

Supplementary materials

Figure S1. MS/MS spectra and fragmentation patterns of selected phenolic components.

Table S1. Disease activity index (DAI) scoring criteria.

Table S2. Primer sequence for quantitative PCR.

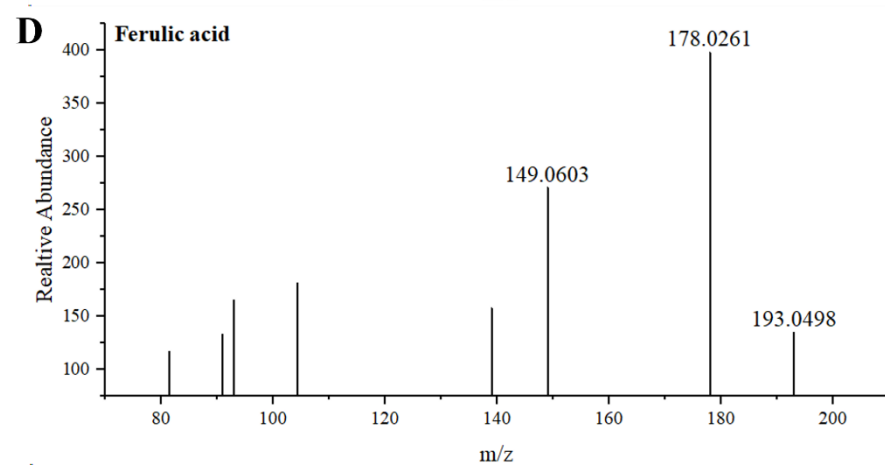
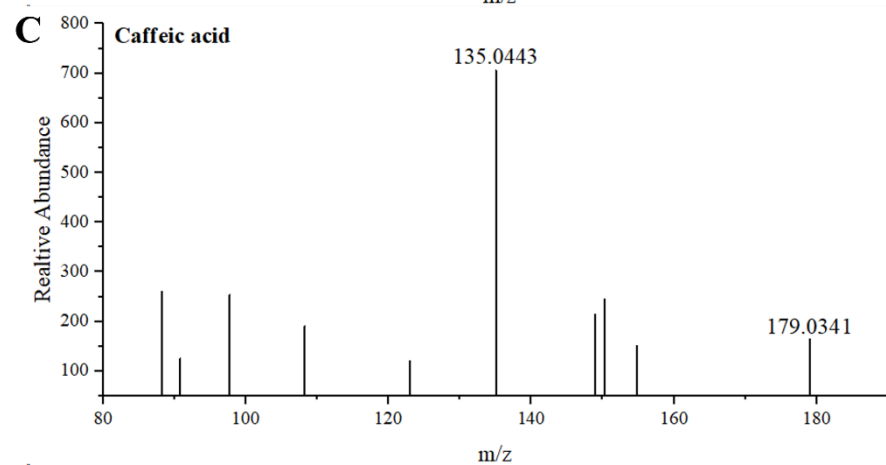
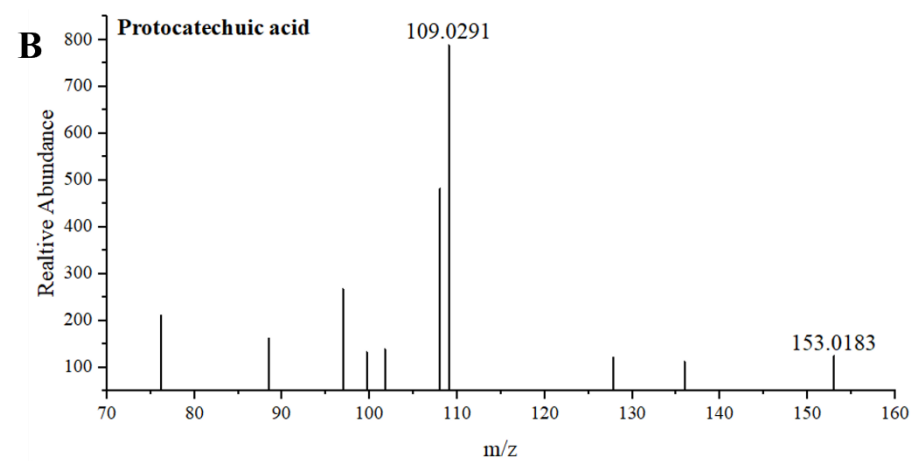
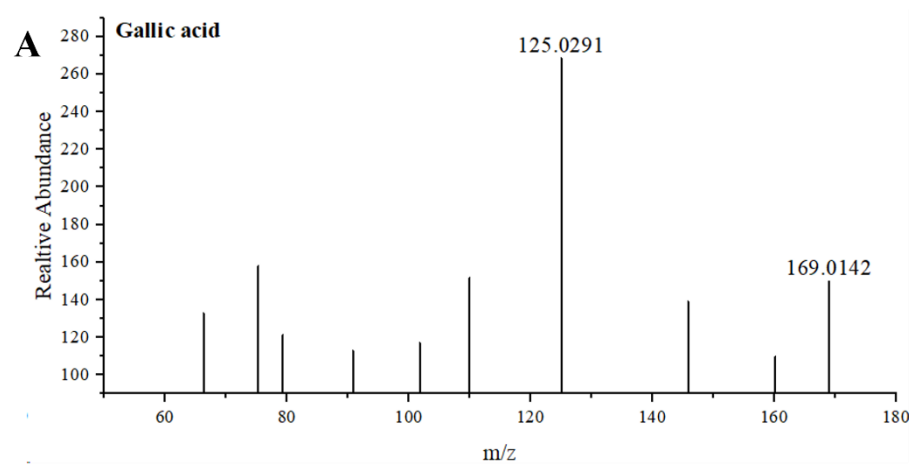


Figure S1. MS/MS spectra and fragmentation patterns of selected phenolic components. (A) Gallic acid; (B) Protocatechuic acid; (C) Caffeic acid; (D) Ferulic acid.

Table S1. Disease activity index (DAI) scoring criteria.

Body weight loss (%)	Stool consistency	Retal bleeding	Activity states	Score
No change	Normal	No observable blood	Normal	0
1-5	Loose stool	Small amount of blood in some stool	Some signs of poor condition	1
5-10	Mild diarrhea	Blood in stool regularly seen	Moderately poor condition	2
≥ 10	Diarrhea	Blood in all stool	Bad condition	3

Table S2. Primer sequence for quantitative PCR.

Gene	Primer	Sequence (5'-3')
β -actin (Mouse)	Forward	ATGGAGCCGGACAGAAAAGC
	Reverse	TGGGAGGTGTCAACATCTTCTT
ZO-1 (Mouse)	Forward	CCAGAGCCTCAGAAACCTCA
	Reverse	GCAGGAAGATGTGCAGAAGG
Occludin (Mouse)	Forward	GCGGAAAGAGTTGACAGTCC
	Reverse	TGCCTGAAGTCATCCACACT
Claudin-1 (Mouse)	Forward	ACGGTCTTTGCACTTTGGTC
	Reverse	GGGAGAGGAGAAGCACAGTT
I κ B- α (Mouse)	Forward	TGAAGGACGAGGAGTACGAGC
	Reverse	TGCAGGAACGAGTCTCCGT
NF- κ B p65 (Mouse)	Forward	TGTTCCAAAGAGCACCCAGG
	Reverse	TGCCAGGTCTGTGAACACTC
ERK (Mouse)	Forward	GGTTGTTCCCAAATGCTGACT
	Reverse	CAACTTCAATCCTCTTGTGAGGG
JNK (Mouse)	Forward	GTGGAATCAAGCACCTTCACT
	Reverse	TCCTCGCCAGTCCAAAATCAA
MAPK p38 (Mouse)	Forward	TGACCCTTATGACCAGTCCTTT
	Reverse	GTCAGGCTCTTCCACTCATCTAT