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17 **Keywords**: diabetic wound, keratinocyte, antimicrobial peptide, high glucose, angiogenesis

Abstract

19 Impaired keratinocyte functions are major factors that are responsible for delayed

20	diabetic wound healing. In addition to its antimicrobial activity, the antimicrobial peptide
21	derived from insulin-like growth factor-binding protein 5 (AMP-IBP5) activates mast
22	cells and promotes keratinocyte and fibroblast proliferation and migration. However, its
23	effects on diabetic wound healing remain unclear. Human keratinocytes were cultured in
24	normal or high glucose milieus. The production of angiogenic growth factor and cell
25	proliferation and migration were evaluated. Wounds in normal and streptozotocin-
26	induced diabetic mice were monitored and histologically examined. We found that AMP-
27	IBP5 rescued the high glucose-induced attenuation of proliferation and migration as well
28	as the production of angiogenin and vascular endothelial growth factor in keratinocytes.
29	AMP-IBP5-induced activity was mediated by the epidermal growth factor receptor,
30	signal transducer and activator of transcription 1 and 3, and mitogen-activated protein
31	kinase pathways, as indicated by the inhibitory effects of pathway-specific inhibitors. In
32	vivo, AMP-IBP5 markedly accelerated wound healing, increased the expression of
33	angiogenic factors and promoted vessel formation in both normal and diabetic mice.
34	Overall, the finding that AMP-IBP5 accelerated diabetic wound healing by protecting
35	against glucotoxicity and promoting angiogenesis suggests that AMP-IBP5 might be a
36	potential therapeutic target for treating chronic diabetic wounds.

1 Introduction

 Diabetes mellitus is a serious public health problem associated with long-term consequences that impacts the quality of life of individuals and their families. The prevalence of diabetes is increasing and is estimated to be 10.2% (578 million) by 2030 and 10.9% (700 million) by 41 2045.¹ The global prevalence of diabetic foot ulcers is as high as 6.3% due to the increasing 42 worldwide prevalence of diabetes and the prolonged life expectancy of diabetic patients.² It has been reported that every 30 seconds, the lower limb or part of a lower limb is amputated in 44 diabetic patients due to foot ulcers worldwide.³ Although a wide range of treatment strategies, such as customized dressings, negative pressure wound therapy, hyperbaric oxygen treatment, debridement, topical growth factor application and stem-cell therapy, have been proposed for diabetic wound therapy, the etiological complexity of impaired wound healing in patients with diabetes often leads to unsatisfactory results, as there is a lack of holistic strategies to resolve 49 this issue.⁴⁻⁶ Therefore, there is a high unmet need for the development of novel therapeutic strategies for the treatment of diabetic wounds.

 Impaired diabetic wound healing involves multiple factors, including hypoxia, epidermal cell dysfunction, impaired angiogenesis and neovascularization, infections, glucotoxicity, 53 decreased host immune resistance, and neuropathy.⁷⁻¹⁰ Keratinocytes play a crucial role in re- epithelialization and angiogenesis via migration, proliferation, and the secretion of cytokines 55 and antimicrobial peptides $(AMPs)$.^{11,12} Several hypotheses suggest that hyperglycemia-

- 58 important factors that contribute to poor diabetic wound healing.^{7,13,14}
- Antimicrobial peptide derived from insulin-like growth factor-binding protein 5 (AMP-IBP5) is a newly discovered AMP produced by the defined proteolytic processing of insulin-like growth factor-binding protein 5 (IGFBP5) via serine proteases such as prohormone convertases 62 and carboxypeptidase.¹⁵ Although AMPs were initially well known for their antimicrobial 63 activities, increasing evidence suggests that these molecules, such as human β -defensin (hBD)- 2, hBD-3 and cathelicidin LL-37, also exert multiple immunomodulatory effects, including the modulation of inflammation, promotion of cell proliferation and migration, induction of 66 angiogenesis, and improvements in skin barrier function and wound healing.^{8,16,17} LL-37 promotes keratinocyte migration through activation of the epidermal growth factor receptor 68 (EGFR) and signal transducer and activator of transcription (STAT) pathways.¹⁸ Furthermore, recent studies indicated that AMP-IBP5 induces the proliferation and migration of keratinocytes and fibroblasts, enhances the secretion of angiogenin (ANG) and vascular 71 endothelial growth factor (VEGF), and promotes the activation of mast cells, $19-21$ indicating that AMP-IBP5 might play an important role in the wound healing process.

 Although AMP-IBP5 is thought to contribute to wound healing, its potential effects on diabetic wounds and the underlying mechanisms remain unexplored. We hypothesized that

 AMP-IBP5 might counteract the negative effects of high glucose on keratinocytes and promote diabetic wound healing through the activation of EGFR/STAT and MAPK pathways as well as promote proliferation, migration and angiogenesis in keratinocytes.

2 Materials and methods

2.1 Reagents

 AMP-IBP5 (AVYLPNCDRKGFYKRKQCKPSR-NH2) was obtained from the Peptide Institute (Osaka, Japan). Rabbit anti-CD31 antibody was obtained from Abcam (Tokyo, Japan). IgG isotype control and antibodies against phosphorylated and unphosphorylated EGFR, STAT1, STAT3, ERK, JNK and p38 were purchased from Cell Signaling Technology (Beverly, MA). Goat anti-rabbit Alexa Fluor 594 was purchased from Invitrogen (Carlsbad, CA). Normal goat serum was obtained from Vector (Burlingame, CA). Mouse anti-Ki67 monoclonal antibody was obtained from Invitrogen. Streptozotocin was purchased from Sigma–Aldrich (St Louis, MO). AG1478 was obtained from Santa Cruz Biotechnology (Dallas, TX). Fludarabine and cryptotanshinone were purchased from Cayman Chemical (Ann Arbor, MI). U0126, SB203580 and JNK inhibitor II were obtained from Calbiochem (La Jolla, CA). Enzyme- linked immunosorbent assay (ELISA) kits were obtained from R&D Systems (Minneapolis, MN). Mitomycin C and crystal violet were obtained from the Fujifilm Wako Pure Chemical Corporation (Tokyo, Japan).

2.2 Mouse model and treatment

 cut and washed 3 times with PBS. The skin specimen was placed upside down on a transparent Petri dish for cell/tissue culture and macroscopically visualized for subcutaneous vascular formation.

2.4 Total RNA extraction and real-time quantitative PCR

 Total RNA was extracted from the skin tissue using the RNeasy Plus Universal Mini kit (Qiagen). cDNA was obtained using a ReverTra Ace qPCR RT kit (Toyobo, Osaka, Japan). Real-time PCR was performed with an Applied Biosystems StepOnePlus Real-time PCR system (Thermo Fisher Scientific, Waltham, MA) by using TaqMan Universal PCR Master Mix or SYBR Premix Ex Taq (Takara, Tokyo, Japan). TaqMan assay primers and probe mixes of the genes for murine *Ang* (Mm00833184_s1), *Egf* (Mm00438696_m1) and *Vegf* (Mm00437306_m1) were obtained from Applied Biosystems assays-on-demand. The data 142 were normalized to β-actin or endogenous RPS18 expression ($2^{-\Delta\Delta CT}$ method). The primer information for SYBR Premix Ex Taq is shown in Supplementary Table 1.

2.5 Culture of primary human keratinocytes

 Primary human epidermal keratinocytes from neonatal foreskin were purchased from Kurabo Industries (Osaka, Japan) and were cultured in HuMedia-KG2 (Kurabo Industries) as 147 described previously.²³ Cells were maintained at 37° C and serially passaged at 60-70% confluence. The normal glucose concentration in the human epidermis is close to that in plasma (5.8 mM) :²⁴ thus, we considered 6 mM to be normoglycemic. For hyperglycemic conditions, **2.6 ELISA**

 Keratinocytes were cultured under normal conditions or 38 mM glucose for 48 hours and 154 then incubated with 10 μ M AMP-IBP5 for 48 hours. The amounts of ANG, EGF and VEGF in the cultured supernatants were measured with appropriate kits. In some experiments, keratinocytes were pretreated with various inhibitors for 2 hours before stimulation, and ELISA quantification was performed as described above.

2.7 Proliferation and migration assay

 Keratinocyte proliferation was assessed using a 5-bromo-2'-deoxyuridine (BrdU) labeling and detection kit III (Roche Diagnostics, Indianapolis, IN) according to the manufacturer's protocol. Briefly, keratinocytes cultured in normal or HG conditions were trypsinized and 162 seeded into 96-well plates $(1 \times 10^4 \text{ cells/well})$. Cells were stimulated with 5 μ M AMP-IBP5 163 for 48 hours and incubated with 10 μ M BrdU for 4 hours. The colorimetric reaction product was measured using a microplate reader at a wavelength of 450 nm.

165 Keratinocytes were seeded into collagen I-coated 96-well plates $(0.5 \times 10^5 \text{ cells/well})$ and cultured for 3 hours. Keratinocyte monolayers were scratched using a 96-well wound marker 167 (Essen BioScience, Ann Arbor, MI). To exclude the influence of cell proliferation, $1 \mu g/ml$ 168 mitomycin C (Fujifilm, Tokyo, Japan) was added for 2 hours before stimulation with 10 μ M

2.8 Western blotting

 Equal amounts of protein extracts were fractionated by 12.5% SDS–PAGE. After protein transfer, the polyvinylidene difluoride membranes (Millipore, Billerica, MA) were incubated with appropriate antibodies, developed with the Luminata Forte Western HRP substrate (Millipore, Billerica, MA) and visualized using Fujifilm LAS-4000 Plus (Fujifilm, Tokyo, Japan). Densitometric analysis was performed using ImageJ.

2.9 Statistical analysis

 The statistical analysis was performed using either one-way ANOVA followed by the Dunnett's post hoc test for multiple groups or Student's *t* test for 2 groups (Prism 7, GraphPad Software, San Diego, CA). The results are presented as the means ± standard deviations. The number of independent experiments carried out is indicated as *n*. *P* < 0.05 was considered significant.

3 Results

3.1 IGFBP5 was downregulated in diabetic mouse skin and in keratinocytes cultured in

 To evaluate whether AMP-IBP5 plays a role in diabetic wounds, we investigated the expression of IGFBP5, the parent protein of AMP-IBP5. The expression of *Igfbp5* was significantly decreased by 54.1% in the skin tissues of diabetic mice compared with those of normal control mice (Figure 1A). In addition, pretreatment of keratinocytes with 38 mM glucose for 48 hours to mimic diabetic conditions resulted in the downregulation of *IGFBP5* expression by 74.7% compared with that of the vehicle control (Figure 1B). Therefore, AMP- IBP5 might be reduced in the skin of diabetic subjects. **3.2 AMP-IBP5 promoted wound healing in diabetic mice** To determine the effect of AMP-IBP5 on *in vivo* wound healing, AMP-IBP5 or 0.01% acetic

 acid (vehicle) was topically applied to the wound area. AMP-IBP5-treated wounds exhibited accelerated wound closure compared with vehicle-treated controls in both normal and diabetic mice (Figure 2A). In control mice, AMP-IBP5-treated wounds started to significantly heal on day 4 and were completely healed at day 12, whereas in diabetic mice, the effect of AMP-IBP5 was first observed on day 8, and the wounds were completely healed at day 16 (Figure 2A, B). Of note, we observed that spontaneous wound healing was complete on day 16 in normal mice, while in diabetic mice, this was observed on day 20 postinjury (data not shown). Histological comparison of wounds further confirmed delayed wound closure in diabetic mice and remarkable promotion of wound healing by AMP-IBP5 in both normal and diabetic mice (Figure 2C, D). This finding demonstrated that AMP-IBP5 effectively accelerated delayed wound healing in diabetic mice.

3.3 AMP-IBP5 rescued HG-induced impairments in angiogenesis

214 Impaired angiogenesis plays a critical role in the pathogenesis of diabetic wound healing.²⁶ We investigated the expression of angiogenic growth factors in wound tissues obtained at day 4 posttreatment. AMP-IBP5-treated wound tissues showed enhanced expression levels of *Ang*, *Egf* and *Vegf* (Figure 3A). Interestingly, in diabetic mice, vehicle-treated wounds exhibited markedly reduced expression levels of *Ang*, *Egf* and *Vegf*, and this attenuated expression was ameliorated in AMP-IBP5-treated wound tissues. The expression of other growth factors, such as fibroblast growth factor, transforming growth factor and platelet-derived growth factor, was not observed (data not shown).

 The ability of AMP-IBP5 to rescue HG-induced angiogenesis impairment in diabetic mice was further confirmed by *in vitro* experiments. In keratinocytes cultured under normoglycemic conditions, AMP-IBP5 induced the production of ANG and VEGF, while an effect on EGF was not observed. The spontaneous secretion of ANG and VEGF was significantly reduced under HG conditions, and the addition of AMP-IBP5 restored this secretion (Figure 3B). This finding suggests that AMP-IBP5 improves neovascularization in diabetic wounds.

 Indeed, immunofluorescence staining of CD31, a marker of endothelial cells, showed that AMP-IBP5-treated wounds displayed increased numbers of CD31-positive blood vessels compared with vehicle-treated wounds in both normal and diabetic mice at day 4 postinjury (Figure 3C). Compared with wounds in normal mice, diabetic wounds showed decreased numbers of blood vessels in both the vehicle-treated and AMP-IBP5-treated groups. Similar data were observed by immunohistochemical staining with CD31 antibody (Figure S1). Furthermore, macroscopic observation of vessels in wounds also revealed that treatment with AMP-IBP5 increased the number of vessels in both normal and diabetic mice compared with vehicle treatment (Figure 3D). Collectively, AMP-IBP5 may promote neovascularization in diabetic subjects.

3.4 AMP-IBP5 attenuated HG-induced inhibition of keratinocyte proliferation and migration

 Both keratinocyte proliferation and migration are indispensable in the wound repair process. The hyperglycemic milieu resulted in reduced cell proliferation (Figure 4A). The addition of AMP-IBP5 under both normoglycemic and hyperglycemic conditions markedly increased cell proliferation. Meanwhile, as shown in Figure 4B, increased proliferation was observed in the AMP-IBP5-treated wounds of normal and diabetic mice compared to that in vehicle-treated wounds, as confirmed by Ki67-positive cells (Figure 4C). In addition, the inhibition of proliferation in the diabetic mouse wound area was rescued by AMP-IBP5 (Figure 4B, C). Moreover, similar to the effects on cell proliferation, HG inhibited spontaneous cell migration, and this inhibition was as high as 40%. As expected, AMP-IBP5 not only increased keratinocyte migration under normal conditions but also rescued the HG-induced inhibition of cell migration (Figure 4D). Similarly, as demonstrated by the wound scratch assay, AMP-IBP5- treated wounds closed rapidly when cells were cultured under normoglycemic and hyperglycemic conditions compared with their respective controls (Figure 4E, F). Mannitol was used as an osmotic control and did not affect cell proliferation or migration.

3.5 AMP-IBP5 restored HG-induced keratinocyte dysfunction by activating the EGFR and STAT pathways

 A previous study suggested that LL-37, which is involved in wound healing, induced 257 keratinocyte migration via EGFR activation.¹⁸ Furthermore, HG was reported to suppress 258 EGFR signal activation in epithelial cells.²⁷ Thus, we hypothesized that AMP-IBP5 could activate the EGFR pathway, thereby resulting in the inhibition of HG-induced keratinocyte dysfunction. In keratinocytes cultured under normoglycemic conditions, stimulation with AMP-IBP5 for 15 minutes increased the phosphorylation of EGFR by nearly 2-fold compared with that of control cells (Figure 5A, left panel). Under hyperglycemic conditions, spontaneous

 phosphorylation of EGFR decreased by nearly 30%; however, as expected, the addition of AMP-IBP5 markedly restored HG-induced EGFR inhibition.

 STAT1 and STAT3 are two intracellular transcription factors that are known to be involved 266 in the wound healing process.^{28,29} Similar to the effect on EGFR, stimulation with AMP-IBP5 for 15 minutes or 2 hours induced the phosphorylation of STAT1 (Figure 5A, middle panel) and STAT3 (Figure 5A, right panel), respectively, under both normoglycemic and hyperglycemic conditions. We also observed that AMP-IBP5-treated normal wounds showed stronger expression of p-EGFR, p-STAT1 and p-STAT3 than vehicle-treated wounds, while diabetic wounds displayed weaker staining than normal wounds. Similar to the *in vitro* results, the treatment of diabetic wounds with AMP-IBP5 rescued the expression of p-EGFR, p-STAT1 and p-STAT3 (Figure 5B). Furthermore, the AMP-IBP5-induced production of ANG and VEGF was significantly suppressed by AG1478 (EGFR inhibitor), fludarabine (STAT1 inhibitor) and cryptotanshinone (STAT3 inhibitor) (Figure 5C). Similarly, EGFR, STAT1 and STAT3 inhibitors markedly reduced AMP-IBP5-mediated proliferation (Figure 5D) and migration (Figure 5E, F) under HG conditions, suggesting crucial roles of the EGFR and STAT pathways in the AMP-IBP5- mediated restoration of keratinocyte functions in a hyperglycemic environment. **3.6 AMP-IBP5-mediated recovery of HG-induced keratinocyte dysfunction requires the**

activation of MAPK pathways

 We previously reported that AMP-IBP5 promoted the phosphorylation of MAPKs, which 283 are necessary for VEGF production by keratinocytes.¹⁹ Moreover, recent studies demonstrated 284 that HG-induced cell migration impairment was associated with JNK and p38 suppression.^{24,30} Therefore, we examined the effect of AMP-IBP5 on MAPK activation in keratinocytes cultured in HG conditions. Spontaneous phosphorylation of ERK, JNK and p38 was significantly reduced in HG-treated cells; however, this inhibitory effect was largely abrogated by the addition of AMP-IBP5 (Figure 6A). Next, we confirmed that the MAPK pathways were necessary for AMP-IBP5-mediated activation of keratinocytes under HG conditions. U0126 (ERK inhibitor), JNK inhibitor II and SB203580 (p38 inhibitor) noticeably suppressed AMP- IBP5-induced VEGF production, whereas only U0126 and JNK inhibitor II suppressed ANG production (Figure 6B). Moreover, these MAPK inhibitors attenuated AMP-IBP5-induced keratinocyte proliferation (Figure 6C) and migration (Figure 6D, E), indicating that MAPKs are involved in the protective effect of AMP-IBP5 on keratinocytes under hyperglycemic conditions.

4 Discussion

 Impaired diabetic wound healing is attributed to numerous risk factors, including 298 hyperglycemia, hypoxia, inflammation, angiogenesis dysfunction and neuropathy. $9-12$ Therefore, it is necessary to understand the complex pathogenesis of diabetic wound healing and develop holistic strategies to improve diabetic wound treatment. In this study, we

 Previous studies have demonstrated downregulated expression of AMPs in patients with diabetes, which explains why diabetic wounds show frequent bacterial infections that 322 contribute to tissue destruction and delayed wound healing.^{18,34} In this study, we also confirmed that the expression of IGFBP5, the parent protein of AMP-IBP5, was decreased under *in vivo* and *in vitro* diabetic conditions, suggesting that the absence of AMP-IBP5 could contribute to the impairment of diabetic wound healing.

 Several AMPs activate the EGFR pathway to induce cell migration, which is an indispensable step in the wound healing process. Tokumaru et al. indicated that LL-37 induced 328 keratinocyte migration via the transactivation of EGFR;¹⁸ Grazia et al. suggested EGFR involvement in frog skin-derived AMP esculentin-1a (1-21) NH2-induced migration of A549 330 cells.³⁵ In the current study, AMP-IBP5 not only activated EGFR but also rescued the attenuated activation of EGFR in keratinocytes cultured in HG conditions. However, further studies are needed to investigate how AMP-IBP5 activates EGFR.

 STAT family proteins are intracellular transcription factors that act downstream of tyrosine kinase receptors, including EGFR, and play essential roles in cell proliferation, differentiation 335 and angiogenesis.^{36,37} Among the STAT family members, STAT3 is well known for its role in 336 angiogenesis and wound healing.³⁷ In fact, STAT3 upregulates VEGF expression, thereby 337 inducing angiogenesis, and the blockade of STAT3 signaling inhibits VEGF expression.³⁸ 338 Furthermore, the levels of STAT1 are increased at wound sites after injury in a mouse model.³⁹

 such as the concentration of glucose and the duration of exposure to the HG environment. In addition, subsequent to EGFR activation, MAPK pathways have been shown to act 360 downstream and engage in the wound healing process.⁴⁸ Therefore, the ability of AMP-IBP5 to activate the EGFR, STAT and MAPK pathways may explain the significant improvements in diabetic wound healing.

 In summary, AMP-IBP5 promoted keratinocyte proliferation and migration in an HG environment and accelerated delayed angiogenesis and wound healing in diabetic mice. The molecular mechanism of AMP-IBP5 involves the EGFR, STAT and MAPK pathways. The potency of AMP-IBP5 might be due to its protective effect against glucotoxicity. We propose that AMP-IBP5 might be a new therapeutic candidate for the treatment of diabetic wounds. **Acknowledgments**: The authors wish to thank Michiyo Matsumoto for secretarial assistance and the members of the Atopy (Allergy) Research Center for discussion. **Funding**: This research was funded by a Grant-in-Aid for Scientific Research from the Ministry of Education, Culture, Sports, Science and Technology, Japan (Grant number: 26461703 and 21K08309 to F.N.) and by the Atopy (Allergy) Research Center, Juntendo

University, Tokyo, Japan.

 Author Contributions: Hainan Yue and François Niyonsaba designed the study, analyzed the data, conducted the experiments, and wrote the manuscript. Pu Song, Nutda Sutthammikorn and Yoshie Umehara contributed to the animal experiments. Juan Valentin Trujillo-Paez and

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 Figure 1 IGFBP5 expression was decreased in diabetic mouse skin and in high glucose-induced keratinocytes

(A) Biopsies were obtained from the dorsal skin of normal mice and streptozotocin-induced

diabetic mice. *Igfbp5* mRNA expression was evaluated by quantitative real-time quantitative

517 analysis using SYBR Premix Ex Taq. The results are shown as the mean \pm SD. ### $P < 0.001$,

- 518 $n = 4$. (B) Human epidermal keratinocytes were cultured in normal (6 mM) and high glucose
- (HG, 38 mM) conditions for 48 hours. *IGFBP5* mRNA expression was evaluated by
- quantitative real-time quantitative analysis using SYBR Premix Ex Taq. The values represent
- 521 the mean \pm SD. #### $P < 0.0001$, $n = 3$.

Figure 2 AMP-IBP5 promoted wound healing in normal and diabetic mice

 Dorsal full-thickness dermal wounds were created on normal mice and streptozotocin-induced 524 diabetic mice and were treated with 0.01% acetic acid (vehicle) or 100 μ M AMP-IBP5. (A) Representative images of the wound area from day 0 to day 16 postinjury in normal mice and 526 diabetic mice. Scale bar $= 1$ mm. (B) The average wound area after different treatments was calculated by ImageJ software. (C) Hematoxylin & eosin staining of wounds postinjury. 528 Arrows indicated wound edges. Scale bar = 0.2 mm. (D) Quantitative analysis of the percentage 529 of re-epithelialization postinjury. The values represent the mean \pm SD. * *P* < 0.05, ** *P* < 0.01, **** *P* < 0.0001 for comparisons between vehicle-treated wounds and AMP-IBP5-treated 531 wounds in normal mice. $\# P < 0.05$, $\#H P < 0.01$, $\#H H P < 0.001$ for comparisons between 532 vehicle-treated wounds in normal mice and those in diabetic mice. & *P* < 0.05, && *P* < 0.01 533 for comparisons between vehicle-treated wounds and AMP-IBP5-treated wounds in diabetic 534 mice, $n = 3$ wound areas/group.

535 **Figure 3 AMP-IBP5 rescued high glucose-induced impaired angiogenesis in diabetic** 536 **mice**

537 (A) The mRNA expression of *Ang*, *Egf*, *Vegf* in wounded skin from normal mice and diabetic 538 mice treated with 0.01% acetic acid (vehicle) or 100 μ M AMP-IBP5 at day 4 postinjury. (B) 539 Keratinocytes were cultured in 6 mM normal medium (NM) or 38 mM high glucose medium 540 (HGM) for 48 hours and stimulated with 10 μ M AMP-IBP5 for 48 hours. Mannitol (38 mM) 541 was used as an osmotic control. The levels of ANG, EGF and VEGF were measured by ELISA. 542 (C) Representative images of wounds from normal and diabetic mice treated with 0.01% acetic 543 acid (vehicle) and 100 μ M AMP-IBP5. On day 4 postinjury, sections were stained with 544 immunofluorescent anti-CD31 antibody, and CD31-positive staining is indicated by yellow 545 arrows; scale bar = 50 μ m. (D) Representative images of the macroscopic appearance of blood 546 vessels at the wound sites on day 4, day 8, day 12, and day 16 postinjury in normal mice and 547 diabetic mice treated with 0.01% acetic acid (vehicle) and 100 μ M AMP-IBP5. Scale bar = 1 548 mm. ** $P < 0.01$, *** $P < 0.001$, *** $P < 0.0001$ for comparisons between vehicle and AMP-549 IBP5 in normal mice or normal medium (NM). # *P* < 0.05, ## *P* < 0.01, ### *P* < 0.001 for 550 comparisons between vehicle treatment in normal mice or normal medium (NM) and diabetic 551 mice or high glucose medium (HGM). $\& P < 0.05$, $\& \& P < 0.01$, $\& \& P < 0.001$ for 552 comparisons between vehicle and AMP-IBP5 in diabetic conditions, $n = 3$.

553 **Figure 4 AMP-IBP5 attenuated high glucose-induced inhibition of keratinocyte** 554 **proliferation and migration**

555 Keratinocytes cultured in normal medium (NM), high glucose medium (HGM) or mannitol-556 containing medium for 48 hours were trypsinized and used for different assays. (A) Cells were 557 stimulated with 5 μ M AMP-IBP5 in normal medium (NM) or high glucose medium (HGM) 558 for 48 hours, and cell proliferation was assessed. (B) Immunohistochemical staining of Ki67 559 in the wounds 4 days after treatment. Scale bar = 200 μ m. (C) The number of Ki67-positive 560 cells was counted in 3 high-power fields of wound sections. (D) Keratinocytes in the upper 561 wells of the microchamber were allowed to migrate for 6 hours toward vehicle or 5 μ M AMP-562 IBP5 in normal medium (NM) or high glucose medium (HGM). Migrated cells were counted 563 in 3 high-power fields (HPFs) under a light microscope. (E) Keratinocytes were treated with 1 564 µg/ml mitomycin C for 2 hours. A scratch assay was performed. After treatment with mannitol 565 or 10 μ M AMP-IBP5 in normal medium (NM) or high glucose medium (HGM) for 48 hours, 566 keratinocytes were stained with crystal violet, and images were recorded, scale bar = $200 \mu m$. 567 (F) The average wound areas were calculated using ImageJ software. The results are shown as 568 the mean \pm SD. * *P* < 0.05, ** *P* < 0.01, *** *P* < 0.001 for comparisons between vehicle and

 AMP-IBP5 in normal mice or normal medium (NM). # *P* < 0.05, ## *P* < 0.01, ### *P* < 0.001 for comparisons between vehicle treatment in normal mice or normal medium (NM) and diabetic mice or high glucose medium (HGM). & *P* < 0.05, && *P* < 0.01, &&& *P* < 0.001 for 572 comparisons between vehicle and AMP-IBP5 in diabetic conditions, $n = 3-4$.

Figure 5 AMP-IBP5 restored high glucose-induced keratinocyte dysfunction by activating the EGFR and STAT pathways

 (A) Keratinocytes were cultured in normal, high glucose or mannitol-containing medium for 576 48 hours, stimulated with 10 μ M AMP-IBP5 for 15 minutes for EGFR and STAT1 and 2 hours for STAT3, and then subjected to Western blotting using antibodies against phosphorylated or unphosphorylated EGFR, STAT1 or STAT3. Bands were quantified using densitometry. The 579 results are expressed as the mean \pm SD. *** $P < 0.001$, **** $P < 0.0001$ for comparisons between nonstimulated and AMP-IBP5-stimulated cells in normal medium. # *P* < 0.05, ## *P* < 0.01 for comparisons between nonstimulated cells in normal medium and high glucose medium (HGM). & *P* < 0.05, && *P* < 0.01, &&&& *P* < 0.0001 for comparisons between nonstimulated cells and AMP-IBP5-stimulated cells in high glucose medium, *n* = 3. (B) Immunohistochemistry staining of wound sections 4 days postinjury. (C), (D) and (E) Keratinocytes cultured under high glucose (38 mM) conditions were pretreated with 0.1% DMSO (vehicle), 50 nM AG1478, 100 μM fludarabine and 2.5 μg/mL cryptotanshinone (CT) 587 for 2 hours and exposed to 10 μ M AMP-IBP5 for 48 hours. (C) The levels of ANG and VEGF in culture supernatants were measured by ELISA. (D) Keratinocyte proliferation was assessed using the BrdU labeling detection kit. (E) Keratinocyte migration was analyzed by scratch 590 wound assays *in vitro*; scale bar = 200 μ m. (F) The average wound areas were calculated using 591 ImageJ software. The results are expressed as the mean \pm SD. * $P < 0.05$, ** $P < 0.01$, **** P 592 \lt 0.0001 for comparisons between vehicle and AMP-IBP5. # $P \lt 0.05$, ## $P \lt 0.01$, ### $P \lt$ 0.001, #### *P* < 0.0001 for comparisons between AMP-IBP5-treated cells with or without 594 inhibitors, $n = 3$.

Figure 6 AMP-IBP5-mediated amelioration of high glucose-induced keratinocyte dysfunction requires the activation of MAPK pathways

 (A) Keratinocytes were cultured in normal, high glucose or mannitol-containing medium for 598 48 hours, stimulated with 10 μ M AMP-IBP5 for 15 minutes for JNK, 30 minutes for ERK and 60 minutes for p38, and then subjected to Western blotting using antibodies against phosphorylated or unphosphorylated ERK, JNK and p38. Bands were quantified using 601 densitometry. The results are shown as the mean \pm SD. * *P* < 0.05, *** *P* < 0.001, **** *P* < 0.0001 for comparisons between nonstimulated and AMP-IBP5-stimulated cells in normal 603 medium. $\#P < 0.05$ for comparisons between nonstimulated cells in normal medium and high glucose medium (HG). && *P* < 0.01, &&& *P* < 0.001 for comparisons between nonstimulated cells and AMP-IBP5-stimulated cells in high glucose medium, *n* = 3. (B), (C) and (D) Keratinocytes incubated with 38 mM glucose were pretreated with 0.1% DMSO (vehicle), 10 607 μM U0126, 10 μM SB203580 and 10 μM JNK inhibitor II (JNK inh II) for 2 hours and exposed 608 to 10 M AMP-IBP5 for 48 hours. (B) The levels of ANG and VEGF in culture supernatants 609 were measured by ELISA. (C) Keratinocyte proliferation was assessed using the BrdU labeling 610 detection kit. (D) Keratinocyte migration was analyzed by scratch wound assay *in vitro*, scale 611 bar = 200 μ m. (E) The average wound areas were calculated using ImageJ software. The results 612 are expressed as the mean \pm SD. ** *P* < 0.01, **** *P* < 0.0001 for comparisons between vehicle 613 and AMP-IBP5. $# P < 0.05$, $#H P < 0.01$, $#H \# P < 0.0001$ for comparisons between AMP-

614 IBP5-treated cells with or without inhibitors, $n = 3$.

Figure 2

Figure 3

Figure 4

Figure 5

Figure 6