

A Sterol from Soft Coral Induces Apoptosis and Autophagy in MCF-7 Breast Cancer Cells

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Figure	Caption	Page
S1	Inhibition effects of compound 1 relative to DMSO control on the cell proliferation in MDA-MB-231 cells. Cells were treated with compound 1 at indicated concentrations for 24 h or 48 h, and cell proliferation was determined by MTT assay. Points, means; bar, S.D. (n = 4-6). ** <i>p</i> < 0.01.	3
S2	Statistically analysis of cell cycle after the treatment of compound 1 in MDA-MB-231 cells. Cells were treated with compound 1 at the indicated concentrations for 48 h, followed by PI staining and flow cytometric analysis. Treatment with etoposide (ETO) at 10 μM was used as the positive control. Points, mean; bars, S.D. (n = 4). * <i>p</i> < 0.05, ** <i>p</i> < 0.01.	4

S3	Expression of p62 and LC3B in compound 1-treated MDA-MB-231 breast cancer cells for 48 h.	5
S4	<p>Left, Western blot analyses of LC3B in compound 2 or compound 3 or 50 μM troglitazone (TRO) for 48 h in MCF-7 cells. Three independent experiments were performed, and the fold changes of LC3B-II/β-actin is presented in right panel.</p> <p>Points, means; bars, S.D. (n = 3).</p>	6

Supplementary materials and methods

Inhibition effects of compound 1 relative to DMSO control on the cell proliferation in MDA-MB-231 cells.

Cells (5×10^3) were treated with compound 1 or DMSO in 5% FBS-supplemented DMEM/F12 medium in 96-well plates. After 24 h, the medium was removed, replaced by 200 μ L DMEM/F12 containing 0.5 mg/mL of MTT, and cells were incubated in the CO₂ incubator at 37°C for 2 h. Supernatants were aspirated from the wells, and the reduced MTT dye was solubilized in 200 μ L/well DMSO. Absorbance at 570 nm was measured using a plate reader.

Cell cycle analysis of compound 1 in MDA-MB-231 cells

Cells ($2 \times 10^5/3$ mL) were treated with DMSO or compound 1 for 48 h and washed twice with PBS, followed by staining with propidium iodide (PI) and analyzed by flow cytometry and the multicycler (ModFitLT 3.0) software program.

Western blot analysis

Western blot analysis was performed as previously report (Weng et. al, (2017) *Phytother. Res.*, 31, 1722-1730).

Figure S1

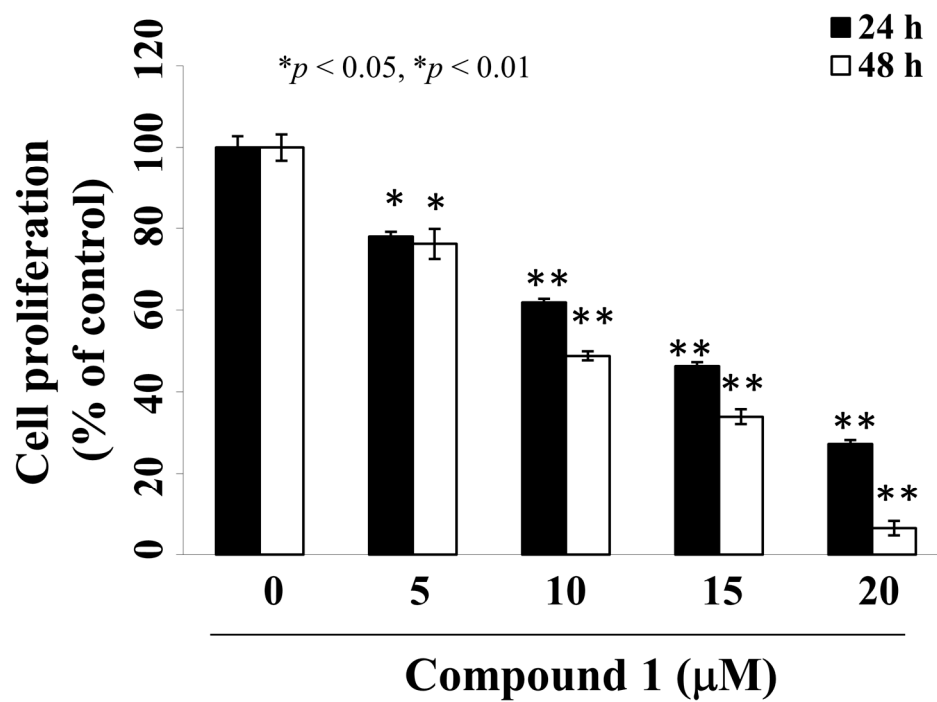


Figure S2

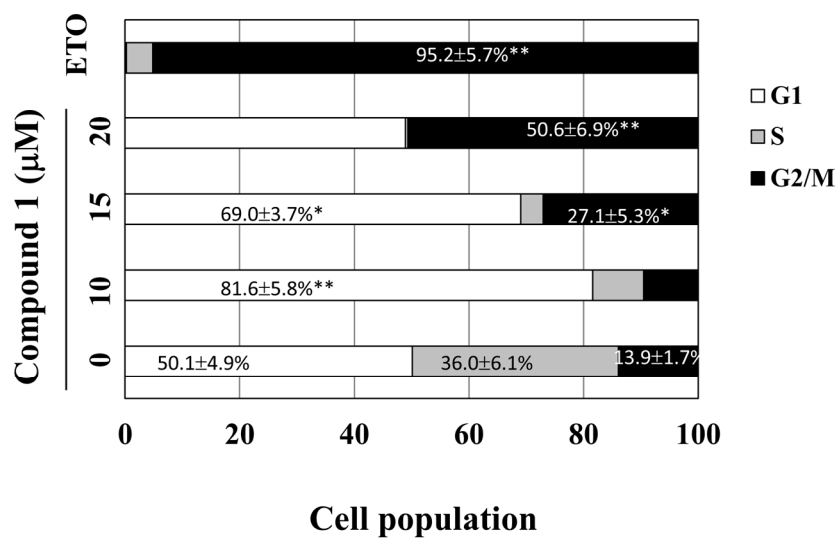


Figure S3

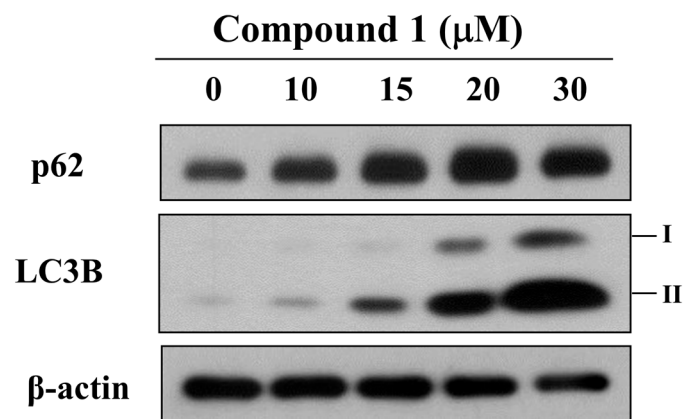


Figure S4

