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8 The optimal concentration of CLT in Raw 264.7 cells. (A) Figure S2 Detected cell viability to screen appropriate intervention concentrations by the 9 CCK8 method. (B) Immunoblot analyses and (C) quantitative determination of iNOS 10 11 in the LPS-induced Raw 264.7, set the CLT concentration gradient to 0, 0.125, 0.25, 12 and 0.5 µmol/L. Statistical differences between treatment groups and CON group 13 were analyzed by student's t-test analysis. All data is represented as mean \pm SD, n=3.***: p < 0.001, vs. CON. #: p < 0.05, ##: p < 0.01, vs. LPS. ns: p > 0.05. 14 15



Figure S3 Morphological changes in Raw 264.7 cells. Treatment of LPS-induced Raw 264.7 cells without or with CLT for 24 h. Representative light microscopy images are presented. Images were captured at ×200 magnification.



Figure S4 Constructing an ECM1 overexpression Raw 264.7 cell line. (A) The mRNA expression of ECM1 in each group of cells; (B) The protein expression level of ECM1 in each group of cells; (C) Quantitative determination of ECM1 protein expression; Statistical differences in multiple groups were determined by one-way ANOVA followed by Tukey's multiple comparisons. All data is represented as mean \pm SD, n=3. **: p<0.01, ***: p < 0.001, vs. CON. ns: p > 0.05.

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Table S1 Primers List 32 primer sequences Gene F-iNOS ATCTTGGAGCGAGTTGTGGATTGTC **R-iNOS** TAGGTGAGGGCTTGGCTGAGTG F-Arg1 AGACAGCAGAGGAGGTGAAGAGTAC R-Arg1 AAGGTAGTCAGTCCCTGGCTTATGG F-IL6 CTTCTTGGGACTGATGCTGGTGAC R-IL6 TCTGTTGGGAGTGGTATCCTCTGTG F-TNF-a CGCTCTTCTGTCTACTGAACTTCGG R-TNF-a GTGGTTTGTGAGTGTGAGGGTCTG GCGAGCTTCCATATCCAGAACAGG F-ECM1 R-ECM1 TCAGGAGACAGGTCACAGCAGAG

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