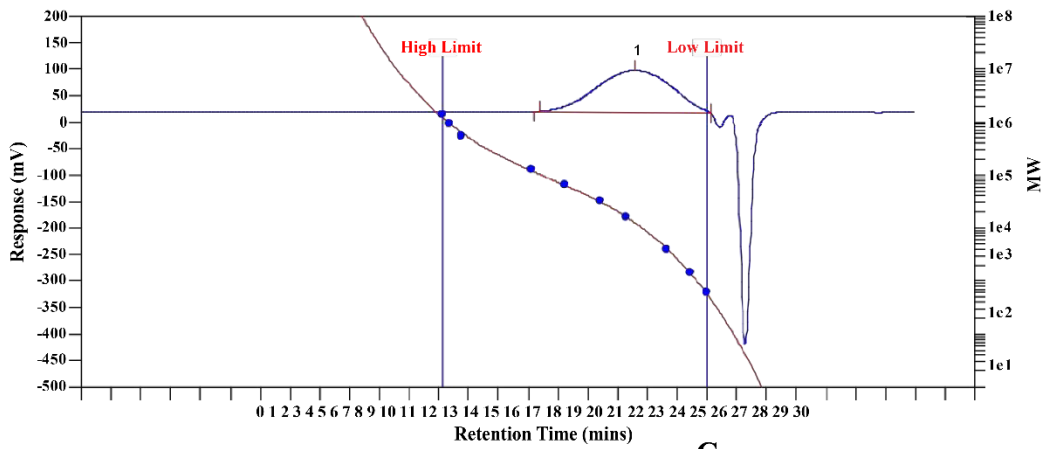
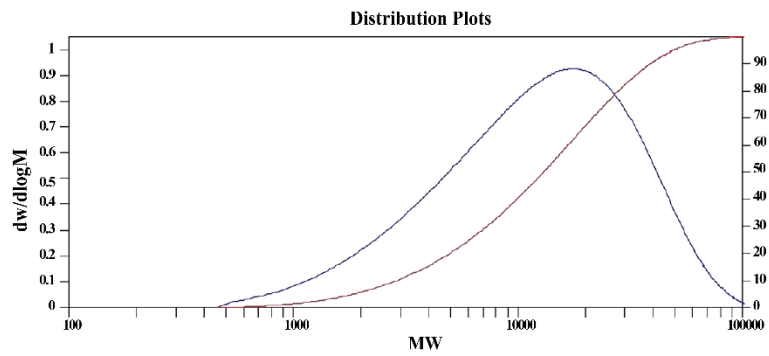


1 Supplementary Tables and Figures

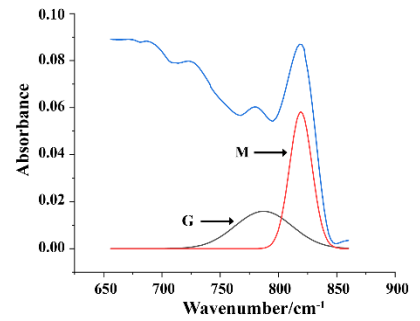
A



B



C



2

3 **Supplemental Figure S1. Characterization of sodium alginate (ALG).** (A) GPC

4 Chromatography of ALG. (B) The distribution plots of molecular weight of ALG. (C) Absorption

5 spectra of ALG by infrared spectra. M and G represent the peak absorption of β -D-mannuronic

6 acid and α -L-guluronic acid, respectively.

7 **Supplemental Table S1. Parameters and criteria of histological damage evaluation^{1,2}**

Parameters	Score	Histological features
	0	No change
(1) Loss of epithelial surface	1	Localized and mild
(2) Destruction of the crypts	2	Localized and moderate
(3) Infiltration of inflammatory cells	3	Localized and severe
	4	Extensive and moderate
	5	Extensive and severe

8 ¹ Each parameter is scored separately on a scale of 0-5. The histological score is the sum of the scores from all three parameters.

9 ² The method was adopted from a previous study with modifications (1).

10

Supplemental Table S2. Scoring of the disease activity index (DAI)^{1,2}

Score	Weight loss (%)	Stool consistency	Fecal blood content
0	None	Normal	Normal
1	0-10		
2	11-15	Loose stool	Occult blood
3	16-20		
4	> 20	Diarrhea	Hemorrhage/Gross bleeding

11 ¹ The DAI score is defined as the sum of three scores from weight loss (%), stool consistency, and fecal blood content.

12 ² The method was adopted from a previous study (2).

Supplemental Table S3. Primers used in this study¹

Target gene	Forward primer (5'-3')	Reverse primer (5'-3')	Annealing temp. (°C)	References
For host genes of mouse				
IL-1 β	CTCGCAGCAGCACATCAACAAG	GGAAGGTCCACGGGAAAGACAC	60	(3)
TNF- α	TGGGAGTAGACAAGGTACAACCC	CATCTTCTCAAAATTCGAGTGACAA	60	(3)
IL-6	ACCACGGCCTTCCTACTT	CACAACCTCTTTTCTCATTCCAC	60	(3)
FXR	TGGGCTCCGAATCCTCTTAGA	TGGTCCTCAAATAAGATCCTTGG	60	(4)
TGR5	CCTGGCAAAGCCTCATCGTC	AGCAGCCCCGGCTAGTAGTAG	60	(4)
β -actin	TGGAATCCTGTGGCATCCATGAAAC	TAAAACGCAGCTCAGTAACAGTCCG	60	(3)
For bacteria				
Total bacteria	GTGSTGCAYGGYYGTCGTCA	ACGTCRTCCMCNCCTTCCTC	60	(5)
<i>baiJ</i>	TCAGGACGTGGAGGCGATCCA	TACRTGATACTGGTAGCTCCA	60	(6)
<i>bsh</i>	ATGGGCGGACTAGGATTACC	TGCCACTCTCTGTCTGCATC	54	(7)
<i>7α-HSDH</i>	GGGTATTGTGTATCAAAAGCTGCGG	TCCGTTGCTATAAGCCCAGGTAAGA	60	This study; MH743112.1
<i>7β-HSDH</i>	GTCGTAAGCAGACTTTTCGCTGC	TTGATCATTGCCTCATGCTTTTCC	60	This study; KF052988.1

14 ¹ IL-6, interleukin-6; IL-1 β , interleukin-1 β ; TNF- α , tumor necrosis factor- α ; FXR, farnesoid X receptor; TGR5, G-coupled protein
15 receptor; *baiJ*, genes encoding bile salt 7 α -dehydroxylase; *bsh*, genes encoding bile salt hydrolase; 7 α -HSDH, genes encoding 7 α -
16 hydroxysteroid dehydrogenase; 7 β -HSDH, genes encoding 7 β -hydroxysteroid dehydrogenase.

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