ERK activating peptide, AES16-2M promotes wound healing through accelerating migration of keratinocytes.

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Supplementary Materials and Methods

Cell proliferation assays.

HaCaT cells (3×10^3 cells/well, 200 µl) were plated into a 96-well plate. After an overnight incubation, media was removed and cells were treated with 200 µl fresh media that included AES16-2M or EGF. Every 24 hr, for 72 hr, 10 µl Cell Counting Kit-8 (CCK-8) (Dojindo, Japan) was added to each well and the plate was incubated for 3 hr at 37°C. Absorbance at 450 nm was measured with an ELISA microplate reader.

Chemicals and antibodies.

The p38 inhibitor, SB203580, was purchased from Calbiochem. Rabbit anti-p38, anti-phosphorylated (p-)p38 (all at 1:1000; 9212, 9211) were purchased from Cell Signaling.

Supplementary figure legends

Figure S1. AES16-2M had an insignificant effect on HaCaT cell proliferation.

HaCaT cells (3×10^3 cells/well) were cultured in 96-well plates with the indicated doses of AES16-2M or EGF (100 ng/ml). Cell proliferation was detected with CCK-8 analysis at 24, 48, and 72 hr. The relative proliferation rate was calculated as the ratio of the AES16-2M or EGF group versus the PBS group at each time point. (*n*=5) Error bars, mean ± SD. ***p*<0.05 (EGF treatment group versus control group), one-way ANOVA followed by Tukey post-hoc test.

Figure S2. HaCaT cell migration induced by AES16-2M was poorly associated with p38 activation.

(A) HaCaT cells were treated with AES16-2M (10 ng/ml), TGF- β (1 ng/ml), and PBS for 30min under serum-free conditions. The p38 inhibitor (SB203580 at 10 μ M) was added to cells 2 hr prior to treatment. Western blotting for phospho-p38, total-p38, and GAPDH was performed. The figure is a cropped image; full-length blots are presented in Figure S4. (B) The band intensity of panel A was measured with Image J and relative density was analysed as the ratio of phospho-p38 to GAPDH (*n*=3). Error bars, mean ± SD. **p*<0.05 (TGF- β or SB203580 treatment group versus control group), one-way ANOVA followed by Tukey post-hoc test. N.S.=No significance (AES16-2M versus AES16-2M + SB203580 at 10 μ M) 2 hr before AES16-2M treatment. Scale bar: 500 μ m. (D) The residual area of the wound was measured by Image J and the relative wound area was calculated as the ratio of the remaining area to that

at 0 hr (*n*=6). Error bars, mean \pm SD. ***p*<0.01, ****p*<0.001 (AES16-2M, AES16-2M + SB203580 or TGF- β treatment group versus control group). N.S.=No significance (AES16-2M versus AES16-2M + SB203580 treatment group), one-way ANOVA followed by Tukey posthoc test.

Figure S3. Full-length blots for Figure 4A.

All dashed lines indicate the area cropped and used in the figures. The number displayed is the size of the marker (kD). (A) phospho (p)-ERK. (B) ERK. (C) GAPDH. Blots B and C were performed following stripping the membrane shown in panel A. The arrow indicated nonspecific binding (bands remaining after stripping ERK from the membrane). (D) Digitized image for maker detection.

Figure S4. Full-length blots for Figure S2A.

All dashed lines indicate the area cropped and used in the figures. The number displayed is the size of the marker (kD). (A) phospho-p38. The arrow indicated non-specific binding. (B) p38. Blot B was performed following stripping the membrane shown in panel A. (C) Digitized image for marker detection of panels A and B. (D) GAPDH. (E) Digitized image for marker detection of panel D.

Figure S1



Figure S2



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Figure S3



Figure S4

