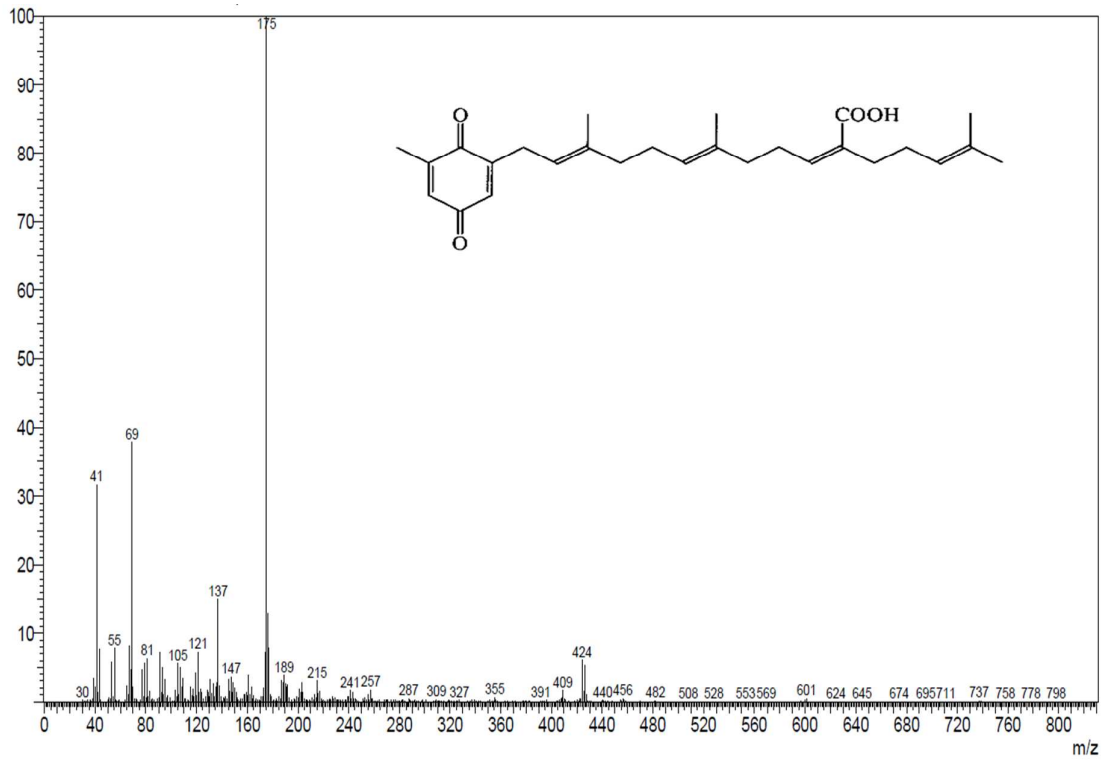
~~Supplementary~~ Table 1.

Primer	Sequence
ICAM-1	sense, 5’-ACC ATC TAC AGC TTT CCG GC-3’ antisense, 5’-GGA GAG CAC ATT CAC GGT CA-3’
VCAM-1	sense, 5’-GGC GCC TAT ACC ATC CGA AA -3’ antisense, 5’-TGC TTC TAC AAG ACT ATA TGA CCC C-3’
IL-8	sense, 5’-CTG TGT GAA GGT GCA GTT TTG C-3’ antisense, 5’-CAA CCC TCT GCA CCC AGT TT-3’
MCP-1	sense, 5’-CTG CTC ATA GCA GCC ACC TT-3’ antisense, 5’-ACA GGA TGT CTG GGG AAA GC-3’
GAPDH	sense, 5’-GAC CCC TTC ATT GAC CTC AA-3’ antisense, 5’-CTT CTC CAT GGT GGT GAA GA-3’

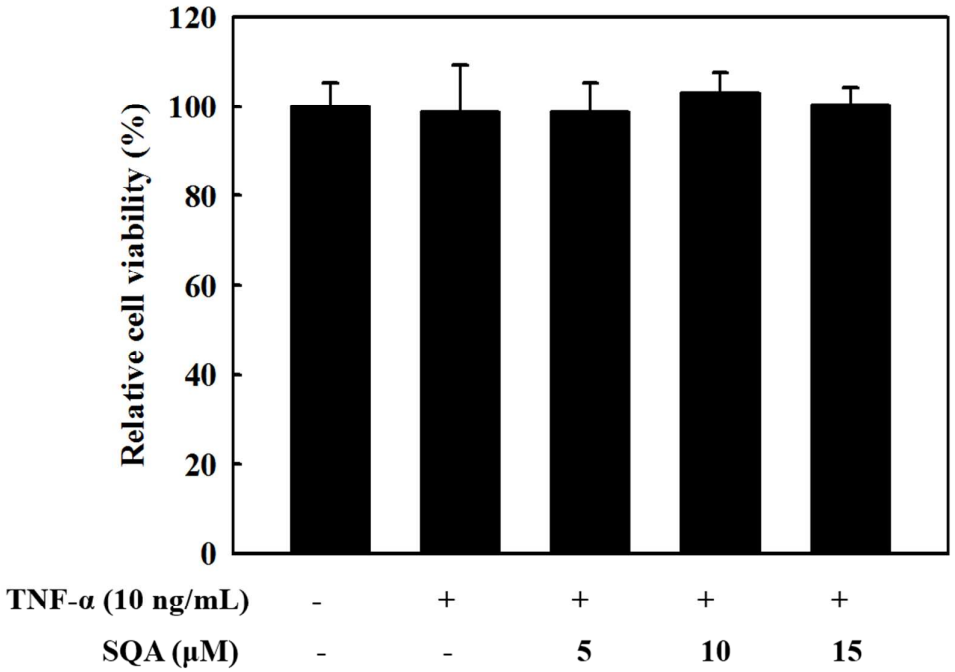
~~Supplementary~~ Table 1. Oligonucleotide primer used for RT-PCR

~~Supplementary Figure 1.~~



~~Supplementary Figure 1.~~ Mass Spectrometry Analysis of SQA, *m/z* 424.

~~Supplementary Figure 2.~~



~~Supplementary Figure 2.~~ Effect of SQA on cell viability in HUVECs. Cytotoxic effect of SQA was measured by using a 3-(4, 5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium (MTS) assay. Cells pretreated with different concentrations (5, 10, and 15 μM) of SQA for 1 h were stimulated with TNF- α (10 ng/mL) for 24 h. The values are the mean ± S.D. of three independent experiment.