

Figure S1. Method of administration. dioscin (80 mg/kg body weight) or vehicle (CMC-Na) was orally administered to C57BL/6 mice every day after saline or CS injection (n=10 per group).

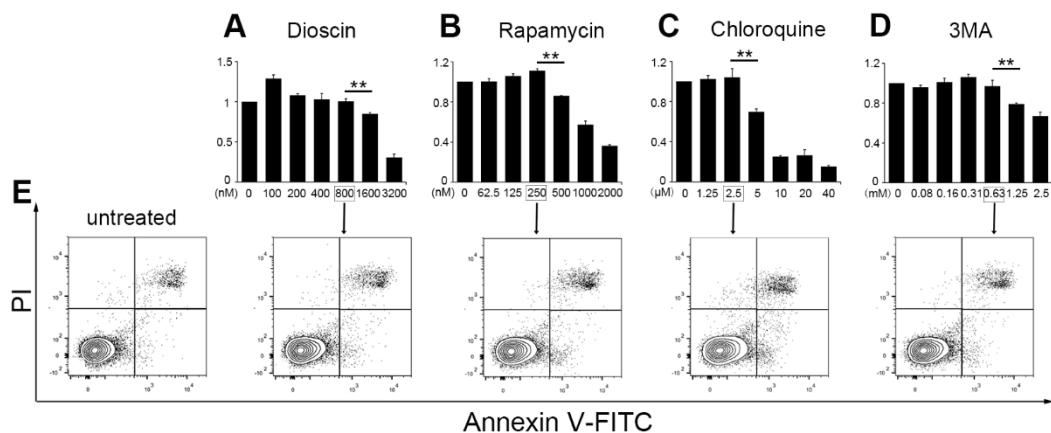


Figure S2. Effects of chemicals on cell viability. (A) MTT assay of dioscin-treated cells. (B) MTT assay of Rapamycin-treated cells. (C) MTT assay of Chloroquine-treated cells. (D) MTT assay of 3MA-treated cells. (E) Apoptosis assay of drugs used in experiment. *, P<0.05; **, P<0.01. Error

bars indicate mean \pm SEM. All experiments were repeated three times.

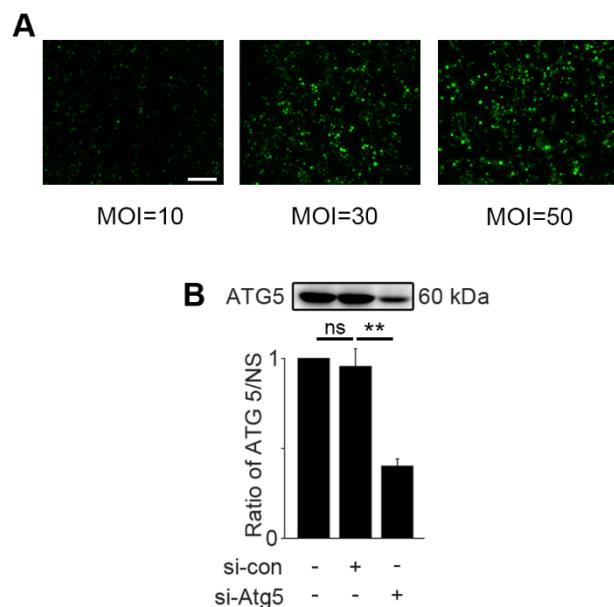


Figure S3. Transfection efficiency of lentivirus with Atg5 siRNA sequence.

(A) Multiplicity of Infection (MOI) gradient of mRFP-GFP-LC3 lentivirus infecting MH-S cells at 24 hours post infection, scale bar indicates 100 μ m. (B) Transfection efficiency was tested by measuring ATG5 by immunoblot assay (n=3). *, $P<0.05$; **, $P<0.01$. Error bars indicate mean \pm SEM. All experiments were repeated three times.

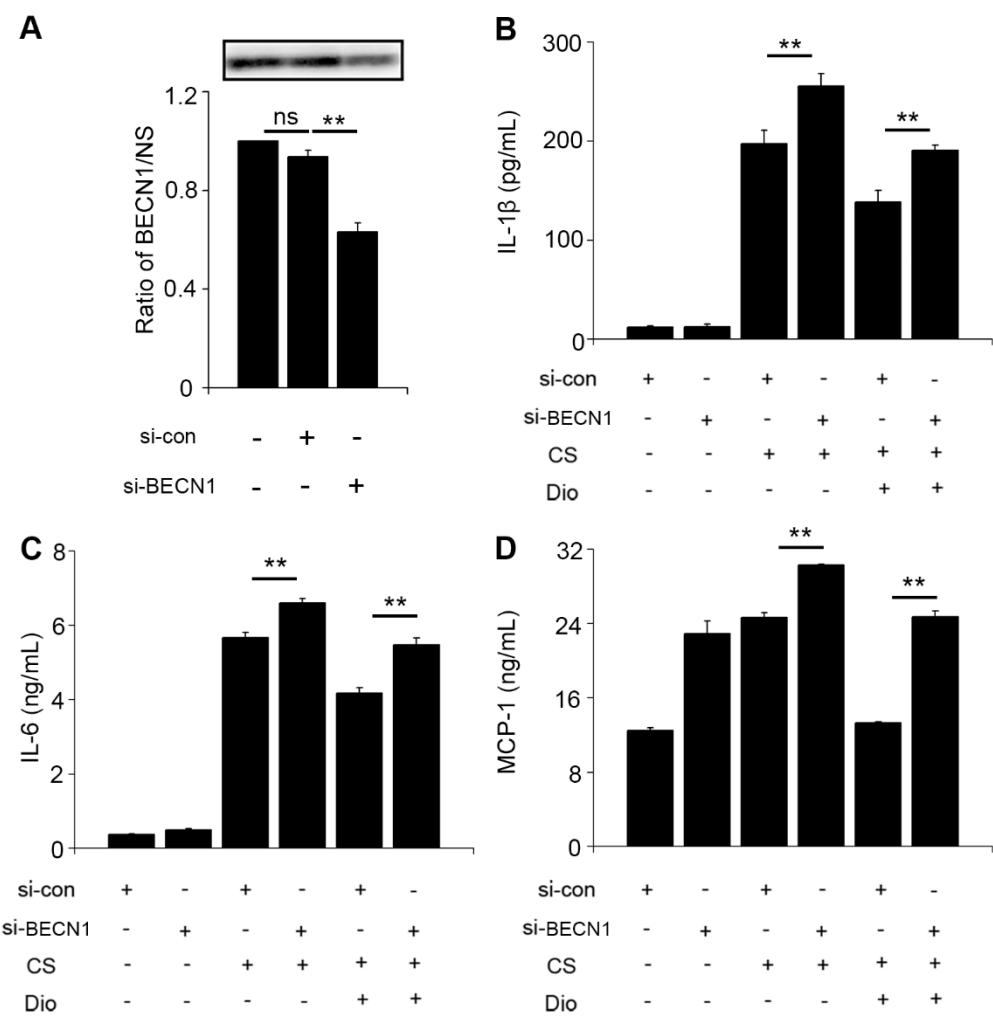


Figure S4. BECN1 knockdown could affect function of dioscin.

(A) Transfection efficiency was tested by measuring Beclin1 by immunoblot assay (n=3) (B-D) ELISA analysis of IL-1 β , IL-6, and MCP-1 in MH-S cell supernatant (n=3). (B-G). *, P<0.05; **, P<0.01. Error bars indicate mean \pm SEM. All experiments were repeated three times with similar results.

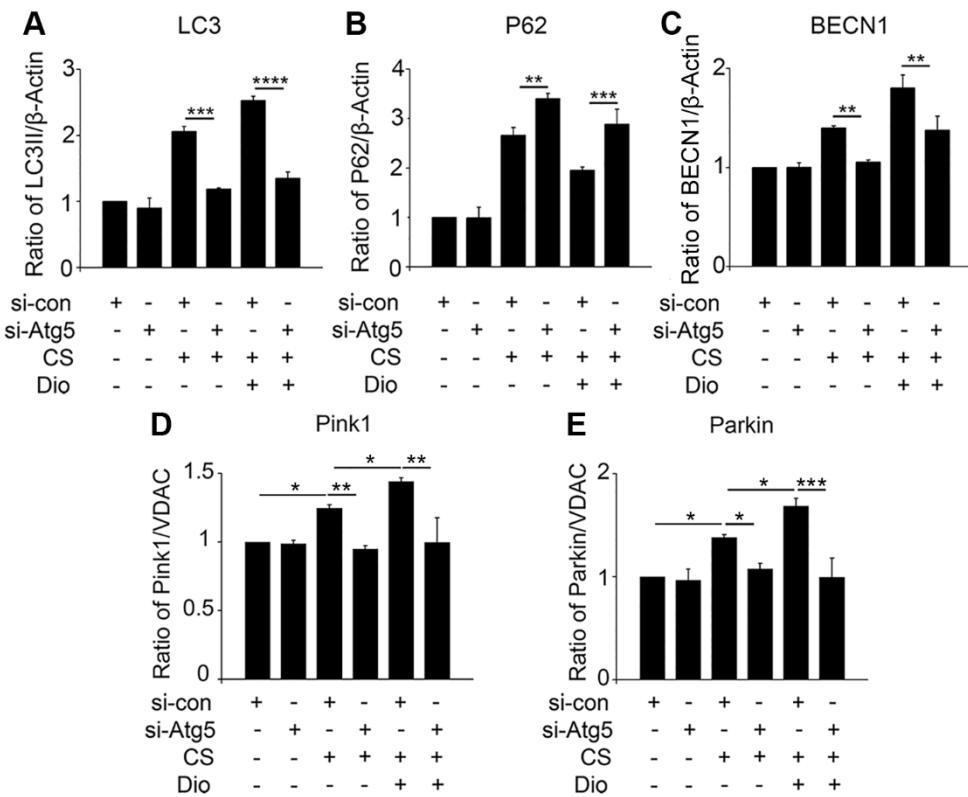


Figure S5. Quantification of autophagy and mitophagy protein. (A) Quantification of LC3II. (B) Quantification of P62. (C) Quantification of BECN1. (D) Quantification of Pink1. (E) Quantification of Parkin. * , $P<0.05$; ** , $P<0.01$; *** , $P<0.001$, **** , $P<0.0001$. Error bars indicate mean \pm SEM. All experiments were repeated three times.

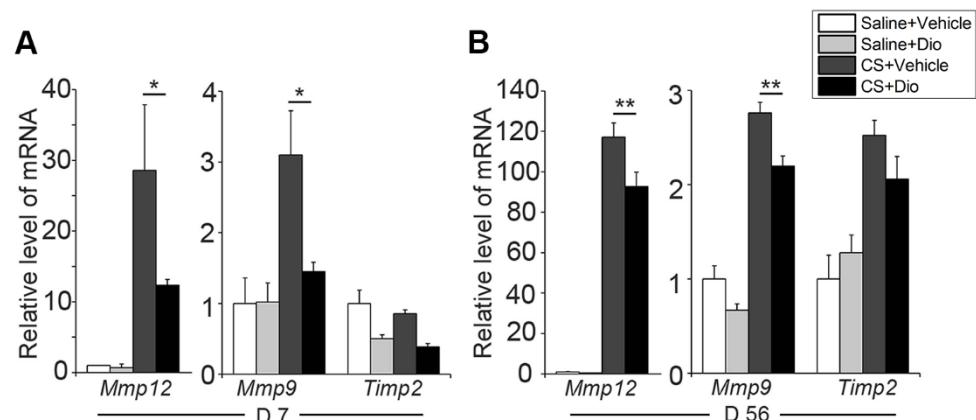


Figure S6. Dioscin treatment reduced the production of MMPs and TIMP

in gene level. **(A)** mRNA level of *Mmp12*, *Mmp9*, and *Tim2* in lung tissue at day 7 (n=5). **(B)** mRNA level of *Mmp12*, *Mmp9*, and *Tim2* in lung tissue at day 56 (n=5). *, P<0.05; **, P<0.01. Error bars indicate mean ± SEM. All experiments were repeated three times.

Table S1. Primer sequences for qPCR in this study.

Gene	Forward (5'-3')	Reverse (5'-3')
<i>BECN1</i>	ATGGAGGGGTCTAAGGCGTC	TCCTCTCCTGAGTTAGCCTCT
<i>P62</i>	AGGATGGGGACTTGGTTGC	TCACAGATCACATTGGGGTGC
<i>Atg5</i>	TGTGCTTCGAGATGTGTGGTT	GTCAAATAGCTGACTCTTGGCAA
<i>Atg7</i>	GTTCGCCCCCTTAATAGTGC	TGAACCTCAACGTCAAGCGG
<i>ULK1</i>	TGGAGGTGGCCGTCAAATG	CGCATAGTGTGCAGGTAGTC
<i>Mmp9</i>	CTGGACAGCCAGACACTAAAG	CTCGCGGCAAGTCTTCAGAG
<i>Mmp12</i>	CTGCTCCCATGAATGACAGTG	AGTTGCTTCTAGCCCAAAGAAC
<i>Il-1b</i>	GCAACTGTTCCCTGAACTCAACT	ATCTTTGGGGTCCGTCAACT
<i>Mcp-1</i>	TTAAAAAACCTGGATCGGAACCAA	GCATTAGCTTCAGATTACGGGT
<i>Il-6</i>	TAGTCCTTCCTACCCCAATTCC	TTGGTCCTTAGGCCACTCCCTC
<i>Col1a1</i>	GCTCCTCTTAGGGGCCACT	CCACGTCTCACCATTGGGG
<i>Col3a1</i>	CTGTAACATGGAAACTGGGGAAA	CCATAGCTGAACTGAAAACCACC