

Table S1. Characterization of probable major metabolites of CPE55 by UPLC/Q-TOF-MS/MS.

No.	Rt (min)	Compound	Probable formula	Measured [M-H] ⁺ (m/z)	Representative fragmentation	References
1	0.91	Phosphatidic acid + glycerol + palmitic acid	C ₂₉ H ₃₉ O ₆	483	136.03 , 137.03, 152.02, 348.05 , 349.05	
2	1.44	gamma-Glutamylisoleucine	C ₁₁ H ₂₀ N ₂ O ₅	260.99	144.92, 189.92, 216.05	
3	1.67	Inosine	C ₁₀ H ₁₂ N ₄ O ₅	268.07	136.03, 137.03	
4	2.72	Methyl stearate	C ₁₉ H ₃₈ O ₂	298.07	136.03 , 146.03, 188.04, 299.07,	
5	4.97	Gluconic acid	C ₆ H ₁₂ O ₇	197.08	105.04, 133.07, 161.06, 179.07 , 251.00 ,	
6	7.28	4,7-Dihydroxy-3-butylphthalid/iso-mer methyl and-O-GluA	C ₁₉ H ₂₄ O ₁₀	413,16	127.00, 139.00, 403.17, 412.66	[1-8],
7	10.29	Lyso-PC (18:3)	C ₂₆ H ₄₈ NO ₇ P	518.32	104.07 , 124.97, 184.04	
8	10.97	Lyso-PC(18:2)	C ₂₆ H ₅₀ NO ₇ P	520.33	104.07 , 124.97, 184.04 , 502.32	
9	12.14	Malvidin-3-O-cis-caffeoyglucoside	C ₂₈ H ₃₁ O ₁₄ N ₂ Cl	655.32	184.04, 449.18, 535.26, 563.26, 595.29, 596.29,	
10	13.83	Naringenin-O-glucoside-O-glucuronide	C ₂₇ H ₃₀ O ₁₆	609	431.17, 447.20 , 519.23, 547.23, 579.26	

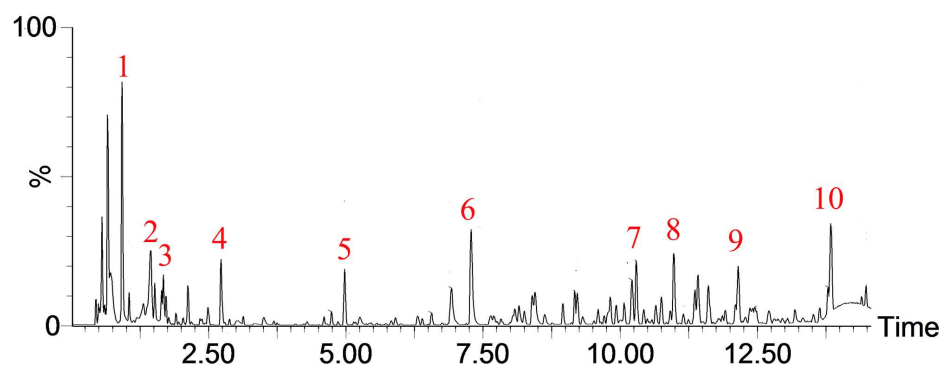


Figure S1. Chromatographic peaks of *Chlorella pyrenoidosa* ethanol extract in UPLC.

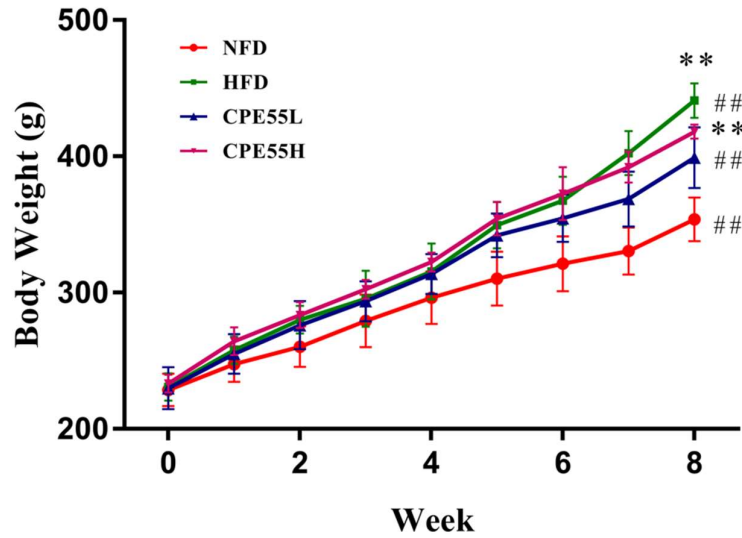


Figure S3. Effect of CPE55 on body weight of high-fat-diet rats during the experimental period. NFD: normal fat diet; HFD: high-fat diet; CPE55L: 150 mg/(kg-day) *C. pyrenoidosa* 55% ethanol extract; CPE55H: 300 mg/(kg-day) *C. pyrenoidosa* 55% ethanol extract; WK: week. NFD group, rats fed NFD and gavaged with 150 mg/(kg-day) normal saline. HFD group, rats fed HFD and gavaged with 150 mg/(kg-day) normal saline. CPE55L group, rats fed HFD and gavaged with 150 mg/(kg-day) *C. pyrenoidosa* 55% ethanol extract in water. CPE55H group, rats fed HFD and gavaged with 300 mg/(kg-day) *C. pyrenoidosa* 55% ethanol extract in water. The differences were assessed by ANOVA and denoted as follows: * $p < 0.05$ versus the NFD group, * $p < 0.05$ versus the HFD group, ** $p < 0.01$ versus the NFD group, and ** $p < 0.01$ versus the HFD group.

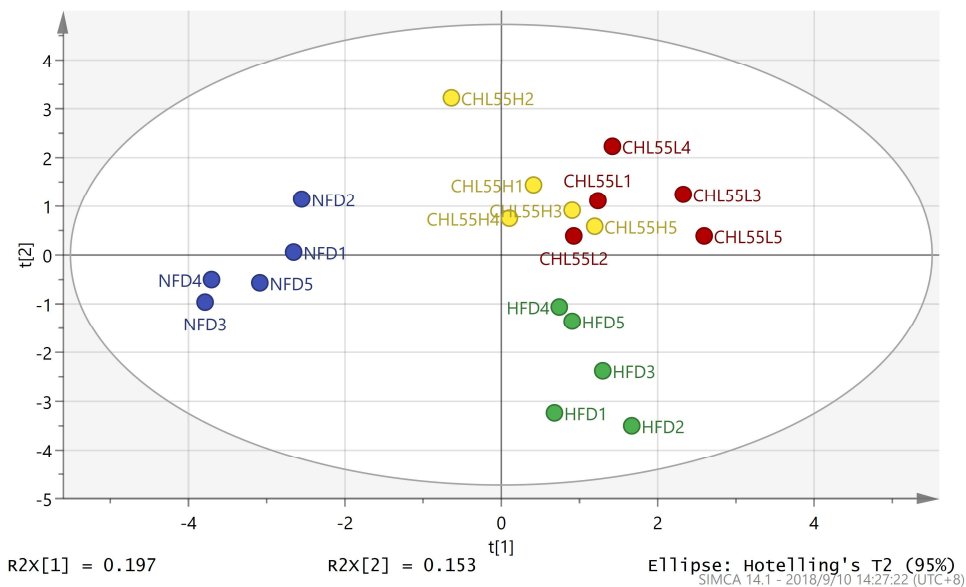


Figure S4. Principal component analysis plots of rat caecal microbiota coloured by diet. Five rats were randomly selected from each experimental group for analysis of caecal microbiota.

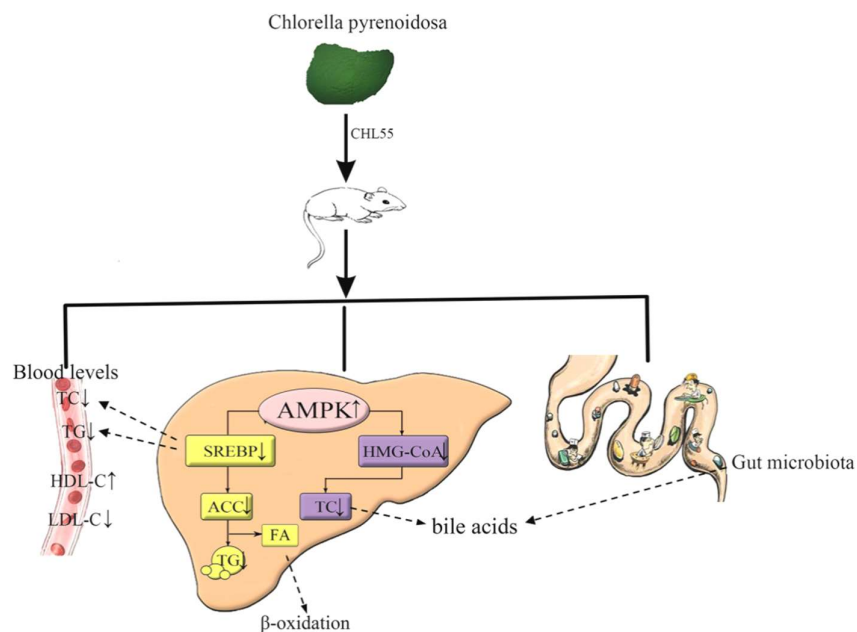


Figure S5. Summary of the mechanism of CPE55 to prevent LMD. Note: FA (fatty acids), stimulatory modification (solid arrow), and indirect modification (dotted arrow).

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