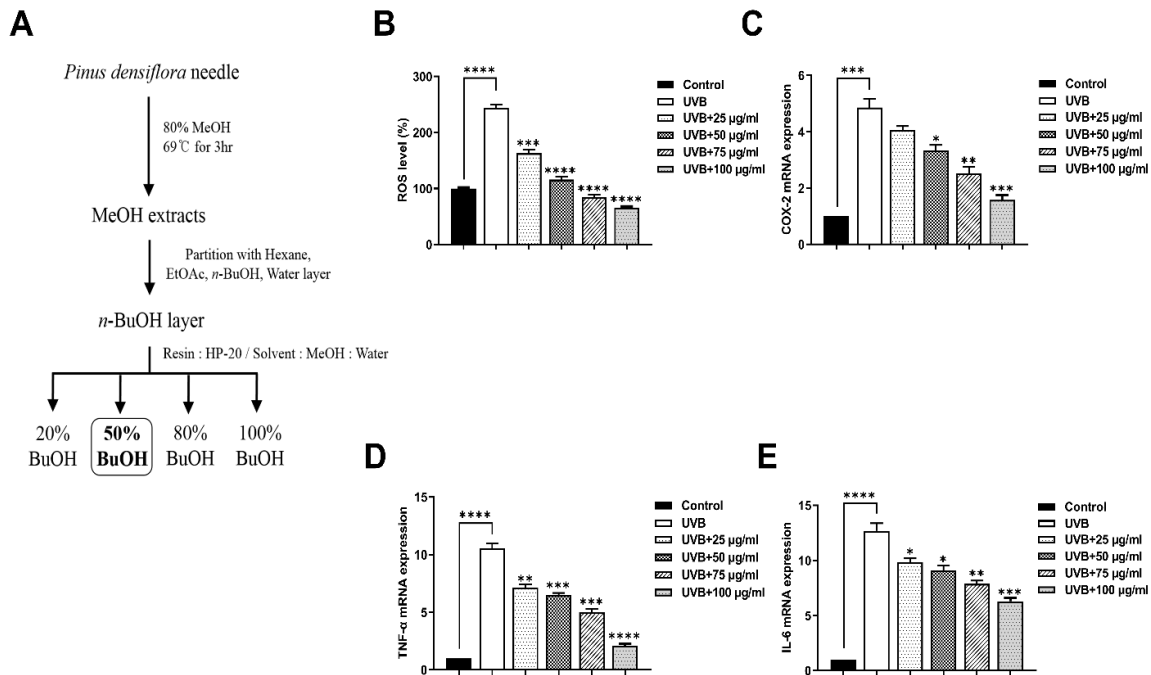


## Supplementary Information

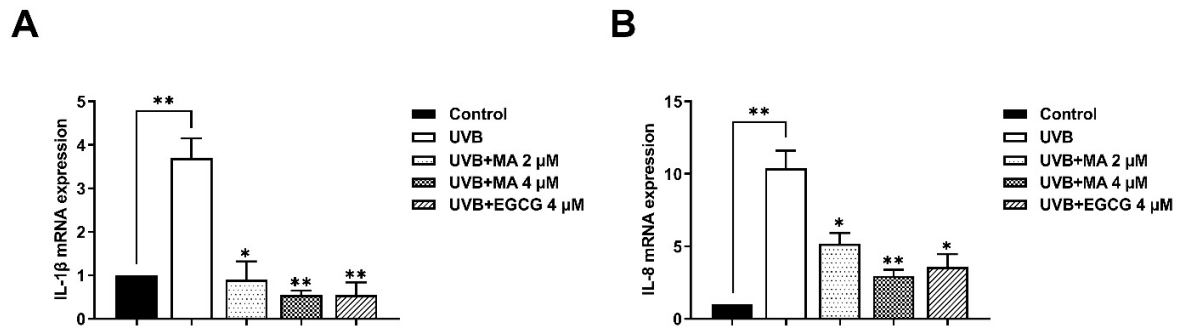
Supplementary Table S1. Primer lists

Genes		sequences
Nrf-2	Sense	5'- CAG CGA CCT TCG CAA ACA AC -3'
	Anti-sense	5'- CAT GAT GAG CTG TGG ACC GT -3'
SOD-1	Sense	5'- ATG GCG ACG AAG GCC GTG TG-3'
	Anti-sense	5'- GAC CAC CAG TGT GCG GCC AA-3'
HO-1	Sense	5'- AGG GAA TTC TCT TGG CTG GC-3'
	Anti-sense	5'- GAC AGC TGC CAC ATT AGG GT-3'
MMP-1	Sense	5'- TCT TGG ACT CTC CCA TTC TAC T-3'
	Anti-sense	5'- TAG CTT ACT GTC ACA CGC TTT T-3'
MMP-3	Sense	5'- TGA GGA CAC CAG CAT GAA CC-3'
	Anti-sense	5'- ACT TCG GGA TGC CAG GAA AG-3'
MMP-9	Sense	5'- GGG CCG CTC CTA CTC TGC CT-3'
	Anti-sense	5'- TCG AGT CAG CTC GGG TCG GG-3'
NF-kB	Sense	5'- TCT TGG ACT CTC CCA TTC TAC T-3'
	Anti-sense	5'- TAG CTT ACT GTC ACA CGC TTT T-3'
P65	Sense	5'- TGA GGA CAC CAG CAT GAA CC -3'
	Anti-sense	5'- ACT TCG GGA TGC CAG GAA AG-3'
P50	Sense	5'- GGG CCG CTC CTA CTC TGC CT-3'
	Anti-sense	5'- TCG AGT CAG CTC GGG TCG GG-3'
TNF- $\alpha$	Sense	5'- AGC TCC CTC TAT TTA TGT TTG C-3'
	Anti-sense	5'- TAC ATG GGA ACA GCC TAT TGT-3'
COX-2	Sense	5'- TCC CAC AGT CAA AGA TAC TCA G-3'
	Anti-sense	5'- CTG TTT AAG CAC ATC GCA TAC T-3'
IL-6	Sense	5'- TCG AGC CCA CCG GGA ACG AAA-3'
	Anti-sense	5'- AGG CAA CTG GAC CGA AGG CG-3'
IL-8	Sense	5'- CCC CAA TAA ATA TAG GAC TGG A-3'
	Anti-sense	5'- GCT GCT TTC ACA CAT GTT ACT C-3'
IL-1 $\beta$	Sense	5'- CAC CTC TCC TAC TCA CTT AAA GC-3'
	Anti-sense	5'- GGC TCT TTT ACA GAC ACT GCT AC-3'
Smad3	Sense	5'-GGC TGC TCT CCA ATG TCA -3'
	Anti-sense	5'- CAC TCT GCG AAG ACC TCC-3'
COL1A1	Sense	5'-TCT GAC TGG AAG AGT GGA GAG TA -3'
	Anti-sense	5'- GTT CTT GCT GAT GTA CCA GTT CT-3'
COL3A1	Sense	5'- CCT GAA CTC AAG AGT GGA GAA TA-3'
	Anti-sense	5'- CTA GAA TCT GTC CAC CAG TGT TT-3'
GAPDH	Sense	5'- TCA ACG GAT TTG GTC GTA TT-3'
	Anti-sense	5'- GGA TTT CCA TTG ATG ACA AG-3'

## Supplementary Figure legends



**Supplementary Figure S1. The 50% *n*-BuOH fraction, which exhibits strong antioxidant activity.** (A) *n*-BuOH layer fractionated using HP-20 column chromatography and a gradient of MeOH:water (from 20:80 to 100:0), to yield four combined fractions. (B-E) HaCaT cells are pretreated for 4 h with the 50% BuOH layer and then irradiated with ultraviolet (UV) B (15 mJ/cm<sup>2</sup>). (B) The levels of reactive oxygen species (ROS) are estimated indirectly by measuring the fluorescence emitted by 2', 7'-dichlorodihydrofluorescein diacetate (DCF-DA), (C-E) mRNA expression of cyclooxygenase-2 (*cox-2*), tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) and interleukin-6 (*il-6*) are evaluated by real-time polymerase chain reaction (PCR). Data are expressed as the mean  $\pm$  the standard error of the mean (SEM). (\*)  $P < 0.05$ , (\*\*)  $P < 0.01$ , (\*\*\*)  $P < 0.001$ , and (\*\*\*\*)  $P < 0.0001$ .



**Supplementary Figure S2 (related to Figure 3C). Effects of malonic acid (MA) on ultraviolet (UV) B-induced proinflammatory mediator expression levels in HaCaT cells. (A-B)** HaCaT cells are pretreated with MA for 4 h and then exposed to UVB irradiation (15 mJ/cm<sup>2</sup>). After overnight UVB exposure, interleukin-1 $\beta$  (*il-1 $\beta$* ) (**A**) and *il-8* (**B**) expression are evaluated by real-time polymerase chain reaction (PCR). Data are expressed as the mean  $\pm$  the standard error of the mean (SEM). (\*)  $P < 0.05$ , (\*\*)  $P < 0.01$ , (\*\*\*)  $P < 0.001$ , and (\*\*\*\*)  $P < 0.0001$ .