

Supporting information for

Original article

Colon-targeted oral delivery of melatonin for regulating macrophage activity and intestinal barrier in ulcerative colitis

Supplementary methods

siRNA transfection on RAW 264.7 macrophages

RAW 264.7 cells (5×10^5 cells/well) were seeded in 6-well plates in 2 mL DMEM overnight. For transfection, 3 μ L of 20 μ M STAT3 siRNA (cat# 20848-3) and 3 μ L lipofectamine 2000 (Invitrogen, cat# 11668-019) were separately diluted in 100 μ l Opti-MEM (Gibco, cat# 31985-070), incubated for 5 min at RT, combined, and further incubated for 20 min to form complexes. Cell culture medium was replaced with 1.8 mL fresh DMEM, and 200 μ L transfection complexes were added dropwise. After a transfection period of 48 h, cells were subjected to downstream assays such as LPS stimulation. Knockdown efficiency was verified by qRT-PCR and Western blot ($> 70\%$ STAT3 reduction). Scrambled siRNA (SN-1001) served as a negative control [1, 2].

Western blot assay

Briefly, RAW 264.7 cells were lysed in RIPA buffer containing 1% protease inhibitor and 1% EDTA (Thermo Fisher Scientific). The lysates were homogenized and centrifuged to collect protein samples, which were then determined and subjected to SDS-PAGE and membrane transfer. The membrane was then blocked with BSA 5% and sequentially incubated with primary antibodies against STAT3 (CST, cat# 4904S), p-STAT3 (CST, cat# 9145S), and GAPDH (ABclonal, cat# A19056), followed by HRP-conjugated secondary antibodies (Invitrogen, cat# 31460). Protein expression was visualized using enhanced chemiluminescence (ECL) detection.

***In vitro* cytotoxic assay**

Caco-2 epithelial cells and RAW 264.7 macrophages were maintained in complete medium. Cells (1×10^4 per well) were seeded in 96-well plates overnight. Subsequently, cells were treated with serial concentrations of free melatonin or Blank-MS for 24 h. Cell viability was then assessed using the WST-8 assay (Biomax, cat# QM2500) following the instructions from the manufacturer.

Synthesis and confirmation of thioketal polymer

The thioketal polymer with ROS-responsiveness was synthesized via step-growth polymerization, as described in our previous study [3]. Briefly, 245 mg 4,4'-bis(mercaptomethyl)biphenyl (Sigma-Aldrich, cat# 716049) was dissolved in 15 mL toluene containing 250 μ L 2,2-dimethoxypropane (DMP; Sigma-Aldrich, cat# D136808). The solution was heated to 75 °C, after which p-toluenesulfonic acid catalyst (2 mg in 250 μ L ethyl acetate; Sigma-Aldrich, cat# 402885) was added. Additional DMP in toluene (500 μ L aliquots, prepared as 500 μ L DMP in 10 mL toluene) was provided every 30 min over 10 h, followed by a final 500 μ L addition of DMP. The reaction was then stirred overnight. Then, cold hexane (−20 °C) was used to precipitate the synthesized polymer. The product was then washed twice with distilled water and dried to obtain a brown viscous product. Polymer structure was analyzed and confirmed by ^1H and ^{13}C NMR spectroscopy (500 MHz, Varian Unity Inova).

Measurement of luminal pH of the gastrointestinal tract in mice

Healthy mice and DSS-induced colitis mice were fasted overnight and euthanized by CO_2 inhalation. The gastrointestinal tract was rapidly dissected, and luminal contents were collected from distinct regions (stomach, small intestine, large intestine). For samples with insufficient volume, 50 μ L sterile water was added, gently homogenized, and mixed. Luminal pH was measured immediately using a calibrated micro pH electrode to limit post-mortem alterations. Region-specific pH gradients were recorded directly from GI contents [4].

***In vivo* safety of META**

Healthy mice received daily oral administration of META for 7 consecutive days. Serum was collected for biochemical analysis of liver function (GOT, GPT; Asan Set Assay kit) and kidney

function (BUN; Colorimetric Detection Kit, Invitrogen). Major organs were harvested for histological evaluation.

Supplementary figures

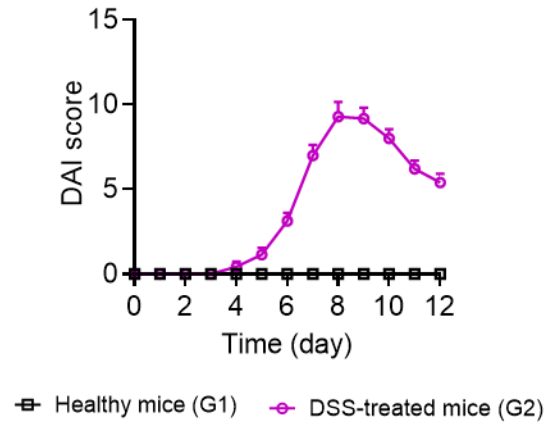


Figure S1. DAI score of DSS-induced mice colitis (n = 4-7 mice). Data are presented as mean $\pm$ s.e.m.

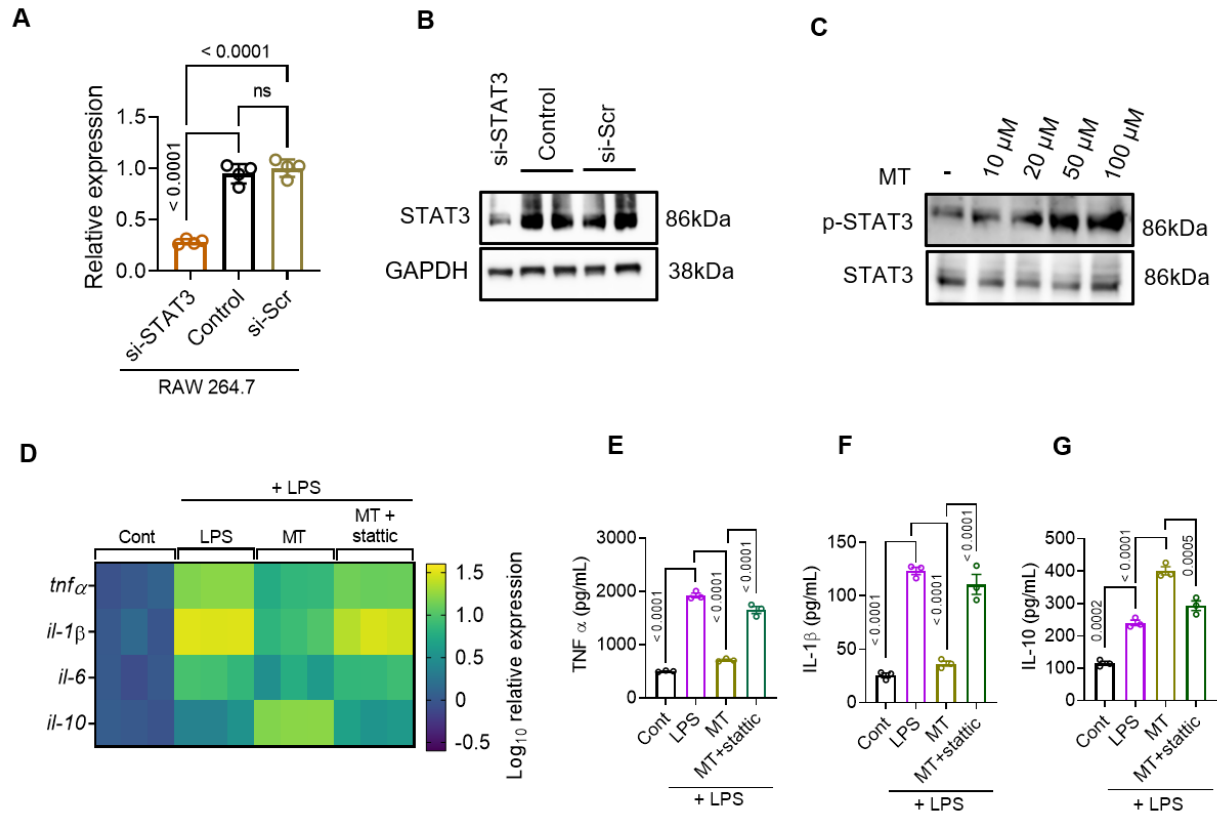


Figure S2. Melatonin regulates macrophage activity through STAT3 signaling. **A**, Expression of STAT3 mRNA in RAW 264.7 cells (n = 4). **B**, Expression of STAT3 protein in RAW 264.7 cells. **C**, Melatonin activates STAT3 in LPS-treated RAW 264.7 cells. **D**, Modulatory effects of melatonin on macrophage polarization after STAT3 inhibition, using static 5 μ M, quantified by qRT-PCR (n = 3). **E-G**, ELISA quantification of extracellular cytokines, including TNF- α , IL-1 β , and IL-10, respectively (n = 3). Data in (**A**, **E**, **F**, **G**) are presented as mean $\pm$ s.d. Data are analyzed using one-way ANOVA with Tukey's multiple comparisons test.

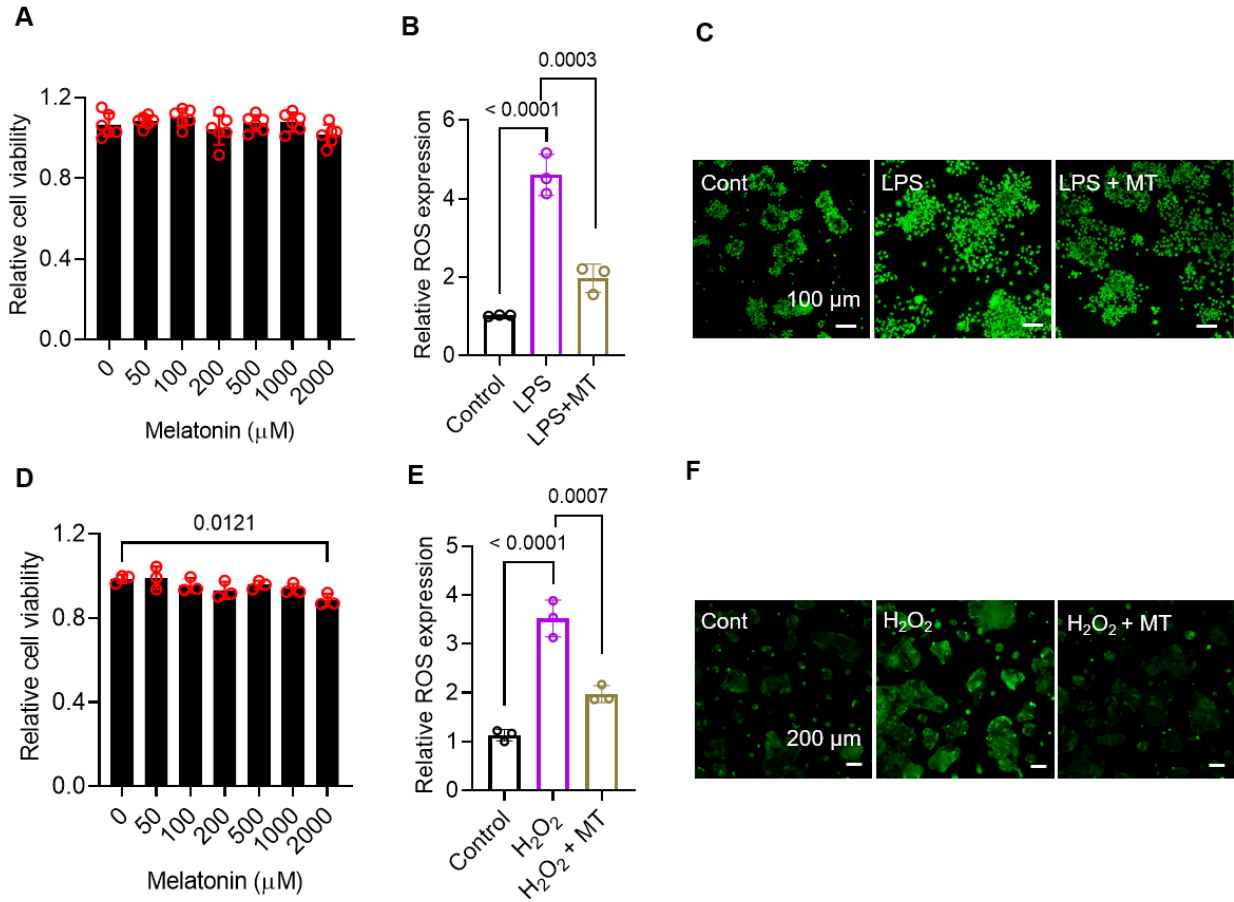
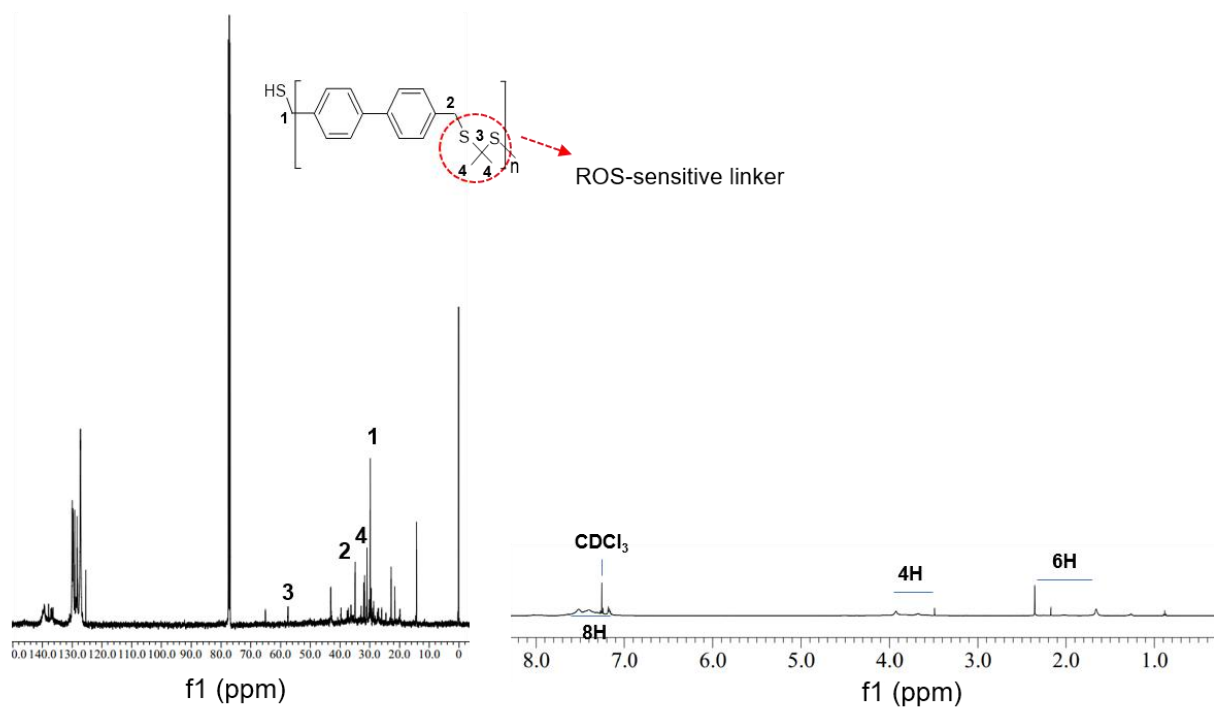


Figure S3. *In vitro* cytotoxicity of melatonin and its antioxidant effects. **A**, Cytotoxicity of melatonin on RAW 264.7 macrophages (n = 6). **B**, The antioxidative capacity of melatonin in RAW 264.7 cells (n = 3). **C**, Imaging ROS levels in RAW 264.7 cells. Scale bar 100 μ m. **D**, Cytotoxicity of melatonin on Caco-2 epithelial cells (n = 3). **E**, The antioxidative capacity of melatonin in Caco-2 cells (n = 3). **F**, Imaging ROS levels in Caco-2 cells. Scale bar 100 μ m. Data in (**A**, **B**, **D**, **E**) are presented as mean $\pm$ s.d. Data in (**A**, **D**) are analyzed using an unpaired two-tailed t-test (significant p values are shown). Data in (**B**, **E**) are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown).

A



B

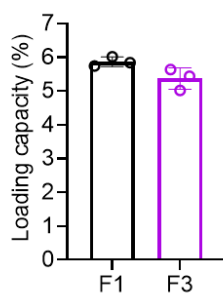


Figure S4. Synthesis and confirmation of thioketal polymer. **A**, Chemical structure of thioketal polymer was confirmed by ¹H and ¹³C NMR spectroscopy. **B**, Image of synthesized thioketal polymer. Chemical formulation was created with ChemSketch software (ACD/Labs).

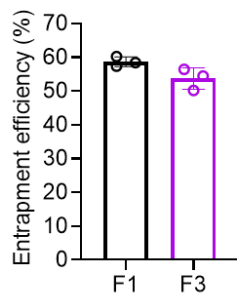
A

Formulation	Melatonin (%)	FS 30 D (%)	Thioketal polymer (%)
F1 (ME-MS)	10	90	0
F2 (META)	10	45	45
F3 (MTK-MS)	10	0	90

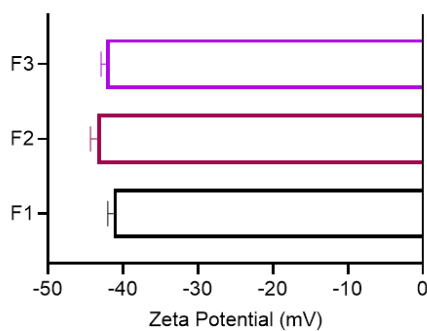
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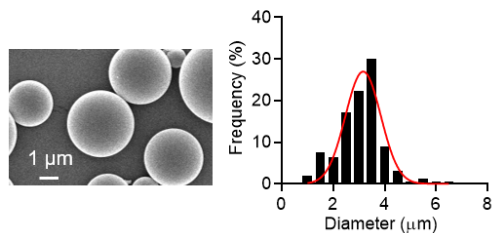
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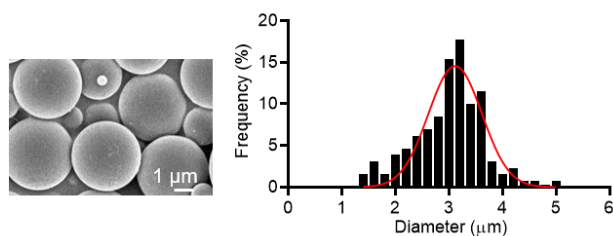
D



E



F



G

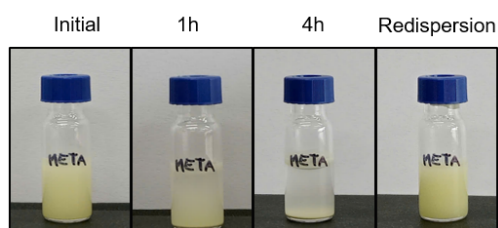


Figure S5. Optimization of microsphere formulations. **A**, Different formulations of microspheres for optimization. **B**, Loading capacity of ME-MS and MTK-MS ($n = 3$). **C**, Entrapment efficiency of ME-MS and MTK-MS ($n = 3$). **D**, Zeta potential of microspheres ($n =$

3). **E**, Morphology and size distribution of ME-MS. **F**, Morphology and size distribution of MTK-MS. Scale bar: 1 μm . **G**, META dispersibility in water.

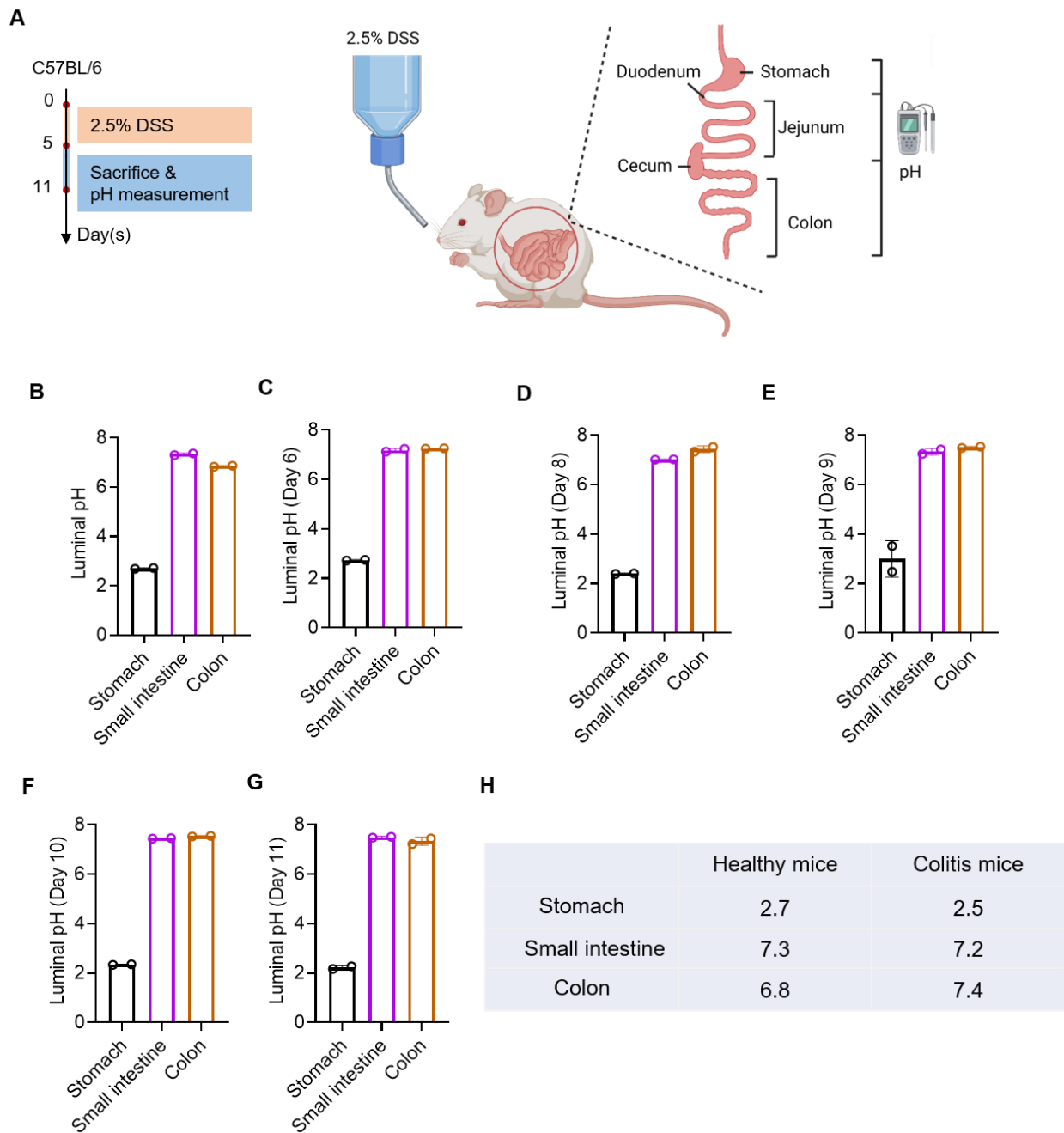


Figure S6. Measurement of luminal pH in the mice GI tract. **A**, Induction of acute colitis model and measurement of luminal pH in GI tract. **B**, Luminal pH in GI tract of healthy mice (n = 2). **C-G**, Luminal pH in GI tract of DSS-induced colitis mice at day 6 (**C**), day 8 (**D**), day 9 (**E**), day 10 (**F**), and day 11 (**G**) (n = 2). **H**, Average of luminal pH in the GI tract of healthy and colitis mice,

calculation based on the average pH value of all assessed mice for each group. Figure **S6A** was created with BioRender.com.

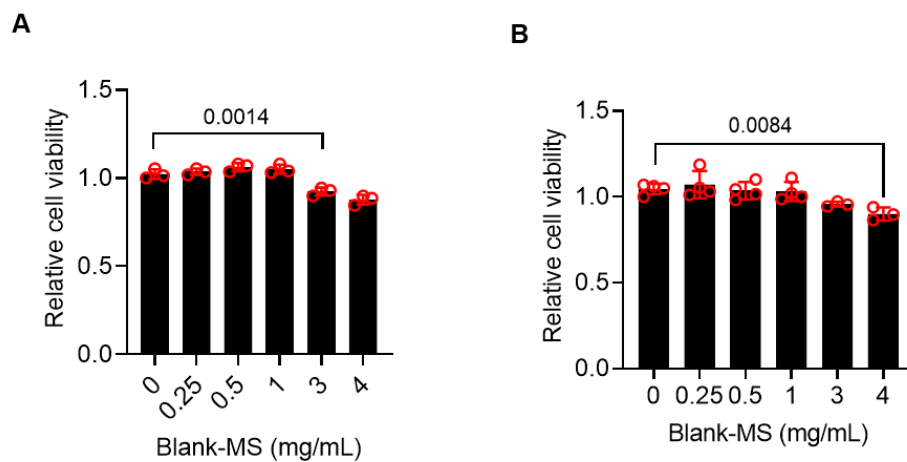
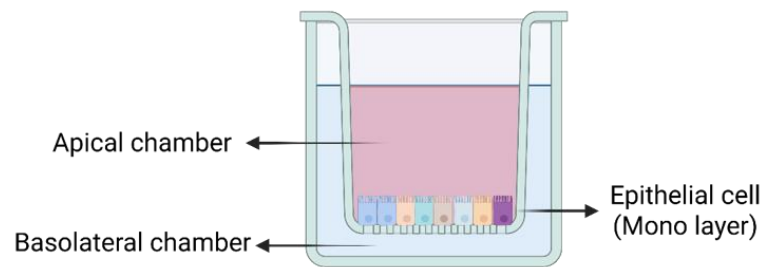


Figure S7. Cytotoxicity of Blank-MS. A, on Caco-2 cells (n = 3). **B,** on RAW 264.7 cells (n = 3-4). Data are presented as mean $\pm$ s.d. Data are analyzed using an unpaired two-tailed t-test (significant p-values are shown). Blank-MS indicating microsphere includes Eudragit FS 30 D and thioketal polymer without melatonin.

A



B



Figure S8. Caco-2 monolayer formation. **A**, Transwell model for Caco-2 layer. **B**, Fluorescence images of Caco-2 monolayer. Scale bar 100 μm . Figure **S8A** was created with BioRender.com.

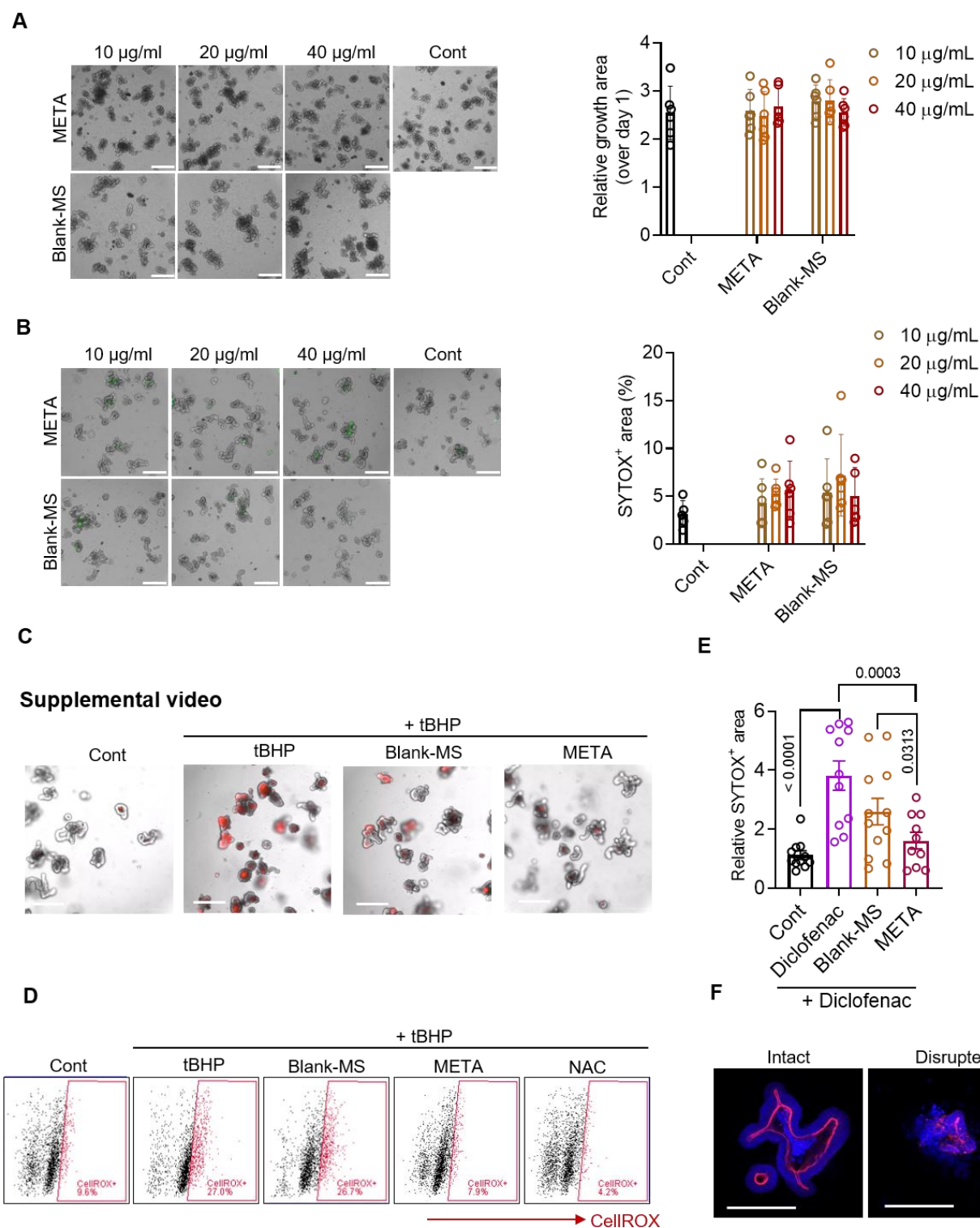


Figure S9. Impact of META on intestinal organoids (IOs). **A**, Images and relative growth area of IOs after treatment of META and Blank-MS (n = 6). Scale bar: 300 µm. **B**, Images and

percentage of SYTOX⁺ area in IOs after treatment of META and Blank-MS (n = 5-6). Scale bar: 300 μ m. **C**, Representative image of supplemental video of ROS levels in tBHP-induced IOs damage, scale bar: 300 μ m. Please see the full movie in the supplemental video file for reference. **D**, Representative results of flow cytometric analysis of ROS-high cells in tBHP-induced IOs damage. N-Acetylcysteine (NAC) was included as a control. **E**, Relative SYTOX⁺ area in diclofenac-induced IOs damage after treatments (n = 10-12). **F**, F-actin staining in healthy and diclofenac-disrupted IOs, scale bar: 300 μ m. Data in (**A**, **B**, **E**) are presented as mean $\pm$ s.d. Data in (**E**) are analyzed using one-way ANOVA with the two-stage linear step-up method of Benjamini, Krieger, and Yekutieli (significant q values are shown).

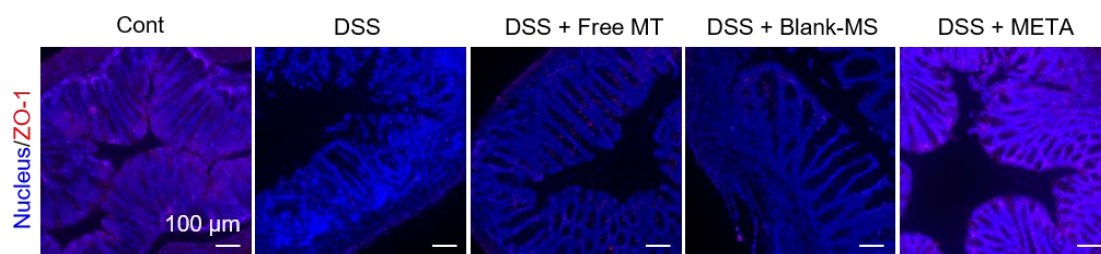


Figure S10. Immunofluorescence staining of ZO-1 protein in colon tissue. Scale bar: 100 μm .

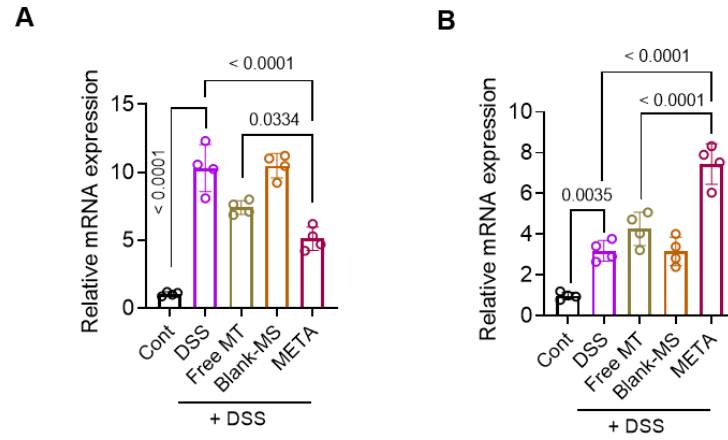


Figure S11. mRNA expression of CD86 and CD206 in the colonic mucosae. A, CD86 and B, CD206 (n = 4). Mononuclear cells were isolated from the colonic lamina propria and subjected to qRT-PCR analysis. Data are presented as mean $\pm$ s.d. Data are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown).

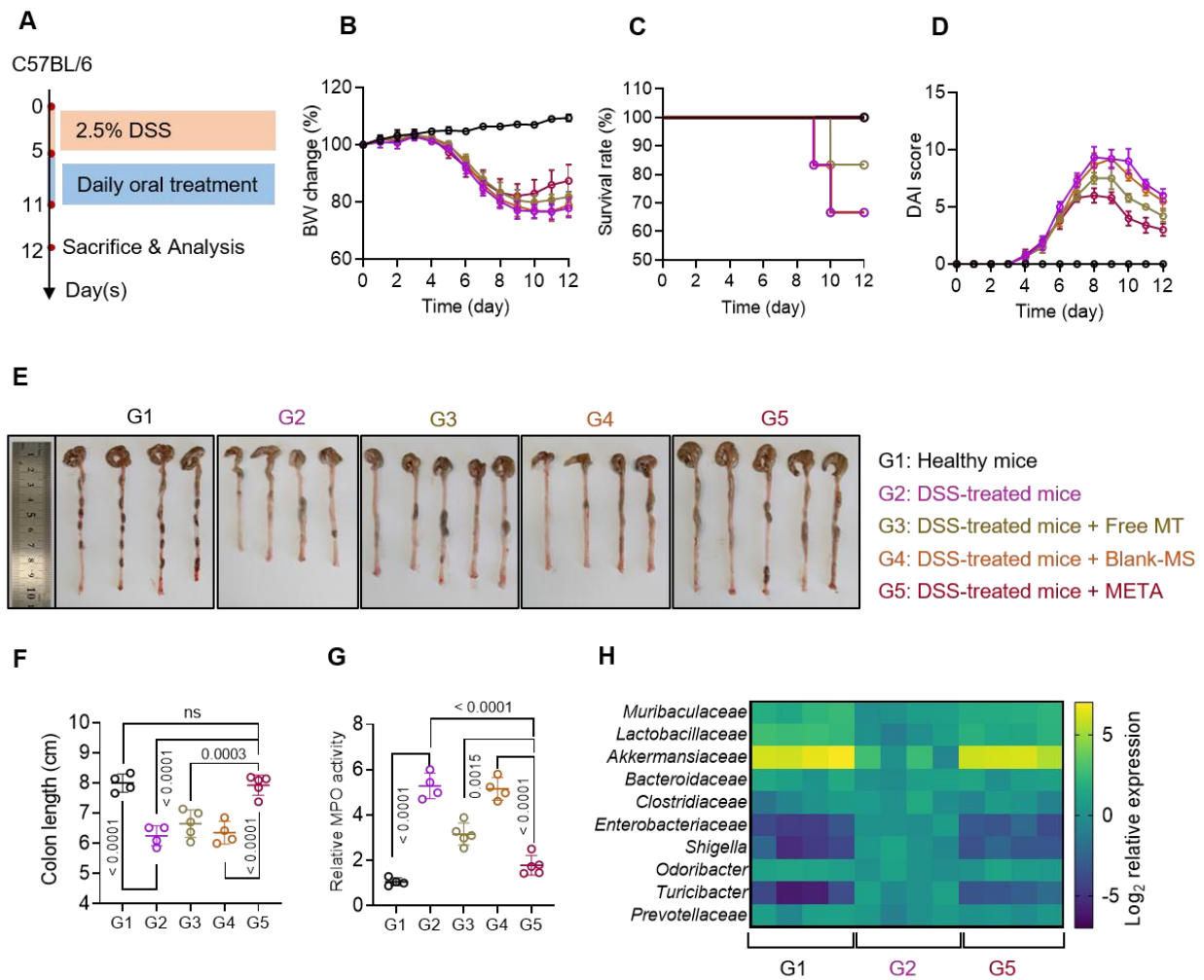


Figure S12. META effectively ameliorated UC. **A**, Induction of the mouse colitis model through oral DSS delivery and subsequent drug treatment. G1: Healthy mice, G2: DSS-treated mice, G3: DSS-treated mice + free melatonin (MT), G4: DSS-treated mice + Blank-MS, G5: DSS-treated mice + META. **B**, Body weight change measurement (n = 4-6 mice). **C**, Survival rate of mice (n = 4-6 mice). **D**, DAI score of mice (n = 4-6 mice). **E**, Images of colon tissues. **F**, Colon length measurement (n = 4-5 mice). **G**, Myeloperoxidase (MPO) activity in colon tissues analysis (n = 4-5 mice). **H**, Abundance of microbiota in the colon was quantified using qRT-PCR (n = 4 mice). Data in (**B**, **D**, **F**, **G**) are presented as mean $\pm$ s.d. Data in (**F**, **G**) are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown).

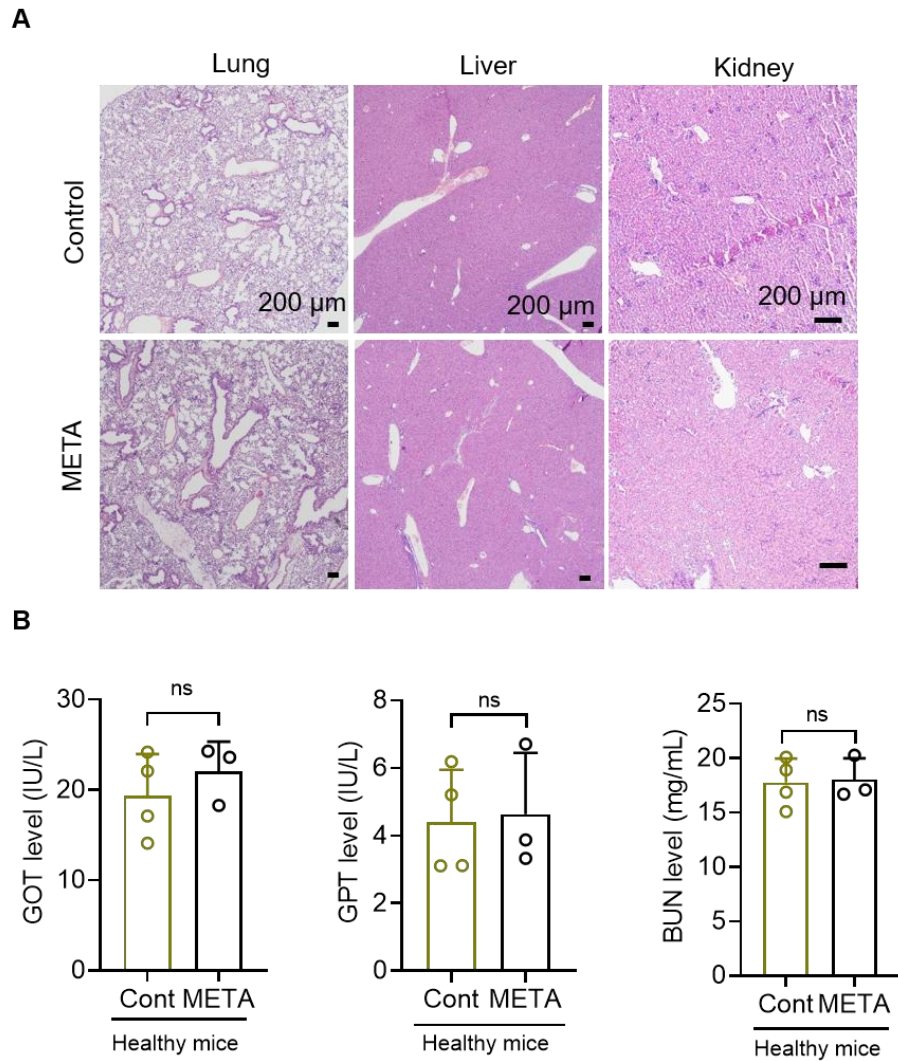


Figure S13. *In vivo* safety of META. **A**, Histological assessments (H&E) of major organs collected from mice after META treatment. Scale bar: 200 μ m. **B**, Serum level quantification of biochemical index in mice, including GOT, GPT (liver function), and BUN (kidney function) (n = 3-4 mice). Data in (**B**) is presented as mean $\pm$ s.d. Data in (**B**) is analyzed using an unpaired two-tailed t-test. ns, no significance.

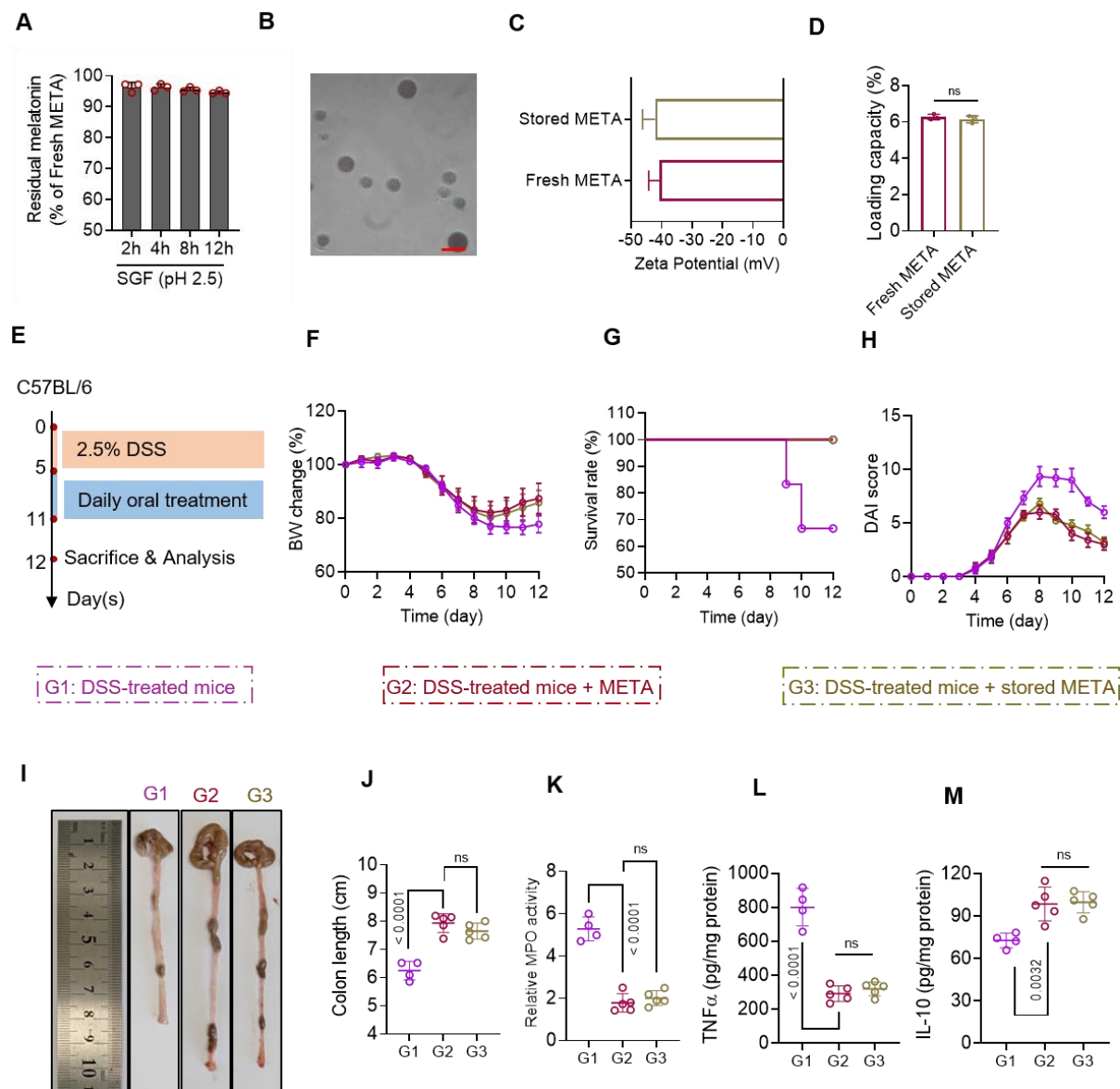


Figure S14. The stability of META. **A**, META stability under SGF conditions (n = 3). **B**, META morphology after 3-month storage at room temperature (RT). Scale bar: 5 μ m. **C**, Zeta potential of META after 3-month storage at RT. **D**, Melatonin loading capacity of META after 3-month storage at RT. **E**, Induction of the mouse colitis model through oral DSS delivery and subsequent drug treatment. G1: DSS-treated mice, G2: DSS-treated mice + fresh META, G3: DSS-treated mice + stored META. **F**, Body weight change measurement (n = 5-6 mice). **G**, Survival rate of

mice (n = 5-6 mice). **H**, DAI score of mice (n = 5-6 mice). **I**, Representative images of colon tissues. **J**, Colon length measurement (n = 4-5 mice). **K**, Myeloperoxidase (MPO) activity in colon tissues analysis (n = 4-5 mice). **L-M**, ELISA measurement of colonic TNF- α (**L**) and IL-10 (**M**) (n = 4-5 mice). Data of **G1 (DSS-treated mice)** and **G2 (DSS-treated mice + META)** in this figure, including data in (**F, G, H, I, J, K**), were extracted from Figure **S12** (Supplementary data) due to the same *in vivo* trial induction. Data are presented as mean $\pm$ s.d. Data are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown).

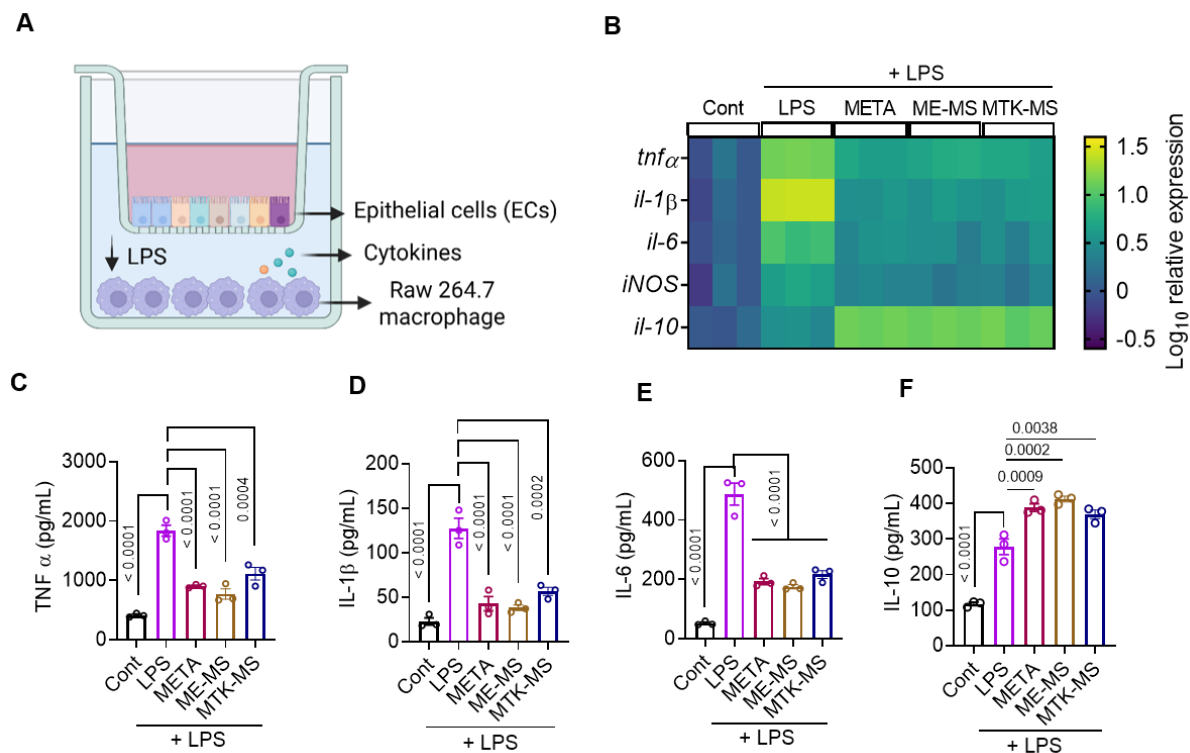


Figure S15. *In vitro* anti-inflammatory effects of individual polymer-based microspheres. A, Co-culture model of the epithelial Caco-2 cells (ECs) and RAW 264.7 macrophages. **B,** Expression of inflammatory and anti-inflammatory genes in RAW cells (n = 3). **C-F,** ELISA quantification of extracellular cytokines, including TNF- α , IL-1 β , IL-6, and IL-10, respectively (n = 3). Data in (C, D, E, F) are presented as mean $\pm$ s.d. Data in (C, D, E, F) are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown). Figure S15A was created with BioRender.com.

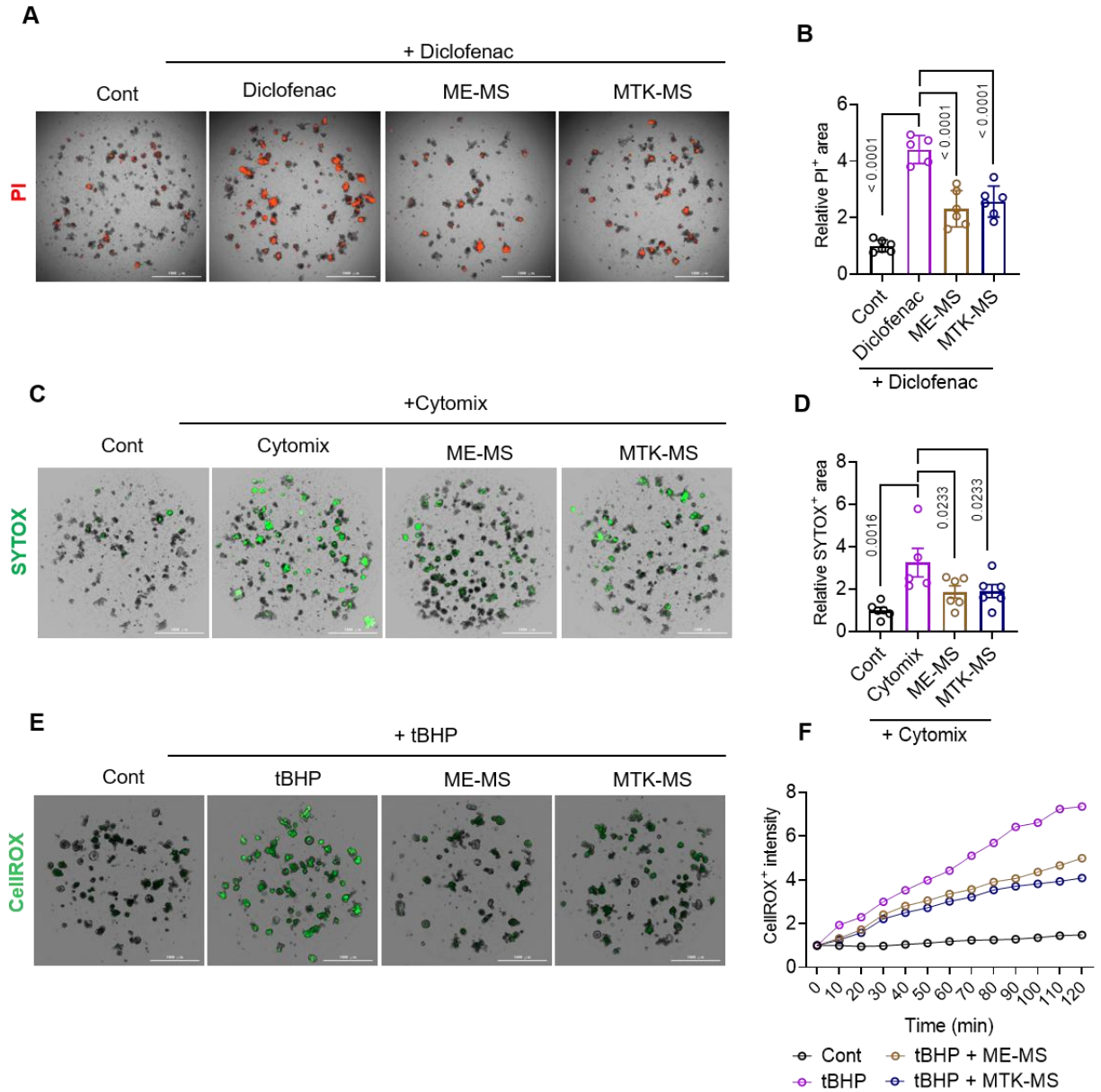


Figure S16. Protective effects of individual polymer-based microspheres on intestinal organoids. **A**, Propidium iodide (PI) staining depicts diclofenac-induced IO damage. Scale bar: 1000 μ m. **B**, PI-positive area of IO measurement (n = 5-6). **C**, Cytomix-induced IOs damage stained with SYTOX. Scale bar: 1000 μ m. **D**, SYTOX-positive area of IO measurement (n = 5-6). **E**, CellROX staining showing oxidative stress-induced damage in IOs treated with tBHP. Scale bar: 1000 μ m. **F**, Time course of CellROX intensity in IOs. Data in (**B**, **D**) are presented as mean

$\pm$ s.d. Data in (**B**, **D**) are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown).

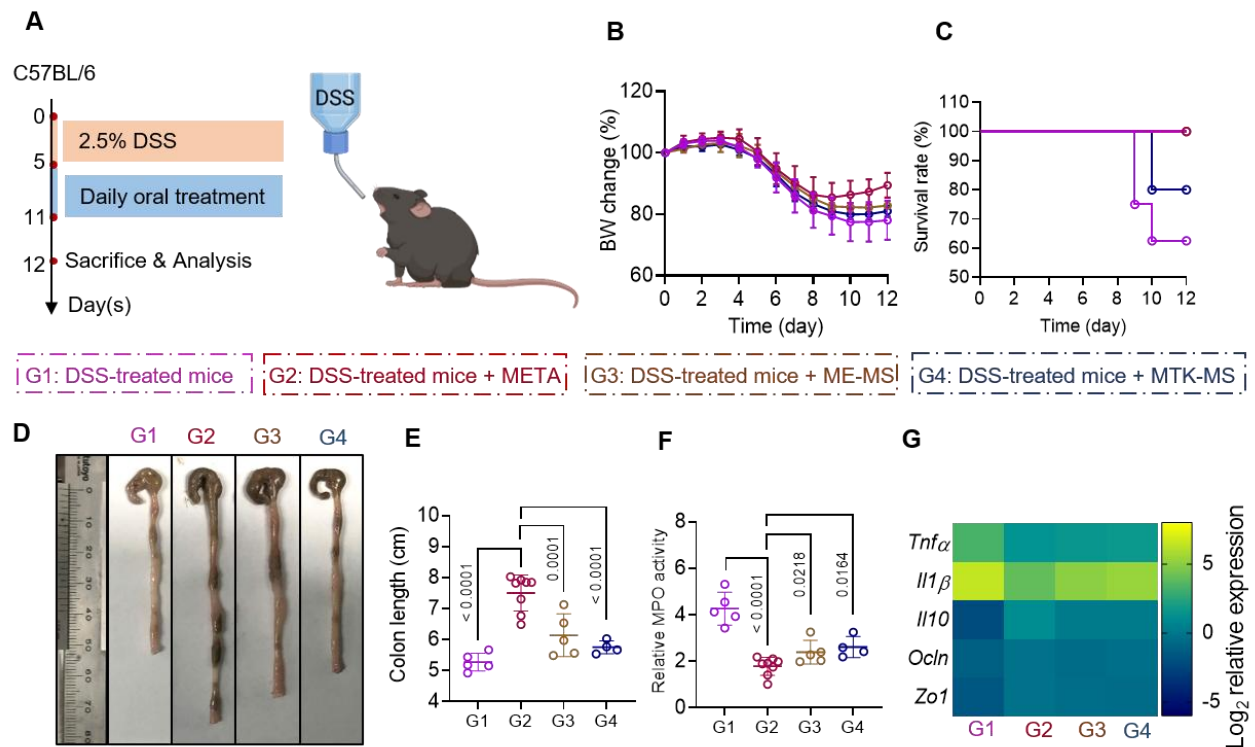


Figure S17. Superior therapeutic effects of META on DSS-induced colitis mice. **A**, Induction of the mouse colitis model through oral DSS delivery and subsequent drug treatment. G1: DSS-treated mice, G2: DSS-treated mice + META, G3: DSS-treated mice + ME-MS, G4: DSS-treated mice + MTK-MS. **B**, Body weight change measurement (n = 4-8 mice). **C**, Survival rate of mice (n = 5-8 mice). **D**, Representative images of colon tissues. **E**, Colon length measurement (n = 4-8 mice). **F**, Myeloperoxidase (MPO) activity in colon tissues analysis (n = 4-8 mice). **G**, Heat map of gene expression in colon tissues (n = 3-5 mice). Data of **G1 (DSS-treated mice)** and **G2 (DSS-treated mice + META)** in this figure, including in (**B**, **C**, **E**, **F**, **G**) were extracted from the main Figure 7 due to the same *in vivo* trial induction. Data are presented as mean ± s.d. Data are analyzed using one-way ANOVA with Tukey's multiple comparisons test (significant p values are shown). Figure S17A was created with BioRender.com.

Supplementary tables

Table S1. Clinical trials of oral melatonin for ulcerative colitis (2005-2025)

Trial	Year	Population	Treatment	Dosage form	Reference
Shahrokh et al.	2021	Mild- moderate UC (n=30)	3 mg/day oral for 3 months + standard therapy	Free melatonin	[5]
Chojnacki et al.	2011	Patients with left-sided UC (n=30)	Oral melatonin (5 mg/day) with mesalazine	Free melatonin	[6]
NCT00790478 Pilot	2008- 2012	Mild- moderate UC	Oral melatonin as an adjunctive	Free melatonin	ClinicalTrials.gov

Only 3 clinical trials were identified (all oral free melatonin, no modified delivery systems).

Doses: 3-5 mg/day (standard human dose). All adjunctive therapy with standard UC treatments.

No nanoparticle/microsphere melatonin clinical trials in UC patients have been found.

Table S2. Formulations used in this study

Objectives	Formulations
Loading capacity (LC%)	$LC\% = \frac{\text{Amount of drug loaded in MS}}{\text{Actual amount of MS}} \times 100\%$
Entrapment efficiency (EE%)	$EE\% = \frac{\text{Amount of drug loaded in MS}}{\text{Initial amount of drug added}} \times 100\%$
Similarity factor f_2	$f_2 = 50 \times \log \left\{ \left[1 + \frac{1}{n} \sum_{t=1}^n (R_t - T_t)^2 \right]^{-0.5} \times 100 \right\}$
Transepithelial electrical resistance (TEER)	$TEER = (R_m - R_i) \times A \text{ (ohm.cm}^2\text{)}$

where n is the number of time points, R_t and T_t are the dissolution values of different formulations at time point t , respectively. R_m is the resistance of the transmembrane, R_i is the intrinsic resistance of the cell medium, and A is the Transwell membrane surface area. MS: Microsphere

Table S3. Disease activity index criteria

Parameters	Criteria	Score
Weight loss	0-1%	0
	1-5%	1
	5-10%	2
	10-20%	3
	> 20%	4
Rectal bleeding	Negative	0
	Minor - moderate	2
	Severe	4
Stool consistency	Hard	0
	Soft	2
	Diarrhea	4

Table S4. Bacterial primer sequences used in RT-PCR analysis

Bacterial genera	Primer	Nucleotide sequence
<i>Total bacteria</i>	F	5'-AGAGTTTGATCATGGCTCAG-3'
	R	5'-ACCGCGACTGCTGCTGGCAC-3'
<i>Muribaculaceae</i>	F	5'-AGGCCCCGGTTCATACAGCTC-3'
	R	5'-GCGAATCCAAGTGCTGTCTGG-3'
<i>Lactobacillaceae</i>	F	5'-AGCAGTAGGGAATCTTCCA-3'
	R	5'-CACCGCTACACATGGAG-3'
<i>Akkermansiaceae</i>	F	5'-CAGCACGTGAAGGTGGGGAC-3'
	R	5'-CCTTGCGGTTGGCTTCAGAT-3'
<i>Bacteroidaceae</i>	F	5'-GGTTCTGAGAGGAGGTCCC-3'
	R	5'-GCTGCCTCCCGTAGGAGT-3'
<i>Clostridiaceae</i>	F	5'-AAATGACGGTACCTGACTAA-3'
	R	5'-CTTTGAGTTTCATTCTTGCGAA-3'
<i>Enterobacteriaceae</i>	F	5'-GTGCCAGCAGCCGCGGTAA-3'
	R	5'-GCCTCAAGGGCACAACCTCCAAG-3'
<i>Shigella</i>	F	5'-CAAGTCCGTAAATTCATTCTCTCTT-3'
	R	5'-GCTGGAAAACTCAGTGCCTCCC-3'
<i>Odoribacter</i>	F	5'-ATGTAATGATGAGCACTCTAACGG-3'
	R	5'-GGCTTTTGAGATTGGCATCC-3'
<i>Turicibacter</i>	F	5'-ACGGGGACAACGATTGGAAA-3'
	R	5'-AGTGATGCCAGGAGCATCTT-3'
<i>Prevotellaceae</i>	F	5'-GGTGTCGGCTTAAGTGCCAT-3'
	R	5'-CGGACGTAAGGGCCGTGC-3'

Table S5. Primer sequences used in RT-PCR analysis

Target gene	Species	Primer	Nucleotide sequence
<i>Actb</i>	Mouse	F	5'-CAT TGC TGA CAG GAT GCA GAA GG-3'
		R	5'-TGC TGG AAG GTG GAC AGT GAG G-3'
<i>Gapdh</i>	Mouse	F	5'-ACC ACA GTC CAT GCC ATC AC-3'
		R	5'-TCC ACC ACC CTG TTG CTG TA-3'
<i>Aanat</i>	Mouse	F	5'-CCA GTG CGT TTG AGA TTG AGC G-3'
		R	5'-AAG GCA GCC TTC CTC AAA CCA G-3'
<i>Hiomt</i>	Mouse	F	5'-GCA GCC TCC TGC TCT ACC TG-3'
		R	5'-ACC TGT AGA TGG CGG TGA AGG-3'
<i>Tph1</i>	Mouse	F	5'-TGT TGA CTG CGA CAT CAG CCG A-3'
		R	5'-GGA AAC CAA GGG ACA GTC TCC A-3'
<i>Tnfa</i>	Mouse	F	5'-GGT GCC TAT GTC TCA GCC TCT T-3'
		R	5'-GCC ATA GAA CTG ATG AGA GGG AG-3'
<i>Il1β</i>	Mouse	F	5'-TGG ACC TTC CAG GAT GAG GAC A-3'
		R	5'-GTT CAT CTC GGA GCC TGT AGT G-3'
<i>P65</i>	Mouse	F	5'-TCC TGT TCG AGT CTC CAT GCA G-3'
		R	5'-GGT CTCA TAG GTC CTT TTG CGC-3'
<i>Cox2</i>	Mouse	F	5'-GCG ACAT ACT CAA GCA GGA GCA-3'
		R	5'-AGT GGT AAC CGC TCA GGT GTT G-3'
<i>Il6</i>	Mouse	F	5'-TAC CAC TTC ACA AGT CGG AGG C-3'
		R	5'-CTG CAA GTG CAT CAT CGT TGT TC-3'
<i>Il10</i>	Mouse	F	5'-CGG GAA GAC AAT AAC TGC ACC C-3'
		R	5'-CGG TTA GCA GTA TGT TGT CCA GC-3'
<i>iNOS</i>	Mouse	F	5'-GAG ACA GGG AAG TCT GAA GCA C-3'
		R	5'-CCA GCA GTA GTT GCT CCT CTT C-3'
<i>Tgfβ</i>	Mouse	F	5'-TGA TAC GCC TGA GTG GCT GTC T-3'
		R	5'-CAC AAG AGC AGT GAG CGC TGA A-3'
<i>Ocln</i>	Mouse	F	5'-TGG CAA GCG ATC ATA CCC AGA G-3'
		R	5'-CTG CCT GAA GTC ATC CAC ACT C-3'
<i>Zo1</i>	Mouse	F	5'-GTT GGT ACG GTG CCC TGA AAG A-3'
		R	5'-GCT GAC AGG TAG GAC AGA CGA T-3'
<i>Ecad1</i>	Mouse	F	5'-GGT CAT CAG TGT GCT CAC CTC T-3'
		R	5'-GCT GTT GTG CTC AAG CCT TCA C-3'
<i>Emr1</i>	Mouse	F	5'-CGT GTT GTT GGT GGC ACT GTG A-3'
		R	5'-CCA CATC AGT TGT TCC AGG AGA C-3'
<i>Stat3</i>	Mouse	F	5'-AGG AGT CTA ACA ACG GGC AGC CT-3'
		R	5'-GTG GTA CAC CTC AGT CTC GAA G-3'
<i>Gapdh</i>	Human	F	5'-GTC GGA GTC AAC GGA TTT GG-3'
		R	5'-GGG TGG AAT CAT TGG AAC AT-3'
<i>Nrf2</i>	Human	F	5'-CAC ATC CAG TCA GAA ACC AGT GG-3'
		R	5'-GGA ATG TCT GCG CCA AAA GCT G-3'
<i>Hol</i>	Human	F	5'-CCA GGC AGA GAA TGC TGA GTT C-3'

		R	5'-AAG ACT GGG CTC TCC TTG TTG C-3'
<i>Ocln</i>	Human	F	5'-ATG GCA AAG TGA ATG ACA AGC GG-3'
		R	5'-CTG TAA CGA GGC TGC CTG AAG T-3'
<i>Cldn4</i>	Human	F	5'-AGT GCA AGG TGT ACG ACT CGC T-3'
		R	5'-CGC TTT CAT CCT CCA GGC AGT T-3'
<i>Zo1</i>	Human	F	5'-GTC CAG AAT CTC GGA AAA GTG CC-3'
		R	5'-CTT TCA GCG CAC CAT ACC AAC C-3'
<i>Ecad1</i>	Human	F	5'-GCC TCC TGA AAA GAG AGT GGA AG-3'
		R	5'-TGG CAG TGT CTC TCC AAA TCC G-3'

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