number of observed amplicon sequencing variants (ASVs) at the genus level. Beta-diversity was measured using the Aitchison distance and differential abundances of ASVs were determined using centred-log ratio transformed count abundance data.

2.6. Covariates

Age at the time of recruitment and sex (male/female/other) were obtained from medical records. A triage nurse collected participant height and weight at their initial outpatient consultation, which were used to calculate body mass index (BMI; kilograms/metre²). Participants self-reported their residential postcode and suburb, which were used to calculate socioeconomic status using an area-based measure called the Index of Relative Socio-economic Advantage and Disadvantage (IRSAD)³². Each suburb has an IRSAD classification ranging from 1-10, where a lower IRSAD score suggests greater disadvantage. Smoking status, lifetime history of medical conditions (including depression), and current medication use were self-reported. Diet quality was measured using the Simple Dietary

Questionnaire³³, where the total score (out of 100) rated dietary adherence to the Australian Dietary Guidelines, higher scores representing greater compliance³⁴. The ROME III Diagnostic Questionnaire for Adult Functional Gastrointestinal Disorders was used to determine if participants met irritable bowel syndrome (IBS) criteria³⁵. Colonoscopy indication and outcomes were obtained from medical records.

2.7. DNA Extraction

Microbial DNA extraction from stool was performed using the commercial QIAamp Fast DNA Stool Mini Kit (QIAGEN, Germany) per manufacturer instructions, with an additional mechanical lysis step using PowerBead tubes (QIAGEN, Germany). Extracted DNA was stored at -80°C until couriered on dry ice to the Australian Genomic Research Facility (AGRF) for sequencing.

2.8. Sequencing and Annotation

Sequencing of the 16S rRNA gene sequence was performed using the Illumina MiSeq platform. The V1–V3 hypervariable region of the 16S rRNA gene was amplified by polymerase chain reaction using 27F (AGAGTTTGATCMTGGCTCAG) and 519R (GWATTACCGCGGCKGCTG) primers with a read length of 300 base pairs. Diversity profiling analysis was performed with QIIME 2 2019.7³⁶. The demultiplexed raw reads were primer trimmed and quality filtered using the cutadapt plugin followed by denoising with DADA2 (via q2-dada2)³⁷. Taxonomy was assigned to amplicon sequence variants (ASVs) using the q2-feature-classifier³⁸ classify-sklearn naïve Bayes taxonomy classifier. Taxonomy was assigned using the SILVA (v.132) database.

2.9. Pre-Processing

Pre-processing and filtering was performed per Callahan et al³⁹. Zero count bacterial features and non-bacterial taxa were removed prior to calculating the alpha diversity metrics Shannon index and observed ASVs. Additional filtering removed low prevalence taxa (those present in less than 5% of samples), and data were centred-log ratio transformed to calculate Aitchison distances (i.e., beta-diversity) and for differential abundance testing. Functional potential was predicted using the Phylogenetic Investigation of Communities by Reconstruction of Unobserved States (PICRUSt2)⁴⁰, which maps to MetaCyc pathways.

2.10. Power Calculation

A sample size of 59 has more than 80% power to detect mean differences on paired before-after comparison in HADS score of 1.5 or greater, which has been considered clinically important⁴¹. We assumed a standard deviation of 3.7 for the differences⁴², and significance level of 0.05 using a two-sided paired t-test was used in the power calculation.

2.11. Statistical Analyses

Statistical analyses were performed within the RStudio⁴³ 4.3.1 environment using complete case analysis. Packages used for analysis are listed within the Supplementary Material. All univariate and multivariate models included age, sex, BMI, diet quality, and IBS at baseline as covariates, as these have previously been associated with depression. Multiple imputation with predictive mean matching was performed for missing covariate data, including age, sex, and BMI as auxiliary variables for precision.

The primary outcome was the change in average HADS depressive symptom scores one week pre- versus one-month post-procedure using complete case analysis. This was analysed using generalised estimating equations (GEE) assuming a Gaussian distribution with an AR(1) correlation structure to account for within-participant autocorrelation. Changes in average depressive symptom severity, anxiety symptoms, quality of life, perceived stress scores, Shannon index, observed ASVs, and the Shannon index of MetaCyc pathways were considered exploratory outcomes and measured with GEEs as above. Robust standard error estimates were reported for all GEE models. Beta-

diversity pre- and post-intervention was plotted using Principal Component Analysis (PCA) of Aitchison distances. Changes in beta-diversity metrics across time points were calculated using pairwise permutational ANOVA with 999 permutations, stratified by participant ID to consider the paired nature of the data. Differential abundance analyses at the genus level and of functional MetaCyc pathways were calculated using linear mixed models in the *Maaslin2* package, with minimum abundance and prevalence set to zero, time point and covariates as fixed effects, participant as the random effect, and a centred-log ratio transformation.

Exploratory linear regression models were used to examine potential associations between the changes in average HADS depressive symptom scores with change in bacterial genera and alpha diversity. A post-hoc sensitivity analysis was also performed, whereby all participants with a baseline sample were included in a modified intention-to-treat analysis with missing follow up data imputed using predictive mean matching, with age, sex, BMI, marital status, employment status, socioeconomic decile, diet quality, IBS, and bowel preparation adequacy used as auxiliary variables.

For differential abundance and exploratory analyses, the Benjamini-Hochberg procedure was applied to control the false discovery rate (FDR), with taxa below an FDR of 0.05 reported in results.

3. Results

3.1. Recruitment

Our study enrolled a total of 136 participants who provided informed consent at the time of their outpatient appointment. Eighty-six participants were successfully contacted, consented, and provided baseline data. Of these, five participants were excluded from analyses due to inadequate bowel preparation (rated as 'poor' by their endoscopist) and two participants were excluded because of a cancer diagnosis post-procedure. A further 20 participants were lost to follow-up between baseline and their one-month follow up. Therefore, 59 participants were included in complete case analyses, and 79 participants were included in modified intention-to-treat sensitivity analyses (Figure 1). In addition, two participants did not have intra-colonoscopy luminal samples collected, and three intra-colonoscopy samples were of too low biomass to yield sufficient DNA, leaving 56 luminal samples for analysis.

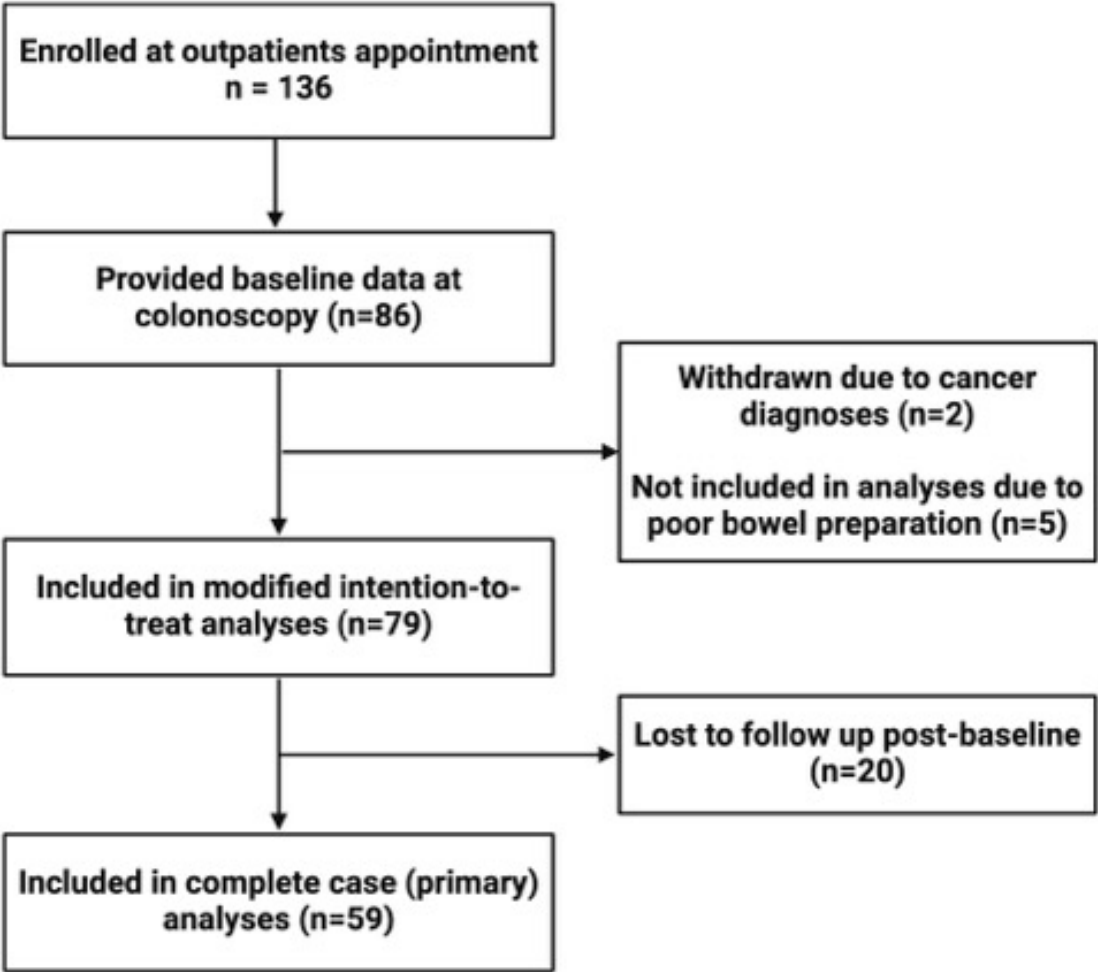


Figure 1. Participant enrolment and participation flow chart.

3.2. Participant Characteristics

Characteristics of this sample are presented in Table 1. Participants had a mean age of 58.5 years, almost equal distributions of sex (54% female) and socioeconomic advantage (56% IRSAD >5), and an average BMI of 29.7. Most of the study participants were taking at least one medication (92%), had poor average diet quality (46.8/100 SDQ points), with 12% current smokers, 22% meeting diagnostic criteria for IBS, and 22% self-reporting a lifetime history of depression. No participants self-reported antibiotic use within one month prior to colonoscopy. Procedural characteristics are presented in Table S1. Faecal samples were collected (on average) seven days pre-procedure, and 33 days post-procedure. The bowel preparation adequacy of most participants was rated as ‘good’ (but not excellent) by the endoscopists (63%), and polyps were the most common finding during colonoscopy (63%).

Table 1. Baseline characteristics of participants included in the present analyses (complete cases) and those lost to follow up.

	Complete cases (N=59)	Lost to follow up (N=20)	All cases (N=79)
Demographic			
Age, years, mean (SD)	58.5 (11.2)	56.7 (15.6)	58.0 (12.4)

	Complete cases (N=59)	Lost to follow up (N=20)	All cases (N=79)
Sex, female, n (%)	32 (54%)	13 (65%)	45 (57%)
IRSAD SES decile > 5, n (%)	33 (56%)	10 (50%)	43 (54%)
Health measures			
BMI, mean (SD)	29.7 (6.42)	27.8 (6.21)	29.2 (6.38)
Current smoking, yes, n (%)	7 (12%)	6 (30%)	13 (17%)
Diet quality, mean (SD)	44.8 (11.9)	30.2 (12.6)	41.7 (13.4)
Missing, n (%)	0 (0%)	4 (20.0%)	4 (5.1%)
Medical			
Self-report depression, yes n (%)	13 (22%)	6 (30%)	19 (24%)
Missing, n (%)	2 (3%)	4 (20%)	6 (7.6%)
IBS, yes n (%)	13 (22%)	6 (30%)	19 (24%)
IBS-constipation predominant	2 (15%)	2 (33%)	4 (21%)
IBS-diarrhoea predominant	6 (46%)	0 (0%)	6 (32%)
IBS-mixed predominant	5 (39%)	4 (67%)	9 (47%)
IBS-undefined predominant	0 (0%)	0 (0%)	0 (0%)
Missing, n (%)	1 (2%)	4 (20%)	5 (6%)
Medications			
Any medication, yes n (%)	54 (92%)	18 (90%)	72 (91%)
Antidepressants	11 (19%)	7 (35%)	18 (23%)
Hyperacidity/reflux medication	22 (37%)	6 (30%)	28 (35%)
Digestive supplements/cholelitholytics	2 (3%)	0 (0%)	2 (3%)
Mental health symptoms, mean (SD)			
HADS depression	3.71 (3.06)	5.45 (4.03)	4.15 (3.39)
HADS anxiety	5.37 (3.74)	7.20 (4.63)	5.84 (4.03)
HADS total	9.08 (6.04)	12.7 (7.98)	9.99 (6.71)
PHQ9 depression severity	3.61 (5.38)	7.20 (7.14)	4.52 (6.03)
Perceived stress	10.7 (7.67)	15.5 (7.14)	11.9 (7.77)
Quality of Life, mean (SD)			

	Complete cases (N=59)	Lost to follow up (N=20)	All cases (N=79)
Total	0.77 (0.19)	0.64 (0.24)	0.74 (0.21)
Psychosocial	0.48 (0.21)	0.36 (0.22)	0.45 (0.22)
Physical	0.71 (0.17)	0.64 (0.23)	0.69 (0.19)

Note: Age is age at time of recruitment; BMI calculated as weight (kg)/height(m)² at time of recruitment; Diet quality calculated from the Simple Dietary Questionnaire based on previous studies ³⁴. Abbreviations: BMI, body mass index; HADS, Hospital Anxiety Depression Scale; IBS, irritable bowel syndrome; PHQ, Patient Health Questionnaire; SD, standard deviation; SDQ, Simple Dietary Questionnaire.

3.3. Change in Mental Health Symptoms

There were decreases in average HADS depressive symptom scores (adjusted β =−0.64; 95%CI: −1.18, −0.11) and PHQ-9 depression severity scores (−0.68; −1.33, −0.02) one-month post-procedure versus baseline (Table 2). There were also increases in the average total (0.02; 0.01, 0.04), psychosocial (0.04; 0.02, 0.06), and physical (0.02; 0.00, 0.04) quality of life scores, but no statistical evidence of changed average HADS anxiety symptom scores or perceived stress scores (Table 2). Results of modified intention-to-treat sensitivity analyses (n=79) were similar; however, after imputation of missing data there was a statistically significant reduction in perceived stress scores (−1.86; −3.21, −0.51) (Table 2).

Table 2. Changes after bowel preparation and colonoscopy.

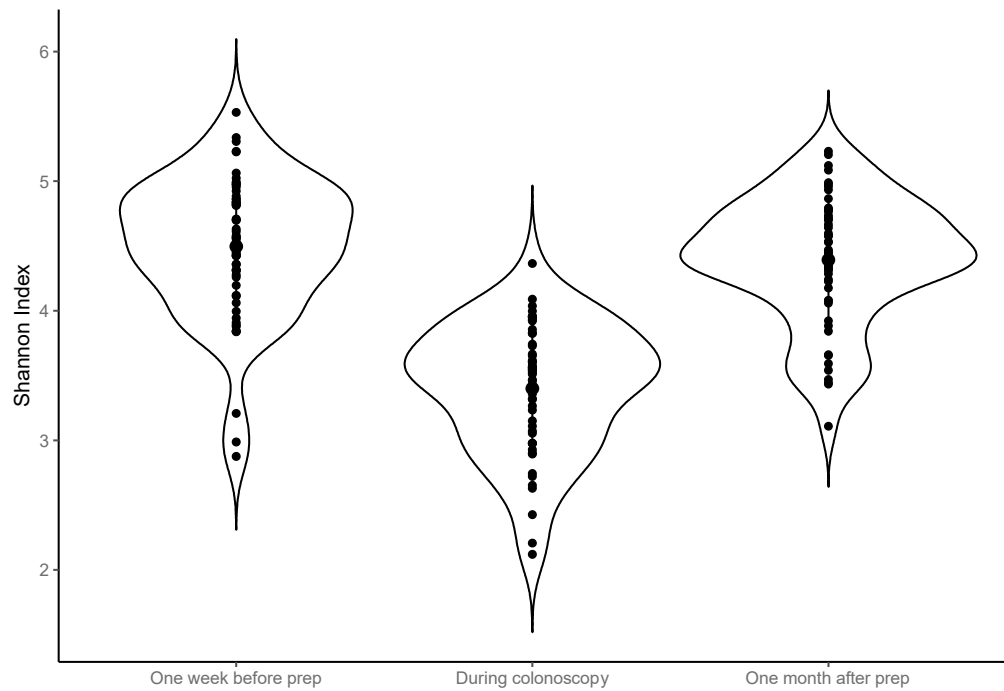
			Complete-case analysis (n=59)		Modified ITT
	Baseline Mean (SD)	Final Mean (SD)	Unadjusted β (95% CI)	Adjusted β (95% CI)	Unadjusted β (95% CI)
Depressive symptoms	3.71 (3.06)	3.07 (2.83)	−0.64 (−1.18 to −0.11)	−0.67 (−1.19 to −0.15)	−0.80 (−1.34 to −0.26)
Depressive symptom severity	3.61 (5.38)	2.98 (5.07)	−0.63 (−1.29 to 0.04)	−0.68 (−1.33 to −0.02)	−0.84 (−1.48 to −0.20)
Anxiety symptoms	5.37 (3.74)	5.27 (4.18)	−0.10 (−0.63 to 0.42)	−0.09 (−0.61 to 0.44)	−0.19 (−0.72 to 0.34)
Perceived stress	10.7 (7.67)	10.4 (7.30)	−0.32 (−1.53 to 0.89)	−0.39 (−1.61 to 0.82)	−0.57 (−1.78 to 0.64)
Total quality of life	0.77 (0.19)	0.80 (0.19)	0.02 (0.01 to 0.04)	0.03 (0.01 to 0.04)	0.03 (0.01 to 0.05)
Psychosocial quality of life	0.48 (0.21)	0.52 (0.22)	0.04 (0.02 to 0.06)	0.04 (0.02 to 0.06)	0.04 (0.02 to 0.06)
Physical quality of life	0.71 (0.17)	0.73 (0.18)	0.02 (0.00 to 0.04)	0.02 (0.00 to 0.05)	0.03 (0.01 to 0.05)
Observed ASVs	476 (143)	341 (98.8)	−135 (−168 to −102)	−137 (−170 to −105)	−123 (−156 to −90)
Shannon index	4.5 (0.54)	4.39 (0.48)	−0.10 (−0.23 to 0.02)	−0.10 (−0.23 to 0.03)	−0.04 (−0.17 to 0.09)
MetaCyc groups	5.17 (0.11)	5.16 (0.12)	−0.01 (−0.04 to 0.02)	−0.01 (−0.04 to 0.02)	−0.01 (−0.04 to 0.02)

^a Post-hoc sensitivity analysis including all participants with a baseline sample with missing follow-up data imputed using predictive mean model; ASVs, average sequence variants; BMI, body mass index; CI, confidence interval; ITT, intention-to-treat; SD, standard deviation.

3.4. Changes in Gut Microbiota Composition

Visual inspection of the data suggested that luminal samples (n=56) collected from colonic aspirates during colonoscopy – as expected – had lower average alpha diversity (i.e., observed ASVs and Shannon index) compared to faecal samples collected pre- and post-procedure (Figure 2).

A)



B)

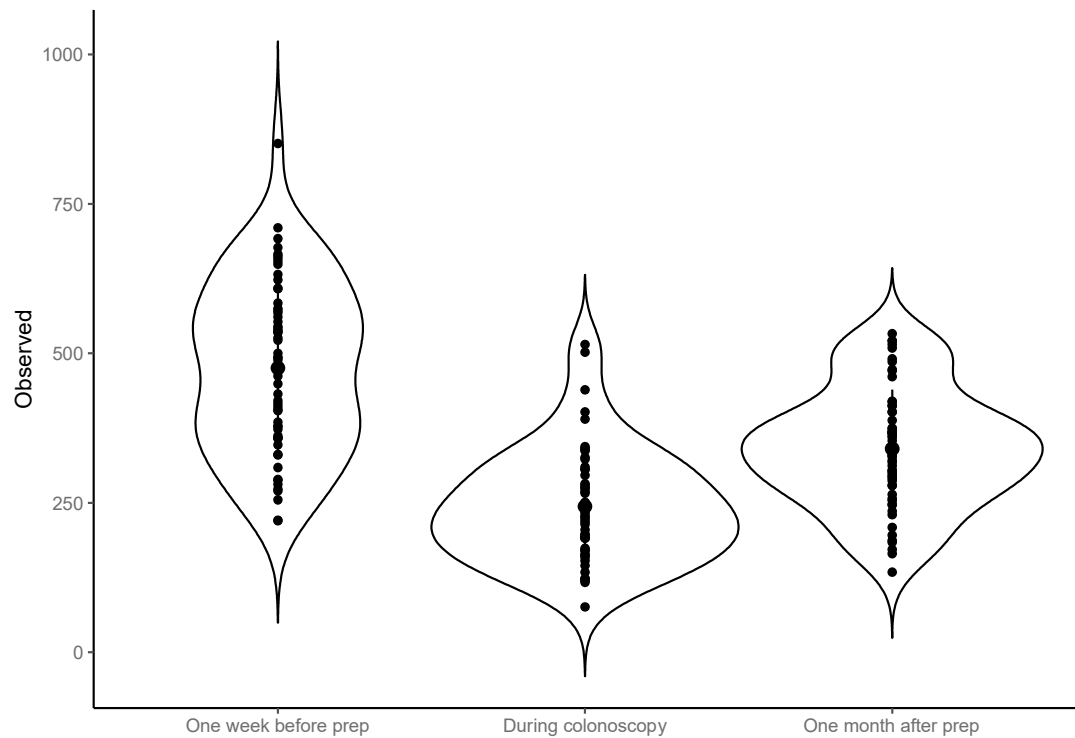


Figure 2. Differences in alpha-diversity metrics of A) Shannon index and B) Observed ASVs one week before bowel preparation and colonoscopy (n=59, orange), during colonoscopy (n=56, purple), and one month after bowel preparation and colonoscopy (n=59, blue).

As luminal samples are not directly comparable to faecal samples, statistical testing of alpha diversity metrics was performed using only the faecal samples collected one week pre- and one-month post-procedure. The average number of observed ASVs was decreased one-month post-procedure compared to baseline (−135; −168, −102), however there was no statistically significant change in the average Shannon index (Table 2). There was a modest change in overall community composition (i.e., beta diversity) one-month post-procedure ($p < 0.001$), however time point only explained 0.4% of the variance after adjusting for baseline covariates (Table S2).

There were 22 differentially abundant genera one-month post-procedure compared to baseline after adjusting for baseline covariates and multiple comparisons ($q < 0.05$) (Table S3). Of these, only an unidentified Lachnospiraceae genus was decreased one-month post-procedure compared to baseline, whereas the other 21 genera were increased post-procedure, and included the following annotated/identified taxa: Cutibacterium, Prevotella 9, Megamonas, Ruminococcaceae UCG-009, Oxalobacter, Lactonifactor, Ruminococcaceae CAG352, Lachnospiraceae GCA900066575, Eubacterium, Gordonibacter, Solobacterium, Rikenellaceae RC9 group, Lachnospiraceae UCG-010, and Megasphaera (Table S3).

There were no statistically significant changes in the average alpha diversity (i.e., Shannon index) of functional MetaCyc pathways (−0.01; −0.04, 0.02) (Table 2). Differential abundance analyses identified increases in ten MetaCyc pathways one-month post-procedure compared to baseline after adjustment for baseline covariates and multiple comparisons ($q < 0.05$), which included lipopolysaccharide lipid A biosynthesis and denitrification as the only annotated pathways (Table S4).

3.5. Exploratory Analyses

As 22% of the cohort met diagnostic criteria for IBS, exploratory analyses were conducted to examine whether IBS at baseline moderated the observed decrease in average depressive symptom scores post-procedure. We observed a statistically significant two-way interaction between IBS and time point ($\beta = 1.78$; 95%CI: 0.292, 3.26; $p = 0.019$). Those without IBS appeared to have reductions in depression symptom scores, whereas those with IBS appeared to have increased depression symptom scores, one-month post-procedure compared to baseline (Figure 3).

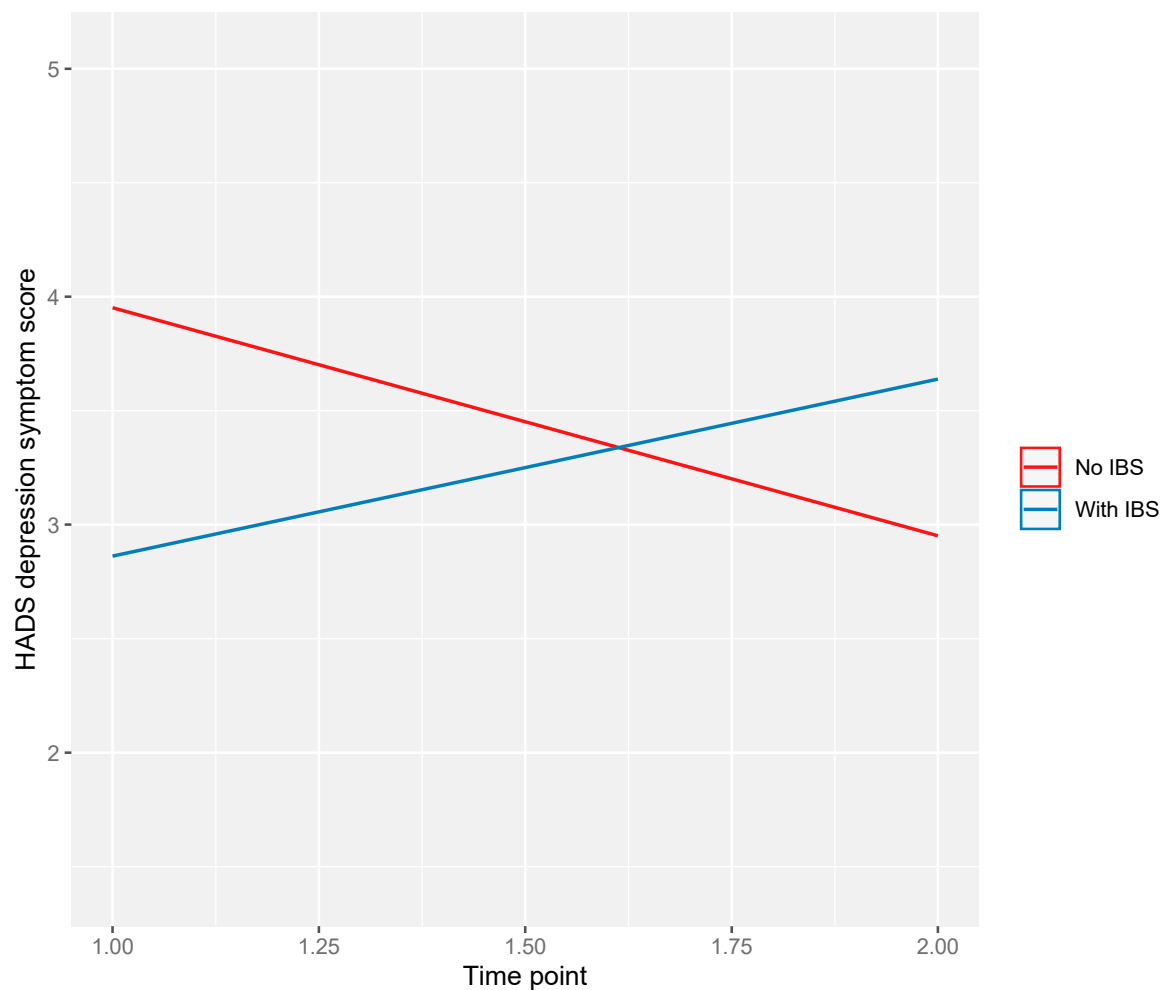


Figure 3. Two-way interaction between average HADS depressive symptom scores and irritable bowel syndrome at baseline. Red represents those without irritable bowel syndrome, blue represents those with irritable bowel syndrome. Shaded regions are confidence intervals for the estimated marginal means.

We also examined whether colonoscopy outcome moderated the observed decrease in average depressive symptom scores post-procedure. Due to our small sample size and some participants receiving more than one colonoscopy outcome (i.e., both polyps and diverticular disease), we dichotomized participants into two groups – those with (80%) and without abnormalities identified. We did not observe evidence of a statistically significant two-way interaction between colonoscopy outcome and time point ($\beta=-0.69$; 95%CI: $-2.11, 0.72$; $p=0.338$).

3.6. Post-Hoc Analyses

We also explored whether there were associations between the changes in gut microbiota composition and changes in average HADS depressive symptoms. There were 8 ASVs whose change was associated with changes in depressive symptoms after adjusting for covariates ($p<0.05$) (Table S5). Of those identified at the genus level, only the inverse association between *Ruminococcaceae* UCG-009 and the change in average HADS depression scores remained significant after adjusting for multiple comparisons ($q=0.028$) (Table S5).

4. Discussion

This is the first study, to our knowledge, to investigate the potential impact of colonoscopy and bowel preparation on depressive symptoms, and to associate changes in depressive symptoms with changes to gut microbiota composition. We observed a decrease in average depressive symptom scores, and increases in average quality of life scores, one-month post-procedure compared to baseline, but there was little evidence to support decreased average perceived stress or anxiety symptom scores. The changes in average depression symptom scores appeared to be moderated by IBS, whereby those with IBS experienced worsening of their depressive symptoms, and those without IBS experienced improvements. Compared to baseline, we also observed changes in gut microbiota composition, including reduced alpha-diversity, modest changes in beta-diversity, and genera and functional pathways that were differentially abundant one-month post-procedure. Finally, we observed that increases in the genus *Ruminococcaceae* UCG-009 were associated with decreases (i.e., improvements) in depressive symptom scores.

The finding that average depressive symptom scores decreased one-month post-procedure is novel and has not been previously explored. This finding is somewhat concordant with studies of other microbiome-modulating interventions that have reported improvements in depressive symptoms, such as probiotics⁹, faecal microbiota transplant^{44,45}, and diet^{6,8}. However, it is plausible that our participants experienced improvements in their depressive symptoms due to no major adverse findings (i.e., cancer) during colonoscopy, although we did not observe any changes in stress and anxiety measures, which might be expected if this were the explanation. Moreover, levels of depressive symptoms at baseline were low, and the average reduction observed, although statistically significant, was very small.

We observed a two-way interaction between time point and IBS status that suggested that those with IBS at baseline had worsening of their depressive symptoms post-procedure whereas those without IBS appeared to experience improvements. Interpretation of this finding is limited by the small number of participants with IBS within the study, however it is possible that the increase in depression symptoms in those with IBS could be due to an aggravation of their symptoms post-procedure. Worsening of gastrointestinal symptoms occurs in ~20% of patients after colonoscopy and bowel prep⁴⁶, and those with IBS are more likely to experience prolonged post-procedural abdominal pain⁴⁷. Worsening of depressive symptoms could also be due to a lack of findings to adequately explain their gastrointestinal symptoms. A previous study of IBS patients showed no improvement in reassurance or health-related quality of life in those that received a negative colonoscopy result⁴⁸. Additionally, a study observed that those with IBS were more likely to experience lower satisfaction, and higher burden and embarrassment, immediately after colonoscopy⁴⁹, and that these perceptions worsened six weeks compared to immediately post-colonoscopy⁴⁹. Whether IBS is an at-risk group that require additional disease management after colonoscopy may be an important focus of further research. Our findings suggest that the gut microbiota may still be re-establishing one month after bowel preparation and colonoscopy, however the health implications are unclear.

Interestingly, many bacterial genera that were higher in abundance one-month post-procedure compared to baseline have potential functionality that may be advantageous for health. We observed an increase in the relative abundance of bacteria with the ability to metabolise dietary polyphenols into metabolites that confer health benefits, including *Gordonibacter*^{50,51} and *Lactonifactor*⁵². A recent Mendelian Randomisation study identified low hippurate, which is produced by bacterial metabolism of polyphenols, as causally related to depression⁵³. There were also increases in bacteria with butyrate-producing capacity, including *Eubacterium*, *Solobacterium*, and *Megasphaera*; health benefits of short-chain fatty acids such as butyrate have been reviewed extensively⁵⁴. Thus, it could be postulated that the flushing out of gut bacteria by bowel preparation and colonoscopy may 'reset' the gut microbiota, resulting in a composition better able to metabolise foods and harness benefits from their diet.

We also observed that as *Ruminococcaceae* UCG009 increased, depressive symptoms decreased (i.e., improved). Our previous systematic review found that *Ruminococcaceae* was consistently lower in mental disorders compared to healthy controls⁴. More recently, a large association study found

Ruminococcaceae UCG002, UCG003, and UCG005 all to be negatively associated with depressive symptoms⁵⁵. The *Ruminococcaceae* family is largely considered beneficial to health, particularly due to the many butyrate-producing genera within this family including *Ruminococcus*, *Faecalibacterium*, *Caproiciproducens*, *Agathobaculum*, *Butyricoccus*, and *Gemmiger*. Future studies employing whole genome metagenomic sequencing with greater resolution at the species and strain level, or that specifically target changes in the abundance of *Ruminococcaceae* taxa, may afford further insight into how these bacteria change after intervention, and mechanistic studies such as reverse translation preclinical models may assist to elucidate how these bacteria influence host health and behaviours, including depressive symptoms.

Our study has notable limitations. We used 16S rRNA gene sequencing, which is subject to variability in sequencing depth and has only genus-level resolution. Our study did not collect any additional biological samples such as blood or urine to investigate changes to bacterial metabolites. Our study had a small sample size, and colonoscopy indications and outcomes were heterogeneous, which limited our statistical power and prevented subgroup investigations. Some of the changes observed may be associated with colonoscopy itself rather than bowel preparation. For example, the use of propofol, a sedative used for colonoscopy in the present study, has been previously found to have antidepressant potential⁵⁶. All participants in this study specifically used a sodium picosulfate-based bowel preparation product, and different results may be observed with alternative bowel preparation products. Those lost to follow-up had poorer mental health compared to those that completed the study, and the impact of their missing data, particularly for perceived stress, is unknown. Finally, we only collected comparable faecal samples at two time points, and greater sampling frequency within and beyond one month post-procedure may provide additional information regarding how the gut microbiota re-establishes.

Our study provides preliminary evidence of a potential impact of bowel preparation and colonoscopy on depressive symptoms that may relate to changes in the gut microbiota. Future research leveraging bowel preparation as a method of gut microbiota modulation may further our understanding of the microbiota-gut-brain axis. To better elucidate any potential mental health impact of bowel preparation itself, clinical trials of this intervention in a population experiencing heightened psychological distress and without colonoscopy are needed. Finally, clinical implications of the potential differential impact of bowel preparation and colonoscopy in those with and without IBS deserves further exploration.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org. Supplementary Table S1. Bowel preparation adequacy and colonoscopy indications and findings. Supplementary Table S2. Unadjusted and adjusted models of the change in beta-diversity of faecal samples collected one week before and one month after bowel preparation and colonoscopy using the Aitchison distance metric. Supplementary Table S3. Adjusted analyses of the change in differential abundance of taxa at the genus level. Taxa were adjusted for age, sex, body mass index, diet quality, and irritable bowel syndrome at baseline, and for multiple comparisons using Benjamini-Hochberg correction ($q < 0.05$). Supplementary Table S4. Adjusted analyses of the change in differential abundance of MetaCyc pathways. Analyses were further adjusted for age, sex, body mass index, diet quality, and irritable bowel syndrome at baseline, and for multiple comparisons using Benjamini-Hochberg correction ($q < 0.05$). Supplementary Table S5. Associations between the change in depressive symptom scores and change in the relative abundance of taxa at the genus level one month after bowel preparation and colonoscopy compared to baseline. Linear regression analyses were adjusted for age, sex, body mass index, diet quality, irritable bowel syndrome at baseline, and for multiple comparisons using Benjamini-Hochberg correction ($q < 0.05$).

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Conflicts of interest: MOH has a financial interest in Prevatech Pty Ltd, a company developing probiotic-based biotherapeutics. AL is a named inventor on a patent relating to Prevotella. FNJ has written two books for commercial publication. The remaining authors declare no conflicts of interest.

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