

1 **Characterizing Date Seed Polysaccharides: A Comprehensive Study on Extraction, Biological**
2 **Activities, Prebiotic Potential, Gut Microbiota Modulation, and Rheology Using Microwave-Assisted**
3 **Deep Eutectic Solvent**

4

5 **Athira Jayasree Subhash^a, Gafar Babatunde Bamigbade^a, Basel al-Ramadi^{b,c}, Afaf Kamal-**
6 **Eldin^a, Ren-You Gan^d, Chaminda Senaka Ranadheera^c, Mutamed Ayyash^{a,e*}**

7

8 ^a Department of Food Science, College of Agriculture and Veterinary Medicine, United Arab
9 Emirates University (UAEU), Al-Ain, UAE

10 ^b Department of Medical Microbiology and Immunology, College of Medicine and Health
11 Sciences, United Arab Emirates University (UAEU), Al-Ain, UAE

12 ^c Zayed Center for Health Sciences, United Arab Emirates University (UAEU), Al-Ain, UAE

13 ^d Singapore Institute of Food and Biotechnology Innovation (SIFBI), Agency for Science, Technology and
14 Research (A*STAR), 31 Biopolis Way, Singapore 138669, Singapore

15 ^e School of Agriculture, Food, and Ecosystem Sciences, Faculty of Science, The University of Melbourne,
16 VIC 3010, Australia

17

18 ***Corresponding author**

19 Mutamed Ayyash: mutamed.ayyash@uaeu.ac.ae

20

21

22

23

24

25

26 **Abbreviations**

27 PS: Polysaccharide; DES: Deep Eutectic Solvents; HBD: Hydrogen Bond Donors; SCFA: Short-
28 Chain Fatty Acids; DSP: Date Seed Polysaccharide; MPS: Microwave Extracted Polysaccharide;
29 CCD: Central Composite Design; TCA: Trichloroacetic Acid; GPC: Gel Permeation
30 Chromatography; TFA: Trifluoroacetic Acid; PMP: 1-phenyl-3-methyl-5-pyrazolone; FTIR:
31 Fourier Transform Infrared Spectroscopy; PDA: Photodiode Array Detector; DSC: Differential
32 Scanning Calorimetry; TGA: Thermo-gravimetric analysis; SEM: Scanning Electron
33 Microscope; WSI: Water Solubility Index; WHC: Water Holding Capacity; OHC: Oil Holding
34 Capacity; DPPH: 1-diphenyl-2-picrylhydrazyl; ABTS: 2,2'-asino-bis(3-ethylbenzene-thiazoline-
35 6-sulphonic acid; SD: Superoxide Dismutase; SAS: Superoxide Anion Scavenging; TAC: Total
36 Antioxidant Capacity; FRAP: Ferric Reducing Antioxidant Power; RP: Reducing Power; ACE:
37 Angiotensin-Converting Enzyme; FLASH: Fast Length Adjustment of Short; PDI: Polydispersity
38 index; RS: Reducing Sugar; TS: Total Sugar; GOS: Galacto-oligosaccharide; NMDS: Non-
39 metric multidimensional scaling; KEGG: Kyoto Encyclopedia of Genes and Genomes; COG:
40 Clusters of Orthologous Groups; PICRUSt: Phylogenetic Investigation of Communities by
41 Reconstruction of Unobserved States

42

43

44

45

46

47

48

49

50 **Abstract**

51 This study investigated the biological activities, prebiotic potentials, modulating gut microbiota,
52 and rheological properties of polysaccharides derived from date seeds via microwave-assisted
53 deep eutectic solvent systems. Averaged molecular weight (246.5 kDa) and a monosaccharide
54 profile (galacturonic acid: glucose: mannose: fructose: galactose), classifying MPS as a
55 heteropolysaccharide. MPS, at concentrations of 125-1000 $\mu\text{g}/\text{mL}$, demonstrates increasing free
56 radical scavenging activities (DPPH, ABTS, MC, SOD, SORS, and LO), potent antioxidant
57 potential (FRAP: 51.2-538.3 $\mu\text{g}/\text{mL}$; TAC: 28.3-683.4 $\mu\text{g}/\text{mL}$; RP: 18.5-171.2 $\mu\text{g}/\text{mL}$), and
58 dose-dependent antimicrobial activity against common foodborne pathogens. Partially-purified
59 MPS exhibits inhibition against α -glucosidase (79.6%), α -amylase (85.1%), and ACE (68.4%),
60 along with 80% and 46% inhibition against Caco-2 and MCF-7 cancer cells, respectively.
61 Results indicate that MPS fosters the growth of beneficial fecal microbiota, including
62 Proteobacteria, Firmicutes, and Actinobacteria, supporting microbes responsible for major
63 SCFAs (acetic, propionic, and butyric acids) production, such as Ruminococcus and Blautia.

64
65 Keywords: Date seeds; green extraction; polysaccharides; prebiotic; probiotic.

66

67

68

69

70

71

72

73

74

75 **1. Introduction**

76 *Phoenix dactylifera* L., commonly known as the date palm, is a crucially cultivated crop in
77 Middle Eastern countries for its fruits, with the date seed constituting 10-15% of the total weight
78 (Ghnimi, Umer, Karim, & Kamal-Eldin, 2017). Industries processing date-based products
79 contribute to an annual post-harvest loss of 852 thousand tons of date seeds (Faostat). Rich in
80 dietary fiber, tannins, proteins, and polyphenols, date seeds exhibit notable antioxidant and
81 antimicrobial potential, addressing conditions such as hypertension, coronary heart disease, high
82 cholesterol levels, and promoting gut microbiota enhancement (Al-Farsi & Lee, 2011). These
83 health benefits facilitate their incorporation into various food products, serving as emulsifiers,
84 stabilizers, and potential substitutes (Al-Khalili, Al-Habsi, & Rahman, 2023). Despite their
85 established functions, date pits are underutilized and often considered waste. Valorizing these
86 date seed wastes could provide a cost-effective source of various natural bioactive compounds.
87 Efforts should focus on utilizing the seeds in their natural form or extracting bioactive
88 components for specific applications.

89

90 Date seeds primarily consist of polysaccharides (approximately 80%), with xylans being the
91 main hemicellulose polymer (Noorbakhsh & Rabbani Khorasgani, 2023). Polysaccharides,
92 macromolecules composed of monosaccharide units linked via glycosidic bonds, have garnered
93 attention for their proven antioxidant, antimicrobial, anticancer, and prebiotic potential
94 (Romdhane et al., 2017). Polysaccharides in food positively influence gut microbiota, promoting
95 the growth of beneficial bacteria like *Bifidobacterium* and *Lactobacillus*, vital for digestion,
96 nutrient synthesis, and immune modulation (Koropatkin, Cameron, & Martens, 2012).
97 Traditional hot-water extraction methods suffer from limitations, leading to a shift towards
98 environmentally friendly approaches. Deep eutectic solvents (DESs), formed by mixing
99 quaternary ammonium salts with hydrogen bond donors, are recognized for their ease of

100 synthesis, non-toxicity, and economic feasibility. Coupled with non-thermal extraction methods,
101 microwave-assisted extraction using DESs has gained prominence (Wu et al., 2022). Microwave-
102 extracted polysaccharides exhibit improved extraction rates due to the disruption of
103 intramolecular hydrogen bonds (Dao, Webb, & Malherbe, 2021).

104
105 Understanding the health-promoting effects of polysaccharides necessitates grasping their
106 interaction with gut microbiota. Polysaccharides reach the colon undigested, fostering the growth
107 of beneficial bacteria and forming short-chain fatty acids (SCFAs), fulfilling the criteria for
108 prebiotics (Han et al., 2022). While conventional methods have been used to extract
109 polysaccharides from date seeds, this study explores the innovative non-thermal green extraction
110 technique of microwave-assisted deep eutectic solvent extraction for date seed polysaccharides
111 (MPS). The study aims to characterize MPS and assess its physicochemical, bioactive,
112 antioxidant, and rheological properties. Furthermore, it examines the fate of MPS during in vitro
113 digestion, evaluates its prebiotic potential on known probiotics, and analyzes its influence on the
114 gut microbiome using in vitro fecal fermentation.

115

116 **2. Materials and Methodology**

117 Whole date seeds (50 kg) were procured from the Al-Foah Date Processing Industry (Al Ain,
118 Abu Dhabi, United Arab Emirates) and stored at room temperature for further analysis. The DES
119 reagents (ChCl and EtGly) were purchased from Sigma-Aldrich Inc. (St. Louis, Missouri, USA).
120 All other chemicals and reagents used were of high-purity analytical grade and were also
121 obtained from Sigma-Aldrich unless otherwise indicated.

122 **2.1 Extraction of Polysaccharides from Date Seeds**

123 **2.1.1 Raw Material Preparation**

124 Date seeds were finely powdered using the Retsch ZM200 DR 100 miller (Germany) and
125 subsequently defatted via double hexane extraction following the method outlined by Yang,
126 Zamani, Liang and Chen (2021). The defatted date seeds were then stored for subsequent
127 extraction and analysis.

128 **2.1.2 Microwave-assisted Deep Eutectic Solvent Extraction of DSPs**

129 The microwave-assisted DES extraction of polysaccharides from date seeds was performed,
130 keeping the optimized DES conditions constant and varying the microwave factors, as per (Guo
131 et al., 2021). Date seed powder was combined with Choline chloride: ethylene glycol (1:1) DES
132 at a solid-to-liquid ratio of 1:10, undergoing primary extraction at 100 °C for 120 min. To
133 analyze the interaction between microwave factors and the extraction yield of polysaccharides, a
134 3-level central composite design (CCD) involving three factors—microwave power (250 W, 400
135 W, and 550 W), temperature (60 °C, 80 °C, and 100 °C), and time (5 min, 17.5 min, and 30
136 min)—was employed. The optimized microwave conditions were used for the extraction of MPS
137 from date seeds. Approximately 50g of defatted date seeds were subjected to microwave-assisted
138 DES extraction using Labotron 12T (Lyon, France). The resulting mixture underwent
139 centrifugation at -4 °C, and the residue was discarded. To eliminate the protein fractions, the
140 supernatant was mixed with 4% trichloroacetic acid (TCA) and stored for 1 h at 4 °C.
141 Subsequently, two volumes of absolute chilled ethanol were added to the supernatant, left for 24
142 h at 4 °C, and then subjected to centrifugation. The precipitate obtained was dissolved in warm
143 deionized water. The mixture was transferred to a 20 kDa MWCO Slide-A-Lyzer G2 dialysis
144 cassette (Thermo Fisher Scientific, USA) and dialyzed against deionized water for 72 h at 4 °C.
145 The yield of MPS was determined using the phenol-sulfuric acid method (Dubois, Gilles,
146 Hamilton, Rebers, & Smith, 1956). A portion of the partially-purified MPS was allocated for

147 bioactivity analysis, while the remainder was freeze-dried and stored at -20°C for further
148 analysis.

149 **2.2 MPS Characterization**

150 Molecular weight, monosaccharides composition, FTIR, zeta potential, particle size, and SEM of the
151 partially-purified MPS were carried out. The methods are detailed in the supplementary materials.

152 **2.3 Functional and Rheological Properties**

153 **2.3.1 Functional Properties of MPS**

154 Functional properties of MPS, specifically the water solubility index (WSI), water holding
155 capacity (WHC), and oil holding capacity (OHC), were determined according to the method
156 outlined by Jiang et al. (2021). The calculations were performed using the following equations:

$$157 \text{ WSI}\% = (\text{Dry solid weight}) / (\text{Total solid weight}) \times 100$$

$$158 \text{ WHC}\% = (\text{Water bound weight}) / (\text{Total solid weight}) \times 100$$

$$159 \text{ OHC}\% = (\text{Oil bound weight}) / (\text{Total solid weight}) \times 100$$

160 **2.3.2 Rheological Properties of UPS**

161 The rheological properties of MPS were assessed using a Rheometer (Discovery Hybrid HR-2,
162 TA Instruments, DE, USA) equipped with a cone plate, Peltier plate Steel (760 mm), in
163 alignment with the method detailed by (Ayyash et al., 2020). The amplitude sweep method was
164 utilized to determine the linear viscoelastic region of the samples at a constant frequency of 1.0
165 Hz, varying the strain between 0.1-10%. A frequency sweep test was conducted to assess the
166 viscoelastic behavior at frequencies ranging from 0.1 to 50 Hz, maintaining a constant strain of
167 0.5% within the linear viscoelastic region. The thixotropic behavior of the MPS sample was
168 evaluated using the time sweep method, recording the storage (G') and loss (G'') moduli at a
169 frequency of 1.0 Hz. Three-time segments were applied: 1) first (200 s, stress 0.8 Pa), 2) second
170 (100 s, stress 50 Pa), and third (400 s, stress 0.8 Pa). The data analysis was performed using
171 TRIOS 5.2 software.

172 **2.4 Biological Activities of MPS**

173 The biological activities methods, including the antioxidants activities (DPPH (%), ABTS (%),
174 TAC ($\mu\text{g/mL}$), RP ($\mu\text{g/mL}$), SAS (%), SOD (%), HRS (%), HP (%), FRAP ($\mu\text{g/mL}$), MC (%),
175 and LO (%)), ACE-inhibition (%), α -amylase and α -glucosidase inhibitions (%), minimum
176 inhibitory concentration (%) against 4 foodborne pathogens, and antiproliferative activities (%)
177 against Caco-2 and MCF-7 cancer cell lines were provided in the supplementary material.

178 **2.5 Digestability of MPS**

179 The *in vitro* digestion of the partially-purified MPS was performed as detailed in (Brodkorb et
180 al., 2019). The digesta was stored at $-20\text{ }^{\circ}\text{C}$ to further investigate the reducing sugar and total
181 sugar content before and after digestion based on the methods described in (Dubois et al., 1956;
182 Han et al., 2022).

183 **2.6 Prebiotic Properties of MPS**

184 The prebiotic effects of MPS on six probiotic strains (*Lactobacillus acidophilus* DSMZ 9126,
185 *Lactobacillus delbrueckii* subsp. *delbrueckii* DSMZ 20074, *Lacticaseibacillus rhamnosus* DSMZ
186 20021, *Lacticaseibacillus paracasei* subsp. *paracasei* DSMZ 20207, *Lactobacillus gasseri*
187 DSMZ 20243) were determined using the method outlined by Yılmaz and Şimşek (2020). The
188 growth kinetics of each probiotic strain with different carbon sources were measured at 600 nm
189 for 24 h at 15-min intervals.

190 **2.7 *In vitro* Fecal Fermentation and Gut Microbiota**

191 **2.7.1 *In vitro* Fecal Fermentation of MPS**

192 Fecal fermentation was performed by modifying the method of Han et al. (2022). Fecal slurry
193 was prepared by diluting freshly collected pooled feces with 10% freshly prepared phosphate
194 buffer solution. 2.5 mL of the slurry was combined with 20 mL of sterilized fermentation
195 medium with 1% MPS and 1% GOS-P (Control group), respectively. The fermentation medium

196 without an additional carbon source (the Blank group) served as the negative control group under
197 the same conditions. Incubation was carried out at 37 °C in a shaking water bath for 24 h.

198 **2.7.2 Fecal Fermentation Parameters**

199 The changes in pH, total sugar, gas production, and reducing sugar during fecal fermentation of
200 MPS were examined at 0, 6, 12, 24, and 48 h, following the procedure of Han et al. (2022).

201 **2.7.3 Short-Chain Fatty Acid Production**

202 After 48 h of fecal fermentation, the broth was centrifuged at 15000×g for 20 min, and the
203 supernatant was filtered using 0.45 µm filters. The Shimadzu HPLC system equipped with an
204 SPD-M20A photodiode array detector (PDA) and Shodex C18M 4E (250 × 4.6 mm, 5µm)
205 column was used. A calibration curve was created using different concentrations of acetic acid,
206 propionic acid, and butyric acid. The results were determined at a flow rate of 1.5 mL/min at a
207 column temperature of 30 °C with the UV detector set at 210 nm.

208 **2.7.4 Gut Microbiota Modulation**

209 The microbial composition of each group (NC, GOS-P, MPS) was analyzed at each time point (0,
210 6, 12, 24, and 48 h) during fecal fermentation. Genomic DNA was extracted using a Genomic
211 DNA Kit (Tiangen, Beijing, China), and V3-V4 regions of 16S rRNA were amplified and
212 analyzed by BGI, Hong Kong. Library construction, concentration, and quality assessment were
213 performed using Agencourt AMPure XP beads and Agilent 2100 Bioanalyzer, respectively. The
214 raw data were filtered using iTools Fqtools fqcheck (v.0.25), and the paired-end reads were
215 merged into a single tag sequence using Fast Length Adjustment of SHort reads (FLASH,
216 v1.2.11). The sequences were clustered into operational taxonomic units (OTUs) with a 97%
217 similarity threshold by UPARSE, and chimeras were filtered using UCHIME (v4.2.40). The
218 OTU representative sequences were mapped to the tags using USEARCH (v7.0.1090) and
219 aligned against the database for taxonomic annotation using the RDP classifier (v2.2) at 60%
220 sequence identity. Alpha and beta diversity were calculated using mothur (v.1.31.2) and QIIME

221 (v1.80). Differential species analysis was carried out using Linear Discriminant Analysis Effect
222 Size (LEfSe) (<https://huttenhower.sph.harvard.edu/galaxy/>). Microbial functional annotation was
223 predicted by PICRUSt2 v2.3.0-b, and correlation analysis and model prediction were performed
224 using R (v3.4.1) and Cytoscape.

225 **2.8 Statistical Analyses**

226 All analyses were conducted in triplicates. The activities of partially-purified MPS at
227 concentrations ranging from 125 to 1000 $\mu\text{g/mL}$ were utilized for statistical comparison. One-
228 way and Two ANOVA tests were employed as deemed necessary, utilizing JMP (SAS Institute,
229 NC, USA). Tukey's test was applied to compare differences between samples with a significance
230 level of $P < 0.05$. Data analysis from HPLC was performed using Shimadzu LabSolutions
231 software (Japan). All other softwares used are appropriately cited in the context wherever
232 necessary.

233 **3. Results**

234 **3.1 Extraction and Purification from Date Seeds**

235 Efficient microwave-assisted extraction of date seed polysaccharides using DES was ensured
236 through process optimization using central composite design (CCD) based on RSM. A 16-run
237 CCD with three independent variables—microwave power (W), time (min), and temperature
238 ($^{\circ}\text{C}$)—studied at three levels, revealed the measured response as the yield of MPS. As per Table
239 S1, the overall MPS yield ranged from 15.1 to 41.2 mg/g. Experimental runs at the center points
240 (#3 and #5) demonstrated minimal variation in MPS yield, indicating good repeatability. Higher
241 extraction temperature and power over longer extraction times decreased MPS yield. The
242 maximum MPS yield (42.1 mg/g) was observed at a microwave power of 250 Hz, a temperature
243 of 60°C , and a time of 5 min. These optimized conditions were employed for MPS extraction,
244 and optimization surface plots are detailed in Fig. S1. The obtained MPS, sparingly soluble in
245 water, appeared as solids dispersed in a liquid medium, potentially due to structural

246 characteristics promoting intermolecular association, such as linear chain arrangement and
247 higher molecular weight.

248 **3.2 Characterization of MPS**

249 Purified, freeze-dried MPS underwent evaluation for molecular weight and monosaccharide
250 profiling as specified in section 2.2.1. The Mw of MPS was distributed in the range of 115 to
251 246.45 kDa (Fig. S2 A and B). The chromatogram of polysaccharides depicted 4 sharp peaks,
252 suggesting the presence of 4 polysaccharide fractions with varied concentrations and molecular
253 weights (Fig. S3B). The major constituents were peaks 1 and 4, constituting 94.12% of the total
254 polysaccharide. The polydispersity index (PDI) of MPS peak 1, calculated based on the ratio of
255 weight average molecular weight (Mw) and average molecular weight (Mn), was 2.84 and 2.09,
256 respectively, categorizing MPS as a widely dispersed polymer (greater than 2).

257 Monosaccharide composition analysis (Fig. S3B) illustrated galacturonic acid and glucose as the
258 primary monosaccharides of MPS, verified by comparing their chromatograms with the PMP-
259 derivatized standard monosaccharide (Fig. S3A). Additionally, mannose, fructose, and galactose
260 were present. The structural attributes of natural polysaccharides can be inferred from their FTIR
261 spectra. The FTIR spectra of MPS (Fig. 1A) revealed broad peaks at 1047, 1439, 1522, 1623,
262 2900 and 3277 cm^{-1} . Weak absorption peaks were also observed within the range of 554-846 cm^{-1}
263 and 1179-1326 cm^{-1} .

264 The thermal properties of the polysaccharide, specifically concerning decomposition temperature
265 and heat absorption, are demonstrated through a DSC curve, showcasing the relationship
266 between temperature ($^{\circ}\text{C}$) and heat flow (W/g) (Fig. 1B). This plot visualizes the changes-
267 exothermic or endothermic-experienced by the MPS as the temperature varies from 0 to 250 $^{\circ}\text{C}$.
268 The DSC thermogram of MPS distinctly exhibits an exothermic phase in the range of 25 to 150
269 $^{\circ}\text{C}$, indicating the sample's thermal decomposition. Notably, the absence of a single endothermic
270 peak in the thermogram indicates the lack of impurities and dehydration in the MPS.

271 Furthermore, the thermal stability of the MPS is elucidated in terms of thermal decomposition
272 temperature, thermally stable temperature, and sample composition, represented through the
273 Thermogravimetric (TG) curve (Fig. 1C). This curve portrays a one-stage weight loss
274 corresponding to the evaporation of water around 50-200 °C. It is observed that the MPS does
275 not decompose before 210 °C, demonstrating a gradual weight loss of 1.5% at 110 °C, followed
276 by a rapid 6% weight loss at 600 °C. Additionally, the particle size and zeta potential of MPS
277 were determined as 5082.5 nm and -391.34mV, respectively.

278 In the morphological and surface characterization through SEM analysis (Fig. 2 A-D), freeze-
279 dried MPS at different magnifications shows a regular, porous surface with defined geometry and
280 irregular pores. This feature signifies a larger surface area due to structural changes induced by
281 microwaves. For hydration and foaming properties, MPS demonstrated an average water holding
282 capacity (WHC) of 3.378 ± 0.068 g/g, oil holding capacity (OHC) of 5.79 ± 0.0155 g/g, and water
283 solubility index (WSI) of $80.9 \pm 0.021\%$.

284 Regarding viscoelastic properties, evaluation through amplitude, frequency, and time sweep tests
285 revealed that as the strain varied from 0.1% to 10%, G' decreased from 10^{-4} MPa to 10^{-5} MPa,
286 while G'' remained constant at 10^{-5} MPa. This indicated the dominance of elastic behavior over
287 viscous behavior in MPS. Frequency sweep test data also showed that G' remained higher than
288 G'' , highlighting the elastic properties of MPS. The complex viscosity decreased from 10^{-3} to 10^{-5}
289 as the frequency increased from 0.1 to 20 Hz. The time sweep test, at 0.8% strain, showed a
290 convergence between G' and G'' , while at 50% strain, G'' surpassed G' , suggesting a transition
291 from elasticity to a more viscous state in MPS.

292 **3.3 Bioactivities of MPS**

293 To assess the overall bioactivity of the polysaccharide, various activities such as antioxidant,
294 antimicrobial, anticancer, ACE inhibition, and antidiabetic potential were evaluated for MPS.
295 Radical scavenging assays were conducted across a concentration range of 125-1000 $\mu\text{g/mL}$

296 (Table 1). The DPPH and ABTS radical scavenging activities increased from 17.9 to 60.8% and
297 11.7 to 67.8%, respectively, with the increase in MPS concentration. The superoxide dismutase
298 and superoxide anion scavenging activities also exhibited increments from 20 to 57.4% and 18.8
299 to 49.8%, respectively, as the MPS concentration rose. Similarly, hydrogen peroxide and
300 hydroxyl radical scavenging activities increased in a dose-dependent manner, reaching 68.8%
301 and 64.6% at the highest concentration of 1000 $\mu\text{g}/\text{mL}$. MPS showed notable antioxidant
302 properties, preventing lipid oxidation by 60% and exhibiting a metal chelation activity of 6343.8
303 $\mu\text{g}/\text{mL}$. Comparing MPS to standard ascorbic acid (1 mg/mL), assays like ferric ion-reducing
304 antioxidant power, Total antioxidant, and reducing power displayed a dose-dependent increase in
305 antioxidant potential. At 1000 $\mu\text{g}/\text{mL}$, the antioxidant potential measured 538.3, 683.4, and 171.2
306 $\mu\text{g}/\text{mL}$, respectively, for FRAP, TAC, and RP. Our results affirm that the antioxidant properties of
307 MPS were dose-dependent and amplified with concentration ($P < 0.05$).

308 The effects of purified UPS on α -glucosidase, α -amylase, and ACE inhibition are depicted in Fig.
309 3A. The inhibition of both enzymes showed a significant increase ($P < 0.05$) with rising MPS
310 concentrations from 125-1000 $\mu\text{g}/\text{mL}$. The maximum inhibition was observed at 1000 $\mu\text{g}/\text{mL}$,
311 with MPS inhibiting α -glucosidase and α -amylase by 79.6% and 85.1%, respectively. At 1000
312 $\mu\text{g}/\text{mL}$, MPS exhibited an ACE inhibition of 68.4%, which was notably lower than the enzyme
313 inhibitions.

314 The evaluation of the anticancer activity of MPS on Caco-2 (colorectal adenocarcinoma) and
315 MCF-7 (breast cancer) cell lines was conducted to ascertain potential anticancer properties.
316 While MPS displayed significant dose-dependent increasing inhibition towards Caco-2 cell lines,
317 inhibition against MCF-7 cancer cell lines (% inhibition) decreased with increasing MPS
318 concentration (Fig. 3A). At 1000 $\mu\text{g}/\text{mL}$, MPS only exhibited 28% inhibition against MCF-7
319 cancer cell lines with an IC_{50} value of 87.6 ($\mu\text{g}/\text{mL}$), whereas 80% inhibition was reported against
320 Caco-2 cell lines with an IC_{50} value of 58.9 ($\mu\text{g}/\text{mL}$) (Fig. 3A).

321 Moreover, the antimicrobial potential of MPS against common foodborne pathogens
322 demonstrated a dose-dependent increase in inhibition (Fig. 3B). At the highest concentration of
323 4.71 mg/ml, MPS displayed significantly higher inhibition percentages: *E. coli* 0157:H7 1934
324 (60%), *S. aureus* ATCC 25923 (66%), *S. Typhimurium* 02-8423 (86%), and *L. monocytogenes*
325 DSM 20649 (71%) (Fig. 3B). The MIC of MPS for the inhibition of all four pathogens was 0.588
326 mg/mL. The robust antimicrobial properties of MPS broaden its applications in functional foods
327 and food preservation.

328 **3.4 *In vitro* Digestion and Prebiotic Potentials**

329 *In vitro* digestion studies revealed decreased total sugar (TS) content from 32.7 to 16.89 mg/mL,
330 and increased reducing sugar (RS) content from 1 to 7.96 mg/mL with digestion (Fig. S4). These
331 results indicate that some parts of MPS might undergo digestion in the digestive tract before
332 reaching the colon for interaction with microflora. Further studies involving the determination of
333 relative molecular weights of MPS fractions during each stage of digestion are recommended for
334 deeper insights.

335 The prebiotic effects of MPS on 6 probiotic strains were investigated by measuring their growth
336 kinetics using MPS as a carbon source (Fig. S5). All strains of probiotics utilized MPS as a
337 carbon source, similar to glucose, indicating their prebiotic potential. However, MPS did not
338 significantly enhance the exponential growth of *L. rahnmosus* and *L. gasseri* compared to
339 glucose. The lag phase of the growth of probiotics when using MPS and glucose was almost
340 identical. Additionally, the optical densities of the media were higher for the growth of all 6
341 strains utilizing MPS than glucose, suggesting comparably higher growth when using MPS as a
342 carbon source. For all cases, the 24-hour period did not indicate the onset of deceleration for
343 microbial strains using MPS and glucose, but did for the strains using GOS-P and no carbon
344 source. This emphasizes the comparable prebiotic potential of glucose and our polysaccharide.

345 **3.5 *In vitro* Fecal Fermentation of MPS**

346 **3.5.1 Fecal Fermentation Broth Properties**

347 The impact of UPS on *in vitro* fecal fermentation broth was assessed by evaluating changes in
348 pH, gas production, total and reducing sugars, as well as the production of SCFAs (acetic,
349 propionic, and butyric acid) (Fig. 4 A-G). Galacto-oligosaccharide (GOS-P) and the blank
350 without a carbon source were used as positive and negative controls (NC), respectively.

351 The pH values (Fig. 2A) for GOS-P initially decreased significantly from 6.7 to 6.2 after 12 h of
352 fermentation. In contrast, NC exhibited an unstable trend in pH, ultimately settling at 6.6,
353 indicating limited fermentation by gut microbiota. MPS demonstrated a sharp decline in pH
354 within the initial 6 h but remained relatively constant until the 24-h mark. At 24 h, MPS
355 displayed the lowest pH, which later increased by 48 h. Throughout the fermentation period,
356 MPS did not exhibit a significant trend ($p < 0.0001$) in pH alteration.

357 Gas production (Fig. 4B) for GOS-P peaked, increasing from 2.5 to 5.5 ml during fermentation.
358 MPS, until 24 h, significantly increased gas production ($P < 0.0001$) compared to NC (2 mL).
359 However, after 24 h, gas production from MPS decreased, suggesting maximal utilization by the
360 gut microbiota.

361 Observations indicated a significant reduction in total sugar content for all samples at 6 h, which
362 remained constant throughout the fermentation period. From 6 to 48 h, GOS-P exhibited the least
363 total sugar content at 4.650 mg/mL, followed by MPS and NC, respectively (Fig 4C). The
364 reducing sugar content of GOS-P notably dropped after 6 h of fermentation, while that of NC and
365 MPS remained consistent until 24 h. A slight increase in RC content for GOS-P and MPS was
366 observed after 48 h of fermentation (Fig. 4D).

367 The decline in pH, gas production, and total sugar content of the fecal broth until 24 h, coupled
368 with the increase in reducing sugar content after 48 h, suggests the effective utilization of MPS
369 by gut microbiota. This is further supported by the production of SCFAs in the broth. Acetic acid,

370 propionic acid, and butyric acid, the primary SCFAs produced, are depicted in Fig. 4 (E-G). NC
371 in the fecal broth demonstrated the highest acetic acid production at the conclusion of 48 h of
372 fermentation (1923 $\mu\text{mol/g}$), followed by GOS-P (1609 $\mu\text{mol/g}$) and MPS (1476 $\mu\text{mol/g}$),
373 respectively (Fig 4 (E)). NC and GOS-P exhibited consistently lower butyric acid production
374 compared to MPS. After 24 h, MPS displayed the highest butyric acid production (1119 $\mu\text{mol/g}$),
375 which declined to 488 $\mu\text{mol/g}$ after 48 h of fermentation (Fig 4(G)). The propionic acid
376 production of NC (1362 $\mu\text{mol/g}$) and GOS-P (1359 $\mu\text{mol/g}$) was almost similar and higher
377 compared to MPS (1220 $\mu\text{mol/g}$) throughout the fermentation period (Fig 4(F)). These results
378 collectively indicate that MPS underwent breakdown in the colon, releasing significant amounts
379 of all three major SCFAs comparable to the standard polysaccharide (GOS-P).

380 **3.5.2 Gut Microbiota and Functional Profile**

381 The impact of partially-purified MPS on the gut microbiota and their functional profile was
382 examined. Functional analysis holds significant biological importance in the analysis of
383 microbial diversity, as it allows the linkage of species and their functions to derive general
384 distribution profiles of community function. A Venn diagram (Fig. 5A) was employed to
385 ascertain the number of OTUs (shared and unique) in the three groups: NC, GOS-P, and MPS.
386 Results indicated 519 OTUs shared among all groups, with 88 OTUs shared between NC and
387 GOS-P, 70 between GOS-P and MPS, and 62 between NC and MPS. MPS exhibited significantly
388 fewer shared OTUs with both NC and GOS-P. Furthermore, MPS showcased 64 unique OTUs,
389 while NC and GOS-P displayed 53 and 36 unique OTUs, respectively, signifying that MPS
390 contributed to an increased variety of gut microflora in comparison to both positive and negative
391 controls.

392 The alpha and beta diversity analyses of the three groups during fermentation are depicted in Fig.
393 5B and Fig. S6. The rarefaction curves (Fig. S6(A-C)) illustrated rich and diverse gut microbiota
394 in all three groups, with MPS and NC presenting better diversity patterns than GOS-P in Fig.

395 S6(D-F). Significant differences ($P < 0.05$) in diversity among groups were observed only in
396 Shannon's and Simpson diversity, as indicated by alpha diversity box plots (Fig. 5B). In all plots,
397 GOS-P exhibited the lowest diversity, followed by MPS and NC. Put differently, GOS-P and
398 MPS clustered away from NC throughout the fermentation period.

399 Subsequently, beta diversity analysis was conducted using NMDS (non-metric multidimensional
400 scaling), UPGMA cluster tree, and order abundance bar plot. The NMDS plot displayed GOS-P
401 and MPS within the same region, while NC separated from them throughout the fermentation
402 period (Fig. 5C). This suggests that GOS-P and MPS might share some common microbial load
403 distinct from NC. PCA, crucial in β -diversity studies, revealed that the microbiota shifted in both
404 MPS and GOS-P. PC1 and PC2 contributed 58.8% and 25.76%, respectively (Fig. 5D),
405 demonstrating that the gut microbiota favored by MPS and GOS-P significantly differed from
406 NC.

407 Through Bray-Curtis and relative abundance studies (Fig. 5E), among order-based
408 categorization, Enterobacterales were notably abundant in MPS throughout fermentation.
409 Simultaneously, Clostridiales, Bifidobacterales, and Lactobacillales were present in all groups.
410 Enterobacterales were predominant in MPS during the fermentation period up to 24 h, replaced
411 by Bifidobacterales after 48 h. Subsequent function prediction studies are necessary to discern if
412 specific bacteria from this phylum were promoted by MPS. The species phylogenetic tree
413 identified the presence of Firmicutes, followed by Proteobacteria, Bacteroidetes, and
414 Actinobacteria (Fig. 5F), as the major species of the phylum present, in alignment with the beta
415 diversity results.

416 The functional gene prediction of gut microbiota based on the Kyoto Encyclopedia of Genes and
417 Genomes (KEGG), Clusters of Orthologous Groups (COG), and MetaCyc metabolic pathways
418 using PICRUSt2 (Phylogenetic Investigation of Communities by Reconstruction of Unobserved
419 States) was executed. The KEGG data illustrated the possible metabolic pathways and their

420 relative likelihood during fecal fermentation. At KEGG level 1 (Fig. 6A), metabolism emerged
421 as the predominant microbial functionality across all three groups. Level 2 of KEGG (Fig. 6B)
422 highlighted that the microbial load predominantly engaged in the metabolism of cofactors and
423 vitamins, carbohydrates, major and other amino acids, terpenoids and polyketides, and energy,
424 aligning with the KEGG level 1 outcomes. KEGG level 3 (Fig. 6C) demonstrated that
425 biosynthetic mechanisms including ansamycin, riboflavin, vancomycin group antibiotics,
426 peptidoglycan, and photosynthesis were the prominent functionalities of the gut microflora
427 influenced by MPS.

428 The COG heatmap was introduced to further elucidate the KEGG pathway levels (Fig. 6D).
429 Analysis revealed that carbohydrates and amino acids transport and metabolism, translation
430 ribosome structure, biogenesis, transcription, and general function predictions were the principal
431 microbial functionalities, followed by minor functions such as cell wall/membrane biogenesis
432 and energy production and conversion. These functions were consistent for both GOS-P and
433 MPS, displaying similar intensities. The MetaCyc heatmap (Fig. 6E) indicated amino acid
434 biosynthesis and biosynthesis of cofactors, prosthetic groups, electron carriers, and vitamins,
435 appearing abundant in MPS and GOS-P.

436 Species Spearman correlation analysis and network analysis hinted at the associations among
437 different species of fecal microbiota in the three groups—NC, GOS-P, and MPS. Fig. 6F outlined
438 the pairwise correlation coefficients among the species during fecal fermentation. Strong positive
439 correlations ($p < 0.05$) were observed between *Bifidobacterium longum* and *Blautia wexlerae*,
440 and significant positive correlations ($p < 0.05$) with Gemmiger. Formicilis and *Romboutsia*
441 *sedimentarium*, while *Blautia luti* demonstrated robust correlation with *Blautia schinkii* and
442 Gemmiger. Formicilis. Conversely, *Escherichia* and *Enterococcus saccharolyticus* exhibited
443 negative correlations with almost all present microflora. Additionally, network analysis (Fig.

444 S7A) highlighted strong positive correlations ($p < 0.05$) between *Enterococcus saccharolyticus*
445 in the fecal flora and all other gut microbiota.

446 Analyzing the relative abundance heatmaps of phylum and species (Fig. S7 (B, C)) revealed that
447 Firmicutes were abundant in fecal samples with NC and GOS-P, whereas Proteobacteria were
448 present in samples containing MPS. Both Firmicutes and Actinobacteria were also found in
449 MPS-incorporated fecal slurry but at comparatively lower levels. After 48 h of fermentation,
450 Actinobacteria became dominant in MPS-containing slurry, impeding the growth of
451 Proteobacteria. The presence of Bacteroidetes was negligible in the relative abundance heatmap.

452 **4. Discussions**

453 **4.1 Extraction and Characterization of MPS**

454 Utilizing microwaves in conjunction with green solvents in a non-thermal extraction process
455 enables direct heat release within the solvent system through ionic conduction and dipole
456 rotation, inducing a rapid temperature rise (Bagade & Patil, 2021). This phenomenon leads to the
457 disruption of weak hydrogen bonds within the samples, thereby enhancing extraction efficiency.
458 Our study observed a notably higher MPS yield, specifically 42.135 mg/g, which can be
459 attributed to the increased solubility of sugars in DES. The liberated microwave energy
460 facilitates the disintegration of cell walls, enabling solvents to penetrate more effectively, thus
461 enhancing the extraction yield (Chen et al., 2022).

462 Moreover, microwave irradiation has the potential to enhance the polarity of DES, thereby
463 maximizing their ionic nature. Previous studies on date seed polysaccharides have reported
464 extraction yields of 17.54 ± 0.07 g/100 g through ultrasonic-assisted alkaline hot water extraction
465 (Noorbakhsh & Rabbani Khorasgani, 2023), and 1.03% with ultrasound-assisted hot water
466 extraction (Dhahri et al., 2023). Notably, increased microwave temperature and duration can lead
467 to PS degradation, consequently affecting the extraction yield. Microwave-assisted deep eutectic
468 treatment emerges as an excellent alternative to conventional extraction techniques for the

469 successful extraction of date seed polysaccharides by breaking noncovalent intra- and inter-
470 molecular bonds, releasing water-soluble polysaccharide molecules (Chen, Zhang, Huang, Fu, &
471 Liu, 2017).

472 The lower molecular weight of MPS fractions could result from the synergistic effect of
473 microwave irradiation and targeted specific DES extraction. Similar results were reported for PS
474 extracted from sweet tea leaves by Guo et al. (2021). DES has a more significant effect on
475 breaking down the complex structure and intra- and intermolecular interactions of
476 polysaccharides compared to water at the same extraction temperature. This results in a relatively
477 higher degree of MPS degradation, breaking down macromolecules into smaller molecules.

478 Typically, most naturally occurring polysaccharides exhibit high PDI values. The MAE method
479 facilitates the production of a larger quantity of low Mw fractions by degrading polysaccharides,
480 converting high-Mw fractions into corresponding low-Mw sub-fractions, consistent with findings
481 reported by Ren et al. (2017). The presence of galacturonic acid in appreciable amounts was also
482 noted by Manhivi, Venter, Amonsou and Kudanga (2018) in cactus polysaccharides, suggesting a
483 pectic polysaccharide.

484 The composition of monosaccharides varies concerning processing methods, conditions, sample
485 types, and varieties. Polysaccharides are often characterized by their distinctive absorption peaks.
486 The α -glycosidic linkage can be identified by peaks at 846 cm^{-1} . In MPS, the observed broad
487 peaks around 1047 cm^{-1} and 3277 cm^{-1} indicate C-O stretching due to cellulose and O-H
488 stretching owing to cellulose and water molecules. Also, the characteristic band at approximately
489 2900 cm^{-1} aligned with the C-H stretching vibration previously described by (Dhahri et al.,
490 2023). Additionally, prominent peaks at 1439 and 1623 cm^{-1} denote C=C skeletal vibrations of
491 lignin, hemicellulose, and C-O stretching of cellulose. Bands at 1326 cm^{-1} could be linked to the
492 O-C=O bending of uronic acids present in date seeds. Ajwa date seed polysaccharide also
493 displayed similar FTIR spectra (Dhahri et al., 2023). The significantly higher water solubility

494 index of MPS aligns with the notion that microwave treatment can enhance the solubility of
495 polysaccharides (Chen et al., 2022).

496 At higher DSC temperatures, high molecular weight polysaccharides tend to degrade, forming
497 low molecular weight oligo and monosaccharides. In the case of MPS, exothermic phases
498 observed in the range of 25 to 150 °C indicated the thermal decomposition of the sample.
499 Polysaccharides typically manifest endothermic peaks when their structural backbone melts .
500 Therefore, the absence of such endothermic peaks suggests that MPS might exhibit higher
501 thermal stability compared to polysaccharides extracted conventionally and via certain non-
502 thermal methods (Chen et al., 2022).

503 The TG curve of MPS depicted a singular stage of weight loss, aligning with the loss of water
504 occurring around 50-200 °C. In this curve, the intra and intermolecular polymer hydroxyls
505 present in polysaccharides underwent condensation, releasing water molecules. The slower
506 weight loss rate observed in MPS implies its superior thermal stability compared to other
507 polysaccharides (Wang et al., 2022).

508 Visual analysis through surface images revealed that MPS exhibited a regular, loose surface with
509 irregular pores and significant aggregation. Similar findings have been observed in the SEM
510 images of microwave/ultrasound-extracted PS of *Panax notoginseng* (Shen et al., 2022). In
511 contrast, Dhahri et al. (2023) reported that Ajwa seed polysaccharides, extracted using hot water
512 extraction, displayed a loose, branched organization with lower aggregation compared to MPS.
513 The increased temperature and pressure in the microwave unit's internal cavity induce the
514 rupturing and release of cellular and surface components, thereby enhancing the yield of PS
515 (Chen et al., 2022).

516 The water-retention properties of polysaccharides are crucial in food preparation, significantly
517 influencing the practical functionality and sensory characteristics of food products. Furthermore,
518 the oil-holding capacity of PS significantly impacts the mouthfeel, texture, and flavor of food

519 items. Our results suggest that MPS demonstrates good water solubility and commendable
520 capabilities in retaining both oil and water, consistent with findings reported by Noorbakhsh and
521 Rabbani Khorasgani (2023).
522 Rheological analysis is vital for understanding the flow properties of any polysaccharide. Such
523 analysis helps in assessing the textural, compositional, and structural changes that may occur
524 during various stages of food processing, packaging, and distribution. The storage and loss
525 modulus denote the solid and liquid-like characteristics of a sample. The higher G' and lower G''
526 values observed for MPS in all three tests performed signify the prevalence of elastic behavior
527 over viscous behavior. Additionally, the decrease in complex viscosity as shear rate and
528 frequency increase suggests a shear-thinning behavior (Shen et al., 2022). Studies by Niknam,
529 Mousavi and Kiani (2020) have also identified prominent solid elastic properties in
530 polysaccharides. However, in the time sweep test, G'' surpassed G' after a specific temperature,
531 indicating the transition from solid elastic properties to a more liquid, viscous nature. Similar
532 observations regarding polysaccharides displaying both solid and liquid characteristics
533 simultaneously have been reported in other studies (Li et al., 2017).

534 **4.2 Bioactivities of MPS**

535 Key bioactive components present in many plant systems, such as antioxidants, protect cells
536 against radical-induced stress and oxidative cell damage, reducing the risk of associated illnesses
537 (Dhahri et al., 2023). The remarkable radical scavenging ability of MPS, as indicated by their
538 heightened DPPH and ABTS scavenging activities, aligns with previous findings (Dhahri et al.,
539 2023; Noorbakhsh & Rabbani Khorasgani, 2023). The extraction method, molecular weight, and
540 monosaccharide composition significantly influence the radical scavenging potential of
541 polysaccharides (Mohammed, Naveed, & Jost, 2021). The microwave-assisted DES extraction
542 method can retain known components of date seeds, such as carbonyl and carboxyl groups,
543 bound flavonoids, and phenolics, responsible for the antioxidant effects observed in MPS

544 (Noorbakhsh & Rabbani Khorasgani, 2023; Wu et al., 2022). The presence of functional groups
545 like C = O and OH, identified through FTIR spectra, also plays a crucial role in the antioxidant
546 properties of polysaccharides. For instance, polysaccharides extracted from date seeds via
547 ultrasonic-assisted alkaline hot water extraction showed antioxidant activity comparable to MPS,
548 implying the influence of extraction method on such properties (Noorbakhsh & Rabbani
549 Khorasgani, 2023).

550 Furthermore, Dhahri et al. (2023) exploring the effects of hot water and ultrasound extraction on
551 date seed polysaccharides reported ABTS and DPPH scavenging activities that are comparable to
552 MPS, highlighting the potential antioxidant capabilities at a high concentration of 1000 µg/ml.
553 Acidic polysaccharides, containing galacturonic acid, exhibit greater antioxidant potential as the
554 acid binds to metal ions and scavenges DPPH free radicals. Heterogeneous polysaccharides like
555 MPS and *Moreinga oleifera* PS (Chen et al., 2017) also showcase higher DPPH radical
556 scavenging and, consequently, possess notable antioxidant potential. Overall, the high
557 antioxidant capacity observed in MPS could be attributed to factors such as lower molecular
558 weight, polydispersity, and varying monosaccharide composition.

559 Inhibition of diabetic-inducing enzymes with increasing concentrations of MPS indicates its
560 potential antidiabetic property. Increased α -amylase and α -glucosidase inhibitory activities were
561 observed, similar to the findings in *Schizophyllum commune* polysaccharide (Chen, Ragauskas,
562 & Wan, 2020). Additionally, the ACE inhibition percentage of MPS (60% at 1000 µg/ml MPS
563 concentration) suggests significant antihypertensive properties, in alignment with prior reports
564 on date seed flour and hydrolysate (60%) (Ambigaipalan & Shahidi, 2015). MPS also exhibited
565 diverse effects on cell viability, decreasing it in CaCO₂ cell lines while increasing it in MCF-7
566 cell lines. The lower IC₅₀ values of MPS, along with maximum inhibition of CaCO₂ cell lines,
567 indicate that MPS is potent even at low concentrations and, when administered, will display
568 lower systemic toxicity. Similar effects were observed with rice bran polysaccharides towards

569 S180 cancer cells (Wang et al., 2020). Microwave-extracted PS also displayed potent antitumor
570 activities against different sarcoma cells compared to conventional extracts. This further
571 emphasizes the impact of molecular weight on the biological activity of polysaccharides (Chen et
572 al., 2022).

573 Antimicrobial activities of polysaccharides vary based on the extraction method used and the
574 structural properties of the polysaccharide. Variations in the structure of cell walls in gram-
575 negative and gram-positive bacteria also affect their interactions with different plant
576 polysaccharides (Noorbakhsh & Rabbani Khorasgani, 2023). The MIC of MPS against common
577 food pathogens was found to be 0.588 mg/mL. *In vitro* digestion studies suggest that MPS
578 undergoes partial degradation to liberate reducing sugars, indicating its digestion behavior, which
579 differs based on monosaccharide compositions and molecular weight profiling. Acidic
580 polysaccharides often remain unchanged during gastrointestinal digestion. Indigestible or
581 partially digestible polysaccharides are likely to undergo enzymatic degradation by gut
582 microbiota, releasing reducing sugars that act as carbon sources for native flora. MPS was found
583 to act as a prebiotic for the effective growth of standard probiotic *Lactobacillus* strains, except
584 for *L. gasseri* and *L. rahnmosus*. A similar effect favoring the growth of *Lactobacillus plantarum*
585 was reported for polysaccharides from *Crataegus azarolus* seeds after 24 h of fermentation
586 (Khakpour, Hojjati, Jooyandeh, & Noshad, 2023).

587 **4.3 Influence of MPS on Fecal Fermentation**

588 A significant decrease in pH, total sugars, and an increase in gas production during the initial 24
589 h of fecal fermentation indicate the effective utilization of MPS by the gut microbiota. This
590 change in pH and gas production reflects the breakdown of undigested food components by gut
591 microbes into corresponding acids. The negligible presence of reducing sugars in the
592 fermentation broth and decrease in gas production after 24 h, suggests maximal utilization of
593 MPS by the gut microbiota. This indicates that polysaccharides are effectively hydrolyzed by the

594 native colonic microbiota, liberating reducing sugars used to produce fermentation by-products
595 like SCFAs (Wu et al., 2022). The production of SCFAs, associated with a reduction in pH
596 throughout fermentation, serves as an indicator of fermentability of polysaccharides and is
597 crucial for maintaining host metabolism, energy homeostasis, intestinal epithelial barrier
598 function, and preventing intestinal diseases (Zeyneb et al., 2021). Our findings align with (Wu et
599 al., 2021a), illustrating the major SCFAs released by polysaccharides—acetic, propionic, and
600 butyric acids.

601 MPS exhibited lower acetic acid (1476 $\mu\text{mol/g}$) compared to NC and GOS-P but higher
602 concentrations of butyric (488 $\mu\text{mol/g}$) and propionic acids (1220.473 $\mu\text{mol/g}$). The increase in
603 acetic acid is related to increased Bifidobacterium and Lactobacillus species in NC and GOS-P
604 compared to MPS, as previously noted by Wu et al. (2021b). The higher concentration of butyric
605 acid in MPS compared to the blank and GOS-P indicates the potential of microwave
606 polysaccharides in preventing insulin resistance and diabetes onset (Chen et al., 2022). The
607 relative abundance of firmicutes in the fermentation broth and the degradation of galactose and
608 galacturonic acid monomers of the polysaccharide likely contribute to the higher level of butyric
609 acid, which serves as an energy source for epithelial cells, aids in host gene regulation, and cell
610 apoptosis (Zeyneb et al., 2021).

611 Polysaccharides, as dietary fiber sources, modify gut microbiota, conferring health benefits to the
612 host by coordinating energy metabolism and immune system functions (Wu et al., 2021b). The
613 increased presence of Enteriobacteriales in MPS during the initial fermentation period followed
614 by their replacement by Bifidobacteriales after 48 h indicates increased acetic acid production at
615 the end of fermentation. Polysaccharides such as MPS stimulate the development of
616 Bifidobacteria, important for the production of acetate, thereby conferring anticancer and
617 probiotic functionality (Wu et al., 2021a). Additionally, the species phylogenetic tree indicates
618 the presence of anaerobes like Firmicutes, followed by Proteobacteria, Bacteroidetes, and

619 Actinobacteria, among which Bacteroidetes are the major gut microbiota species responsible for
620 polysaccharide degradation. Compared to the MPS group, Bacteroidetes were mainly present in
621 both NC and GOS-P groups. After 48 h of fermentation, MPS exhibited Bacteroidetes and
622 Clostridiales as the major gut bacteria responsible for polysaccharide degradation. This alteration
623 in the Bacteroidetes to Firmicutes ratio reduces the risk of insulin resistance and weight gain
624 (Han et al., 2022). PCA analysis results show that MPS and GOS-P shared common microflora
625 compared to NC, supported by microbial cluster analysis where bacteria in NC displayed distant
626 relations with GOS-P and MPS. Therefore, both GOS-P and MPS, when effectively
627 supplemented, will change the host gut microbiota (Wu et al., 2022; Wu et al., 2021a).

628 The species Spearman coefficient of MPS reported a strong positive correlation of
629 *Bifidobacterium longum* with Ruminococcus and Blautia, common SCFA-producing bacteria
630 responsible for degrading and utilizing dietary fiber. Fan et al. (2020) also reported the same
631 microbial correlations. The dominance of Firmicutes in NC and GOS-P and Proteobacteria in
632 MPS, in both the phylum and species heatmap, is consistent with Wu et al. (2021a). These
633 organisms effectively denature dietary fiber to release SCFAs and aid in energy harvest and the
634 elimination of obesity risks (Han et al., 2022). After 48 h of fermentation, Actinobacteria became
635 the major microbial species in fecal slurry supplemented with MPS. Similar reports of a reduced
636 percentage of Bacteroidetes with fermentation, and an increase in Firmicutes and Actinobacteria,
637 were reported by Wu et al. (2021a). Overall, the relative abundance of various beneficial genera
638 suggests that MPS supplementation effectively regulates gut microbial composition, potentially
639 offering health benefits.

640 **5. Conclusions**

641 The production of date seed polysaccharides using microwave-assisted deep eutectic solvent
642 extraction resulted in superior extraction yield and surface characteristics, exhibiting variable
643 molecular weights and diverse monosaccharide compositions resembling heteropolysaccharides.

644 Notably, the polysaccharide displayed significant antioxidant, antidiabetic, antihypertensive, and
645 antimicrobial activities that increased dose-dependent. Moreover, it stimulated the growth of
646 probiotic strains, indicating potential applications in food systems. Through hydrolysis by human
647 fecal microbiota, the polysaccharide released SCFAs such as acetic, butyric, and propionic acids,
648 enhancing overall metabolic functionality. These findings underscore the promising potential for
649 the effective utilization of MPS in functional food systems, aiding in improving overall intestinal
650 health. Future investigations could focus on exploring the impact of in vitro fecal fermentation
651 on the biological activities of MPS to provide a deeper understanding of the structure-function
652 relationship of the polysaccharide and its fermentation characteristics.

653 **Acknowledgment**

654 The authors are thankful to the United Arab Emirates University and Zayed Center for Health
655 Sciences (UAEU) for funding this project (grant number # 12R105). The authors thank Dr. Jaleel
656 and Mr. Mohammed Tarique for their technical support.

657 **Author Contribution**

658 A. Subhash: Writing-Original Draft, Investigation, Visualization, Formal analysis;

659 G. Bamigbade: Methodology, Formal analysis;

660 B. al-Ramadi: Writing-Review & Editing;

661 R.Y. Gan & S. Ranadheera: Writing-Review & Editing

662 A. Kamal-Eldin: Conceptualization, Writing-Review & Editing;

663 M. Ayyash: Conceptualization, Writing-Review & Editing, Supervision, Project administration, Funding
664 acquisition.

665 **Conflict of interest**

666 The authors declare no conflict of interest.

667 **Availability of Data and Materials**

668 All data generated or analyzed during this study are available upon request.

669 6. References

- 670 Al-Farsi, M. A., & Lee, C. Y. (2011). Usage of date (*Phoenix dactylifera* L.) seeds in human health and
671 animal feed. In *Nuts and seeds in health and disease prevention* (pp. 447-452). Elsevier.
672 <https://doi.org/10.1016/B978-0-12-375688-6.10053-2>
- 673 Al-Khalili, M., Al-Habsi, N., & Rahman, M. S. (2023). Applications of date pits in foods to enhance their
674 functionality and quality: A review. *Frontiers in Sustainable Food Systems*, 6.
675 <https://doi.org/10.3389/fsufs.2022.1101043>
- 676 Ambigaipalan, P., & Shahidi, F. (2015). Date seed flour and hydrolysates affect physicochemical
677 properties of muffin. *Food Bioscience*, 12, 54-60. <https://doi.org/10.1016/j.fbio.2015.06.001>
- 678 Ayyash, M., Stathopoulos, C., Abu-Jdayil, B., Esposito, G., Baig, M., Turner, M. S., Baba, A. S.,
679 Apostolopoulos, V., Al-Nabulsi, A., & Osaili, T. (2020). Exopolysaccharide produced by potential
680 probiotic *Enterococcus faecium* MS79: Characterization, bioactivities and rheological properties
681 influenced by salt and pH. *LWT*, 131, 109741. <https://doi.org/10.1016/j.lwt.2020.109741>
- 682 Bagade, S. B., & Patil, M. (2021). Recent advances in microwave assisted extraction of bioactive
683 compounds from complex herbal samples: a review. *Critical reviews in analytical chemistry*,
684 51(2), 138-149. <https://doi.org/10.1080/10408347.2019.1686966>
- 685 Brodkorb, A., Egger, L., Alminger, M., Alvito, P., Assunção, R., Ballance, S., Bohn, T., Bourlieu-Lacanal,
686 C., Boutrou, R., Carrière, F., Clemente, A., Corredig, M., Dupont, D., Dufour, C., Edwards, C.,
687 Golding, M., Karakaya, S., Kirkhus, B., Le Feunteun, S., . . . Recio, I. (2019). INFOGEST static
688 in vitro simulation of gastrointestinal food digestion. *Nature Protocols*, 14(4), 991-1014.
689 <https://doi.org/10.1038/s41596-018-0119-1>
- 690 Chen, C., Zhang, B., Huang, Q., Fu, X., & Liu, R. H. (2017). Microwave-assisted extraction of
691 polysaccharides from *Moringa oleifera* Lam. leaves: Characterization and hypoglycemic activity.
692 *Industrial Crops and Products*, 100, 1-11. <https://doi.org/10.1016/j.indcrop.2017.01.042>
- 693 Chen, X., Yang, J., Shen, M., Chen, Y., Yu, Q., & Xie, J. (2022). Structure, function and advance
694 application of microwave-treated polysaccharide: A review. *Trends in Food Science &*
695 *Technology*, 123, 198-209. <https://doi.org/10.1016/j.tifs.2022.03.016>
- 696 Chen, Z., Ragauskas, A., & Wan, C. (2020). Lignin extraction and upgrading using deep eutectic solvents.
697 *Industrial Crops and Products*, 147. <https://doi.org/10.1016/j.indcrop.2020.112241>
- 698 Dao, T. A. T., Webb, H. K., & Malherbe, F. (2021). Optimization of pectin extraction from fruit peels by
699 response surface method: Conventional versus microwave-assisted heating. *Food Hydrocolloids*,
700 113, 106475. <https://doi.org/10.1016/j.foodhyd.2020.106475>
- 701 Dhahri, M., Sioud, S., Alsuhaymi, S., Almulhim, F., Haneef, A., Saoudi, A., Jaremko, M., & Emwas, A.-
702 H. M. (2023). Extraction, Characterization, and Antioxidant Activity of Polysaccharides from
703 Ajwa Seed and Flesh. *Separations*, 10(2). <https://doi.org/10.3390/separations10020103>
- 704 DuBois, M., Gilles, K. A., Hamilton, J. K., Rebers, P. t., & Smith, F. (1956). Colorimetric method for
705 determination of sugars and related substances. *Analytical chemistry*, 28(3), 350-356.
706 <https://doi.org/10.1021/ac60111a017>
- 707 Fan, Q., Jiang, C., Wang, W., Bai, L., Chen, H., Yang, H., Wei, D., & Yang, L. (2020). Eco-friendly
708 extraction of cellulose nanocrystals from grape pomace and construction of self-healing
709 nanocomposite hydrogels. *Cellulose*, 27(5), 2541-2553. <https://doi.org/10.1007/s10570-020-02977-2>
- 710
- 711 FAOSTAT. *FAOSTAT*. <https://www.fao.org/faostat/en/#home>
- 712 Ghnimi, S., Umer, S., Karim, A., & Kamal-Eldin, A. (2017). Date fruit (*Phoenix dactylifera* L.): An
713 underutilized food seeking industrial valorization. *NFS Journal*, 6, 1-10.
714 <https://doi.org/10.1016/j.nfs.2016.12.001>
- 715 Guo, H., Fu, M. X., Zhao, Y. X., Li, H., Li, H. B., Wu, D. T., & Gan, R. Y. (2021). The Chemical,
716 Structural, and Biological Properties of Crude Polysaccharides from Sweet Tea (*Lithocarpus*
717 *litseifolius* (Hance) Chun) Based on Different Extraction Technologies. *Foods*, 10(8).
718 <https://doi.org/10.3390/foods10081779>
- 719 Han, X., Zhou, Q., Gao, Z., Lin, X., Zhou, K., Cheng, X., Chitrakar, B., Chen, H., & Zhao, W. (2022). In
720 vitro digestion and fecal fermentation behaviors of polysaccharides from *Ziziphus Jujuba* cv.

- 721 Pozao and its interaction with human gut microbiota. *Food Research International*, 162, 112022.
 722 <https://doi.org/10.1016/j.foodres.2022.112022>
- 723 Jiang, L., Ren, Y., Shen, M., Zhang, J., Yu, Q., Chen, Y., Zhang, H., & Xie, J. (2021). Effect of acid/alkali
 724 shifting on function, gelation properties, and microstructure of Mesona chinensis polysaccharide-
 725 whey protein isolate gels. *Food Hydrocolloids*, 117, 106699.
 726 <https://doi.org/10.1016/j.foodhyd.2021.106699>
- 727 Khakpour, S., Hojjati, M., Jooyandeh, H., & Noshad, M. (2023). Microwave-assisted extraction,
 728 optimization, structural characterization, and functional properties of polysaccharides from
 729 Crataegus azarolus seeds. *Journal of Food Measurement and Characterization*, 17(3), 2830-2840.
 730 <https://doi.org/10.1007/s11694-023-01831-w>
- 731 Koropatkin, N. M., Cameron, E. A., & Martens, E. C. (2012). How glycan metabolism shapes the human
 732 gut microbiota. *Nature Reviews Microbiology*, 10(5), 323-335.
 733 <https://doi.org/10.1038/nrmicro2746>
- 734 Li, T., Lyu, G., Liu, Y., Lou, R., Lucia, L. A., Yang, G., Chen, J., & Saeed, H. A. M. (2017). Deep Eutectic
 735 Solvents (DESs) for the Isolation of Willow Lignin (*Salix matsudana* cv. Zhuliu). *Int J Mol Sci*,
 736 18(11). <https://doi.org/10.3390/ijms18112266>
- 737 Manhivi, V. E., Venter, S., Amonsou, E. O., & Kudanga, T. (2018). Composition, thermal and rheological
 738 properties of polysaccharides from amadumbe (*Colocasia esculenta*) and cactus (*Opuntia* spp.).
 739 *Carbohydrate Polymers*, 195, 163-169. <https://doi.org/10.1016/j.carbpol.2018.04.062>
- 740 Mohammed, A. S. A., Naveed, M., & Jost, N. (2021). Polysaccharides; Classification, Chemical
 741 Properties, and Future Perspective Applications in Fields of Pharmacology and Biological
 742 Medicine (A Review of Current Applications and Upcoming Potentialities). *J Polym Environ*,
 743 29(8), 2359-2371. <https://doi.org/10.1007/s10924-021-02052-2>
- 744 Niknam, R., Mousavi, M., & Kiani, H. (2020). New Studies on the Galactomannan Extracted from
 745 *Trigonella foenum-graecum* (Fenugreek) Seed: Effect of Subsequent Use of Ultrasound and
 746 Microwave on the Physicochemical and Rheological Properties. *Food and Bioprocess
 747 Technology*, 13(5), 882-900. <https://doi.org/10.1007/s11947-020-02437-6>
- 748 Noorbakhsh, H., & Rabbani Khorasgani, M. (2023). Functional and chemical properties of Phoenix
 749 dactylifera L. Polysaccharides and the effect of date flesh and seed intervention on some blood
 750 biomarkers: A contrastive analysis. *Food Chemistry: X*, 19, 100834.
 751 <https://doi.org/10.1016/j.fochx.2023.100834>
- 752 Ren, B., Chen, C., Li, C., Fu, X., You, L., & Liu, R. H. (2017). Optimization of microwave-assisted
 753 extraction of *Sargassum thunbergii* polysaccharides and its antioxidant and hypoglycemic
 754 activities. *Carbohydrate Polymers*, 173, 192-201. <https://doi.org/10.1016/j.carbpol.2017.05.094>
- 755 Romdhane, M. B., Haddar, A., Ghazala, I., Jeddou, K. B., Helbert, C. B., & Ellouz-Chaabouni, S. (2017).
 756 Optimization of polysaccharides extraction from watermelon rinds: Structure, functional and
 757 biological activities. *Food Chemistry*, 216, 355-364.
 758 <https://doi.org/10.1016/j.foodchem.2016.08.056>
- 759 Shen, S., Zhou, C., Zeng, Y., Zhang, H., Hossen, M. A., Dai, J., Li, S., Qin, W., & Liu, Y. (2022).
 760 Structures, physicochemical and bioactive properties of polysaccharides extracted from *Panax
 761 notoginseng* using ultrasonic/microwave-assisted extraction. *LWT*, 154, 112446.
 762 <https://doi.org/10.1016/j.lwt.2021.112446>
- 763 Wang, J., Jing, W., Tian, H., Liu, M., Yan, H., Bi, W., & Chen, D. D. Y. (2020). Investigation of Deep
 764 Eutectic Solvent-Based Microwave-Assisted Extraction and Efficient Recovery of Natural
 765 Products. *ACS Sustainable Chemistry & Engineering*, 8(32), 12080-12088.
 766 <https://doi.org/10.1021/acssuschemeng.0c03393>
- 767 Wang, X., Yang, Z., Liu, Y., Wang, X., Zhang, H., Shang, R., Laba, C., Wujin, C., Hao, B., & Wang, S.
 768 (2022). Structural characteristic of polysaccharide isolated from *Nostoc commune*, and their
 769 potential as radical scavenging and antidiabetic activities. *Scientific Reports*, 12(1).
 770 <https://doi.org/10.1038/s41598-022-26802-x>
- 771 Wu, D.-T., Fu, M.-X., Guo, H., Hu, Y.-C., Zheng, X.-Q., Gan, R.-Y., & Zou, L. (2022). Microwave-
 772 Assisted Deep Eutectic Solvent Extraction, Structural Characteristics, and Biological Functions of

773 Polysaccharides from Sweet Tea (*Lithocarpus litseifolius*) Leaves. *Antioxidants*, 11(8), 1578.
774 <https://doi.org/10.3390/antiox11081578>

775 Wu, D.-T., Nie, X.-R., Gan, R.-Y., Guo, H., Fu, Y., Yuan, Q., Zhang, Q., & Qin, W. (2021a). In vitro
776 digestion and fecal fermentation behaviors of a pectic polysaccharide from okra (*Abelmoschus*
777 *esculentus*) and its impacts on human gut microbiota. *Food Hydrocolloids*, 114, 106577.
778 <https://doi.org/10.1016/j.foodhyd.2020.106577>

779 Wu, D. T., Feng, K. L., Huang, L., Gan, R. Y., Hu, Y. C., & Zou, L. (2021b). Deep Eutectic Solvent-
780 Assisted Extraction, Partially Structural Characterization, and Bioactivities of Acidic
781 Polysaccharides from Lotus Leaves. *Foods*, 10(10). <https://doi.org/10.3390/foods10102330>

782 Yang, J., Zamani, S., Liang, L., & Chen, L. (2021). Extraction methods significantly impact pea protein
783 composition, structure and gelling properties. *Food Hydrocolloids*, 117.
784 <https://doi.org/10.1016/j.foodhyd.2021.106678>

785 Yılmaz, T., & Şimşek, Ö. (2020). Potential health benefits of ropy exopolysaccharides produced by
786 *Lactobacillus plantarum*. *Molecules*, 25(14), 3293. <https://doi.org/10.3390/molecules25143293>

787 Zeyneb, H., Pei, H., Cao, X., Wang, Y., Win, Y., & Gong, L. (2021). In vitro study of the effect of quinoa
788 and quinoa polysaccharides on human gut microbiota. *Food science & nutrition*, 9(10), 5735-
789 5745. <https://doi.org/10.1002/fsn3.2540>

790

791

792

793

794

795

796

797

798

799

800

801

802

803

804

805

806

807 **Figure Captions:**

808 Figure 1: Characterization of MPS: FTIR spectra of MPS (A); DSC curve of MPS (B); TGA of
809 MPS (C); MPS Storage Modulus (D) during (a) Amplitude sweep, (b) Frequency sweep, and (c)
810 Time sweep

811

812 Figure 2: SEM images of MPS (A)-(F) at magnifications 250X, 1000X, 1500X, and 3000X,
813 respectively.

814
815 Figure 3: *In vitro* bioactivities of MPS at different concentrations ($\mu\text{g/mL}$). (A) bioactivities
816 expressed as inhibition percentage of GLU: α -Glucosidase, AMY: α -Amylase, ACE:
817 Angiotensin-Converting Enzyme, Caco-2: colon cancer cell line, and MCF-7: breast cancer cell
818 lines. (B) Minimum inhibitory concentration of MPS on different food pathogens, where A, B, C,
819 and D are *E. coli* 0157:H7, *S. aureus*, *S. Typhimurium*, and *L. monocytogenes*, respectively. Bars
820 are means \pm standard deviations (error bars). ^(a-b) Means with different lowercase letters of the
821 same parameter differed significantly ($P < 0.05$). Values are means \pm standard deviation of $n=6$
822

823 Figure 4: Effect of MPS during 0, 6, 12, 24 and 48 h of fecal fermentation for pH (A), gas
824 production (B), total sugar (C), reducing sugar (D), acetic acid production (E), propionic acid
825 production (F), and butyric acid production (G) of the sample groups: NC (negative control),
826 GOS-P (positive control) and MPS (microwave polysaccharide). Bars are means \pm standard
827 deviations (error bars) of $n=3$. ^{a-b} Means \pm standard deviations with different lowercase letters of
828 the same time differed significantly ($P < 0.05$). A-B Means \pm standard deviations with different
829 uppercase letters of the same group differed significantly ($P < 0.05$).

830
831 Figure 5: Effect of MPS on gut microbiome composition during fecal fermentation. Venn
832 diagram (A), box plot of different indices of alpha diversity (B), NMDS plot (C), NMDS plot
833 (D), combination graph of UPGMA cluster tree and order abundance bar plot (E), and species
834 phylogenetic analysis (F) of the sample groups, where A, B, and F are the sample groups: A- NC
835 (negative control), B- GOS-P (positive control), and E- MPS.

836
837 Figure 6: Effect of MPS on different microbial functions during fecal fermentation. Histograms
838 of KEGG pathways abundance at Levels 1-3 (A-C), heatmap of COG pathways (D), heatmap of
839 MetaCyc pathways (E), and species spearman coefficients analysis of OTUs in each sample
840 group (F), where A, B, and F are the sample groups: A- NC- (negative control), B- GOS-P
841 (positive control), and E- MPS.

842
843

844
845
846
847
848
849
850
851

Table 1: *In vitro* antioxidant activities of MPS¹ at different concentrations.

MPS (µg/mL)	DPPH (%) ²	ABTS (%)	SAS (%)	SOD (%)	MC (%)	LO (%)	TAC (µg/mL)	RP (µg/mL)	HRS (%)	HP (%)	FRAP (µg/mL)
125	17.9±1.9 ^{d3}	11.7±3.4 ^d	18.8±1.4 ^d	20.0±0.2 ^d	384.8±55.6 ^d	15.0±0.1 ^d	28.3±4.5 ^d	18.5±0.4 ^d	14.8±0.7 ^d	26.6±1.0 ^d	51.2±5.1 ^d
250	31.7±5.8 ^c	38.4±0.7 ^c	32.0±0.2 ^c	38.4±0.2 ^c	3698.1±155.1 ^c	27.2±0.8 ^c	178.2±19.4 ^c	52.9±7.7 ^c	36.9±0.5 ^c	57.5±0.1 ^c	124.2±7.7 ^c
500	52.9±1.8 ^b	44.1±1.2 ^b	44.2±1.1 ^b	50.8±0.6 ^b	5110.±65.6 ^b	44.5±0.5 ^b	603.6±45.2 ^b	72.1±2.8 ^b	52.8±0.7 ^b	66.1±0.1 ^b	311.4±7.9 ^b
1000	60.8±.16 ^a	67.6±0.5 ^a	49.8±0.2 ^a	57.4±0.2 ^a	6343.8±545.4 ^a	60.0±0.2 ^a	683.4±49.7 ^a	171.2±19.1 ^a	64.6±0.7 ^a	68.8±0.1 ^a	538.3±82.8 ^a
IC ₅₀ (µg/ml)	305.8	78.5	235.1	117.2	66.9	427.6	324.2	478.8	73.6	49.7	580.2

852 ¹ Polysaccharide produced by microwave-assisted deep eutectic solvent extraction of date seeds.

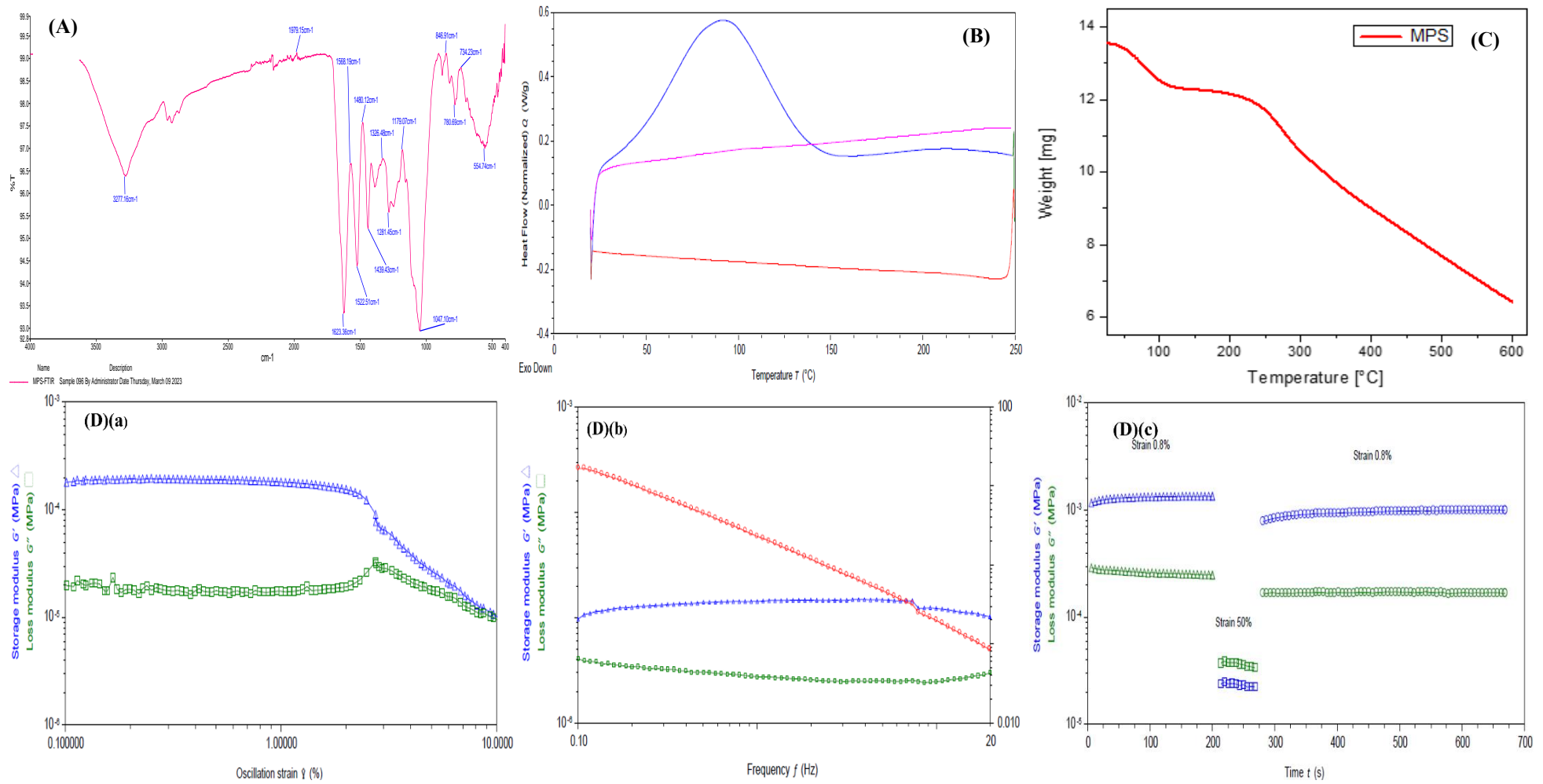
853 ² DPPH: 1-diphenyl-2-picrylhydrazyl; ABTS: 2,2'-asino-bis(3-ethylbenzene-thiazoline-6-sulphonic acid; SD: Superoxide Dismutase; SAS: Superoxide
854 Anion Scavenging; TAC: Total Antioxidant Capacity; FRAP: Ferric Reducing Antioxidant Power; RP: Reducing Power, IC₅₀: Half Maximal Inhibitory
855 Concentration.

856 ³ Values are means ± standard deviation of n=6

857 ^(a-d) Means ± standard deviations with different lowercases, at the same column, differ significantly ($P < 0.05$).

858
859
860
861
862
863
864
865
866
867
868
869
870
871
872
873
874
875
876
877
878

879
880
881
882
883
884
885
886
887
888
889

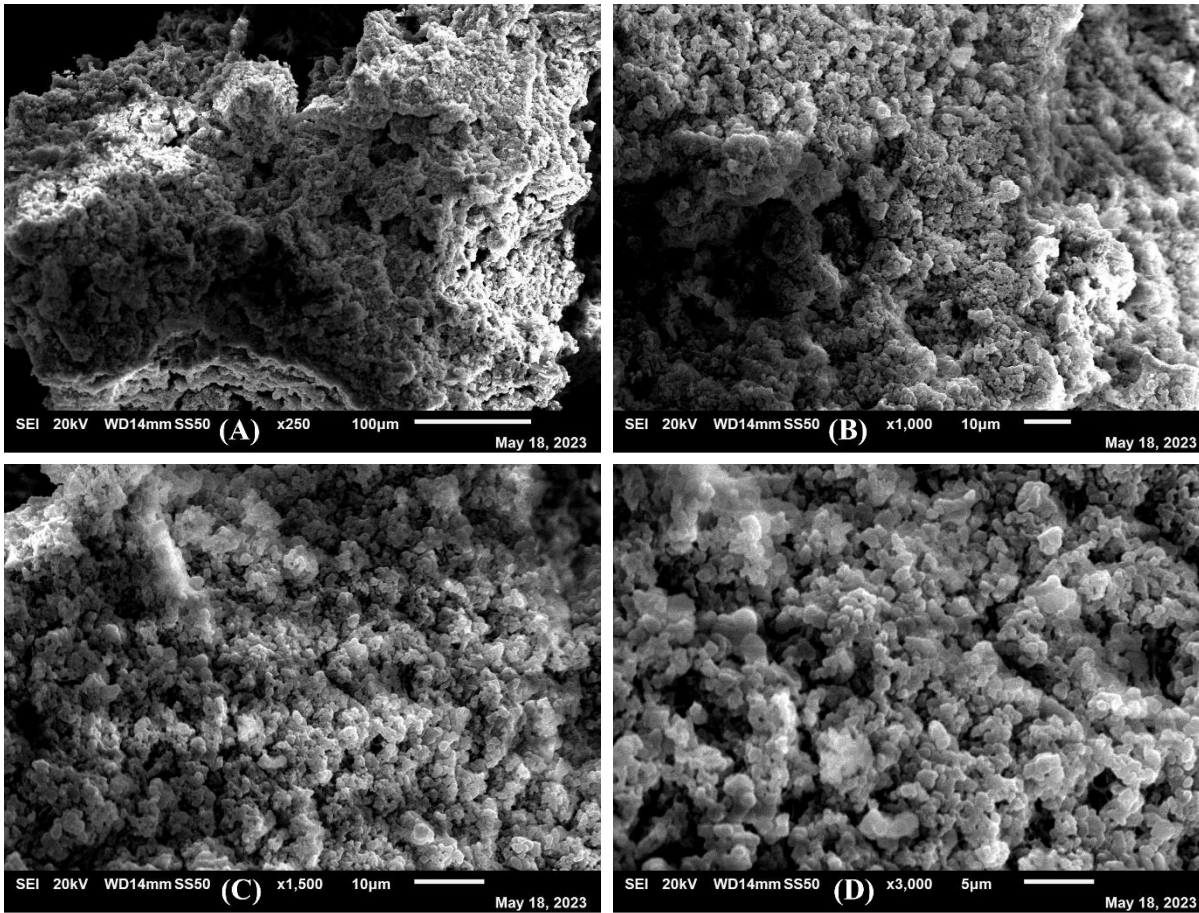


890

891
892
893
894
895
896
897

Figure 1

898
899
900
901
902
903

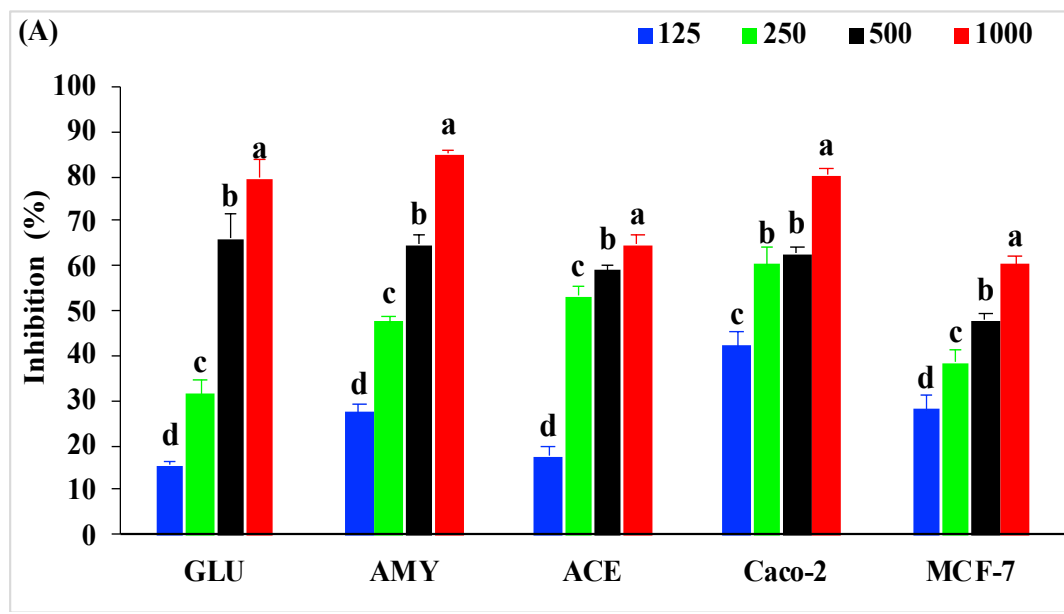


904

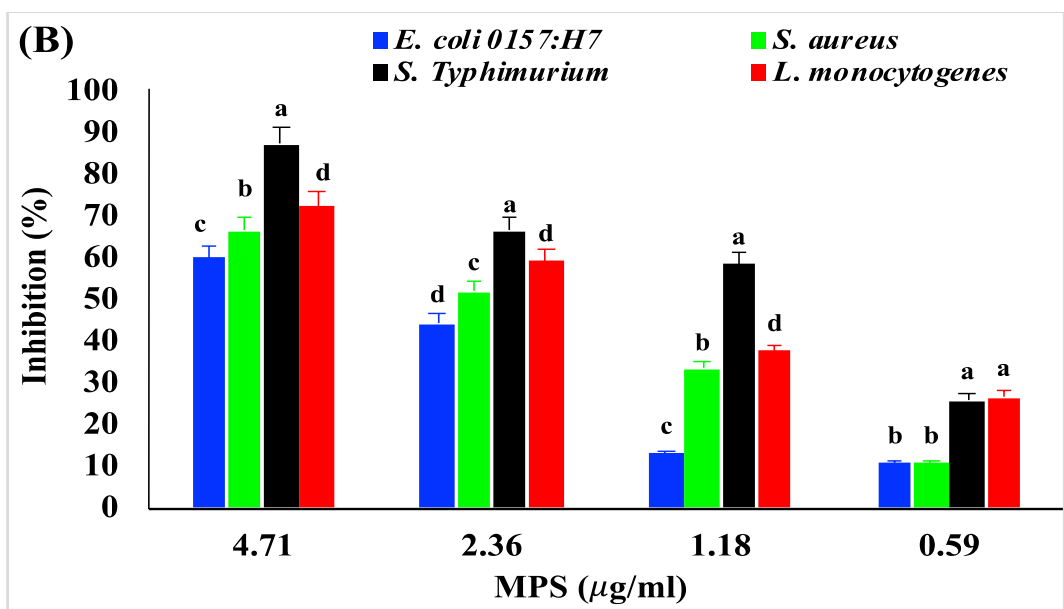
905
906
907
908
909
910
911
912
913
914
915
916
917
918
919
920
921
922
923
924
925
926
927
928
929
930
931

Figure 2: SEM images of MPS (A)-(F) at magnifications 250X, 1000X, 1500X, and 3000X, respectively.

932
933
934
935



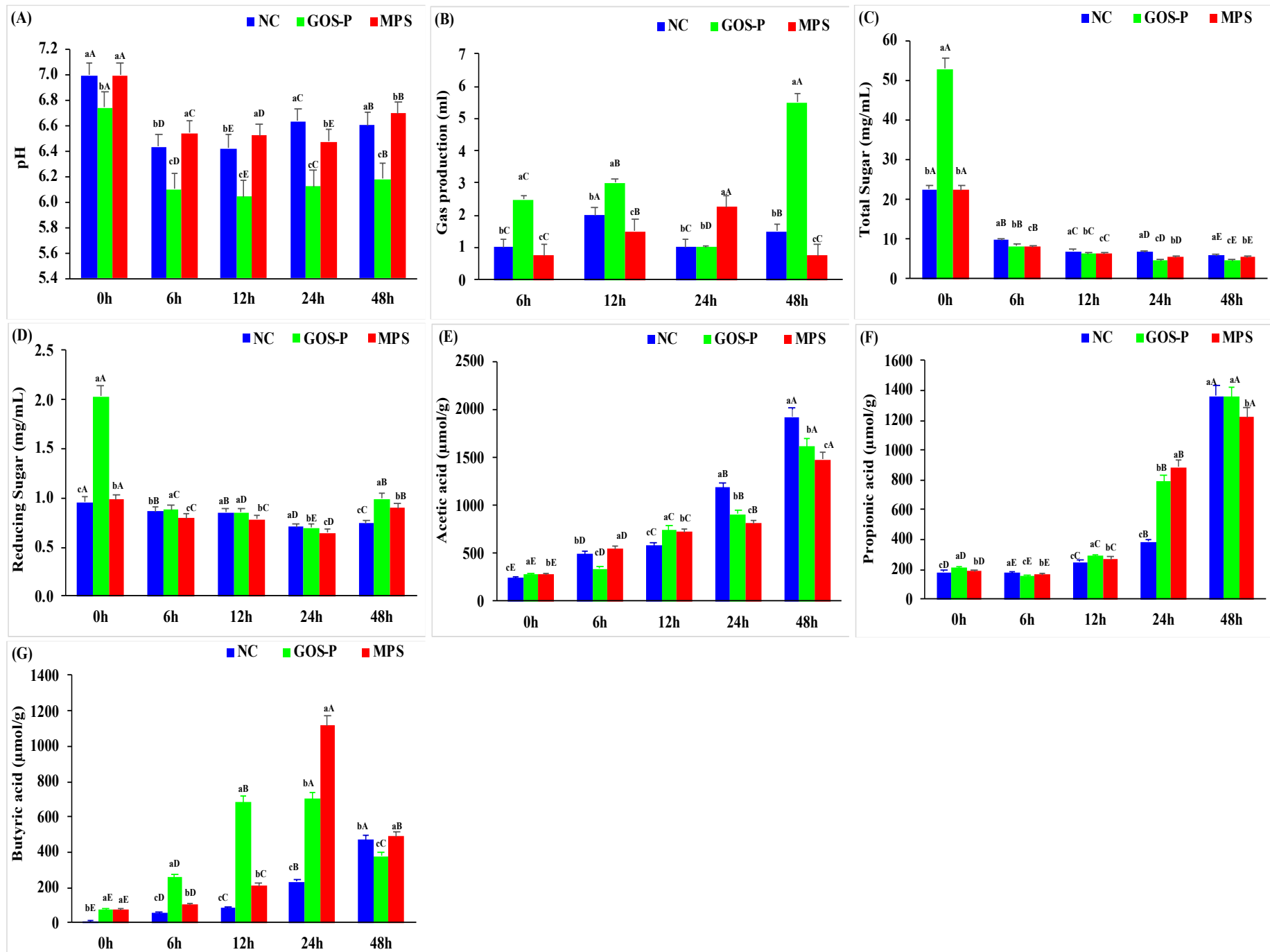
936
937



938
939
940
941
942
943
944
945
946
947
948
949

Figure 3: *In vitro* bioactivities of MPS at different concentrations (µg/mL). (A) bioactivities expressed as inhibition percentage of GLU: α -Glucosidase, AMY: α -Amylase, ACE: Angiotensin-Converting Enzyme, Caco-2: colon cancer cell line, and MCF-7: breast cancer cell lines. (B) Minimum inhibitory concentration of MPS on different food pathogens, where A, B, C, and D are *E. coli* 0157:H7, *S. aureus*, *S. Typhimurium*, and *L. monocytogenes* respectively. Bars are means \pm standard deviations (error bars). ^(a-b) Means with different lowercase letters of the same parameter differed significantly ($P < 0.05$). Values are means \pm standard deviation of n=6

950
951



952

953

954

955 Figure 4: Effect of MPS during 0, 6, 12, 24 and 48 h of fecal fermentation for pH (A),
956 gas production (B), total sugar (C), reducing sugar (D), acetic acid production (E),
957 propionic acid production (F), and butyric acid production (G) of the sample groups: NC (negative control), GOS-P (positive control) and MPS (microwave polysaccharide).
958 Bars are means ± standard deviations (error bars) of n=3. ^{a-b} Means ± standard deviations with different lowercase letters of the same time differed significantly ($P < 0.05$). A-B
959 Means ± standard deviations with different uppercase letters of the same group differed significantly ($P < 0.05$).
960
961