Halting angiogenesis by non-viral somatic gene therapy alleviates psoriasis and murine psoriasiform skin lesions


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Dysregulated angiogenesis is a hallmark of chronic inflammatory diseases, including psoriasis, a common skin disease that affects approximately 2% of the population. Studying both human psoriasis in 2 complementary xenotransplantation models and psoriasis-like skin lesions in transgenic mice with epidermal expression of human TGF-β1, we have demonstrated that antiangiogenic non-viral somatic gene therapy reduces the cutaneous microvasculature and alleviates chronic inflammatory skin disorders. Transient muscular expression of the recombinant disintegrin domain (RDD) of metaggin (also known as ADAM-15) by in vivo electroporation reduced cutaneous angiogenesis and vascularization in all 3 models. As demonstrated using red fluorescent protein–coupled RDD, the treatment resulted in muscular expression of the gene product and its deposition within the cutaneous hyperangiogenic connective tissue. High-resolution ultrasound revealed reduced cutaneous blood flow in vivo after electroporation with RDD but not with control plasmids. In addition, angiogenesis- and inflammation-related molecular markers, keratinocyte proliferation, epidermal thickness, and clinical disease scores were downregulated in all models. Thus, non-viral antiangiogenic gene therapy can alleviate psoriasis and may do so in other angiogenesis-related inflammatory skin disorders.

Introduction

Dysregulated angiogenesis is emerging as a potential new target in inflammatory disorders, among which psoriasis, a common chronic inflammatory skin disorder that affects about 2% of the population, has attracted considerable interest among physicians and researchers alike (1–3). Due to its easy accessibility, psoriasis has increasingly become a model disorder in which to study the basic pathogenesis and therapy of chronic inflammatory diseases (4), genetic inheritance patterns (5, 6), complex cytokine networks (7), and central interactions of immune cells with epithelial tissues (8). In addition, several new therapies targeting immunological key mechanisms have been developed for psoriasis first (9–12).

Angiogenesis and morphological and functional alterations of microvessels are hallmark features of chronic inflammatory disorders, including psoriasis (13–15). It is thought that the proangiogenic microenvironment within psoriatic skin is induced by a T helper cell–initiated inflammatory response, which results in induction and activation of various proangiogenic factors, such as VEGF, HIF, TNF-α, and CXCL-8 (16–19). The VEGF-related angiogenic milieu is also modulated by regulatory T cells (20) and influences key features of psoriasis, including epidermal hyperplasia (14). A single nucleotide polymorphism within the VEGF-encoding gene is associated with severe and early-onset forms of psoriasis (21). Of note, several modern therapies of psoriasis affect not only immunological processes but also reduce pathological vascular functions (22–25). Thus, it is reasonable to assume that direct targeting of angiogenesis will be an effective means to halt the psoriatic disease process, although direct evidence is still scant (17, 26, 27).

Induced expression of adhesion molecules, such as integrins αβ3 and αβ3, on sprouting blood vessels is crucial for angiogenesis and serves as a key when using these receptors as therapeutic targets (28–30). Besides targeting components of the VEGF signaling pathway, adamalysin protein family members, also referred to as a disintegrin and metalloproteinases (ADAMs), are potential players interfering with angiogenesis. Metalloprotease-RGD-disintegrin (metaggin; also known as and referred to throughout as ADAM-15) is the only adalysin known to bind integrins αβ3 and αβ3 (31, 32). Indeed, a recombinant disintegrin fragment of ADAM-15 reduced angiogenesis and tumor growth (33). Therefore, in vivo expression of such factors is predicted to shift the proangiogenic milieu toward a more angiostatic situation, thus interfering with the pathogenesis of inflammatory disorders.

While it appears likely that in vivo expression of antiangiogenic factors would directly normalize the vascular dysregulation in chronic inflammation, gene therapeutic approaches have generally been hampered by the relatively low efficiency of naked DNA and safety concerns when viral vectors are used (34). In this situation, gene delivery by in vivo electroporation shows high transfection...
Figure 1
Expression and functional activity of RDD fusion proteins in vitro. (A) Human fibroblasts transfected with different constructs were analyzed for RDD expression by RT-PCR (lanes: 1, weight marker; 2, untransfected fibroblasts; 3, transfected with pVAX-mock; 4, transfected with pVAX-RDD; 5, transfected with pVAX-RFP-RDD; 6, negative control; 7, plasmid pVAX-RDD [positive control 1]; 8, plasmid pVAX-RFP-RDD [positive control 2]. (B) Murine LL/2 cells were transfected with the pVAX-RFP-RDD or the pVAX-RFP plasmid as indicated. Protein expression was detected by fluorescence microscopy. Original magnification, ×400. (C) HUVEC cultures were scratched in a standardized fashion (wounding of the monolayer), thus creating a cell-free lane (width, 0.5 mm) in each culture. Cultures were then incubated with supernatant from untransfected human fibroblasts (circles), from human fibroblasts transfected with the pVAX-mock plasmid (squares), or the pVAX-RDD plasmid (triangles). Migration of cells into the cell-free lane was monitored. Each data point represents the mean of 10 measurements (± SD). Units on the y-axis: μm. ***P < 0.001. (D) Representative HUVEC cultures described in C. The controls shown here are cultures incubated with supernatant from pVAX-mock–transfected fibroblasts, which were similar to the other controls. The margins of the cell layers are indicated by black lines. Original magnification, ×100. (E) Expression of the RDD target proteins, integrins αVβ3 and α5β1, on HUVEC as detected by flow cytometry. Strong signals are detected with antibodies directed against CD49e (α5 integrin subunit), CD51 (αV integrin subunit), and a combinatorial epitope of CD51 and CD61 (the αVβ3 integrin heterodimer).
efficiency, while avoiding the problem of using viral vectors (34, 35). In addition, given that electric pulse-mediated transfection of therapeutic genes into skeletal muscle is expected to result in long-term expression (36, 37), this strategy appears promising for pilot and proof-of-principle studies.

We have used xenotransplantation of human psoriasis in 2 distinct models (38) as well as a murine psoriasis-like skin disorder in K5.TGF-β1 transgenic mice (39) as independent angiogenesis-related skin disorders (40). In proof-of-principle experiments, we provide the first experimental evidence to our knowledge that non-viral gene therapy by transient expression of the disintegrin domain of ADAM-15 inhibits angiogenesis both in psoriasis and psoriasisiform skin inflammation in K5.TGF-β1 mice, leading to significant improvement of both diseases.

**Results**

A recombinant disintegrin domain inhibits endothelial cell functions in vitro. ADAM-15 is evolutionarily highly conserved; the disintegrin domains of murine and human ADAM-15 share amino acid sequence homologies of 77.6% (41). We have generated a recombinant disintegrin domain (RDD), homologous to murine, human, and bovine ADAM-15 (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI41295DS1) that interferes with αvβ3 and αvβ1 integrins of either species (33). In our first series of experiments, the inhibitory capacity of the RDD molecule was demonstrated on cultured endothelial cells. When monolayers were “wounded” by standardized scratching and closure of the cell-free areas was monitored, RDD significantly inhibited the migration of endothelial cells in a dose-dependent fashion. In 4 independent experiments (16–21 individual cultures for each condition), endothelial cell migration in the presence of 15 μg/ml RDD protein was reduced to 54.5% (± 3.44%); P < 0.0001) as compared with that of untreated control cultures (Supplemental Figure 2). As detected using Western blot analysis, exposure of cultured endothelial cells to RDD reduced expression of several molecules implicated in signaling or migration, including integrin-linked kinase (ILK), focal adhesion kinase (FAK; also known as protein tyrosine kinase-2 [PTK2]), and activated leukocyte cell adhesion molecule (ALCAM; also known as CD166). Decreased activity of MMP-2 was demonstrated by zymography. In contrast, expression of tissue inhibitor of metalloproteinase-2 was increased (data not shown).

To demonstrate that RDD secreted by transfected cells impaired endothelial cell functions, normal human fibroblasts (Figure 1A) were transfected by electroporation with different constructs. The RDD-encoding sequence and several control sequences were cloned into the pVAX1 expression vector (Supplemental Figure 1). The respective plasmids, pVAX-mock, pVAX–red fluorescent protein (pVAX-RFP), pVAX-RDD, and pVAX-RDD-RFP, were obtained from E. coli bacteria after transformation. Each plasmid construct was sequenced, and no sequence alterations were observed (data not shown). Expression of the RFP-coupled fusion protein of RDD (pVAX-RDD-RFP) as well as the control RFP (pVAX-RFP) could be readily detected within transfected murine cells using fluorescence microscopy (Figure 1B). Morphological changes or altered growth characteristics of the transfectants were not detected (data not shown). HUVEC monolayers, which strongly expressed the RDD target proteins, integrins αvβ1 and αvβ3, were incubated with supernatant from different transfected fibroblast populations, and closure of the cell-free lane (artificial wound generated by standardized scratching) was monitored microscopically for up to 29 hours. Cultures incubated with supernatant from RDD-expressing transfectants but not controls showed markedly delayed migration of endothelial cells into the gap (P < 0.0001 at all measured time points, i.e., 6, 12, and 24 hours), indicating functional activity of RDD secreted by transfectants (Figure 1, C and D). RDD target proteins were highly expressed in HUVECs (Figure 1E).

Expression of RDD in vivo. The next series of experiments was designed to achieve in vivo expression of the antiangiogenic peptide by non-viral gene therapy. Toward this end, the plasmids were transfected into the calf muscles (musculus [m.] tibialis cranialis) of mice by DNA electrotransfer, using an adjusted protocol with an initial high-voltage pulse (700 V/cm, 100 μs) and 8 subsequent low-voltage pulses (100 V/cm, 50 μs), a procedure that had been optimized in an extensive series of preliminary experiments (data not shown) (42). This combination of pulses facilitates gentle cell permeabilization, followed by electrophoresis of the plasmids into the cells, while largely avoiding damage to the overlying skin and other adjacent tissues (34, 43). In order to determine the persistence of the transgene in various tissues of the body, both male and female mice were analyzed for up to 4 months after somatic gene therapy with RDD. Vehicle-treated mice were analyzed as controls. It was found that the transgene was below the level of quantification, as determined by quantitative PCR in all tissues, including the gonads. Only in the treated muscles was the transgene detectable by the end of the observation period, albeit expression levels were diminished by approximately 3 logs compared with those at the early time points (data not shown).

Fluorescence microscopy demonstrated clear red signals within the calf muscles of mice transfected with the pVAX-RDD-RFP fusion protein or pVAX-RFP (Figure 2A). In addition, immunohistochemistry was performed to confirm the topographic mapping of the transgene within the skeletal muscle fibers (Figure 2B). Time course experiments revealed that expression of the transgenes was detectable as early as 24 hours after electrotansfer, reached a peak after 3 days and tapered off within 2–3 weeks (data not shown). Inflammatory reactions at the transfection sites or other unwanted side effects were not observed (data not shown). Of note, when organs distant from the transfection site were analyzed in mice transfected with the pVAX-RDD-RFP construct, clear fluorescence signals were detected within the cutaneous connective tissue but not the epidermis (Figure 2A). In contrast, control animals transfected with RFP alone or mock-transfected animals did not show dermal fluorescence (Figure 2A), suggesting that RDD-containing peptides became deposited within the skin. The results obtained by fluorescence microscopy were confirmed using immunohistochemistry, again showing deposition of the RDD-containing products but not the controls in the vicinity of some blood vessels (Figure 2C). Deposition of the RDD-RFP peptide in other organs, including muscle, liver, and kidneys (only single cells showed weak reactivity; data not shown), was weak, if detectable at all, and morphological alterations in other organs were not detected (data not shown). As demonstrated by Western blot analysis using RFP-specific antibodies, secreted protein could be detected within the serum of mice (n = 4 mice) transfected with the pVAX-RDD-RFP construct (Figure 2D). Since no signal was detected when mice were treated with pVAX-RFP (n = 4 mice), these results suggested that RDD-containing sequences were readily secreted (Figure 2D).

To exclude toxicity, a repeated toxicity study (4 intramuscular administrations in 1-week intervals) was performed in both male
and female mice. Animals were sacrificed just after the last administration or 1 month after the last administration. No pathological alterations were detected clinically and histopathologically in various tissues, including the gonads (data not shown). With the exception of a reversible inflammatory reaction within the injected muscle, wild-type immunocompetent mice did not show any signs of adverse effects.

In vivo RDD gene therapy stops psoriasiform changes in K5.TGF-β1 transgenic mice. Having established that non-viral gene transfer could be successfully applied in vivo, the impact of RDD expression on cutaneous vascularization and the course of the psoriasis-like disorder in K5.TGF-β1 transgenic mice (39) was assessed. In a series of preparatory experiments, it was established that K5.TGF-β1 transgenic mice showed significantly increased size and density of cutaneous blood vessels (Figure 3A). As an independent second method, blood flow within the upper plexus of the skin was assessed at several time points for up to 12 days by high-resolution ultrasound. In order to achieve a reliable and reproducible functional quantification of the cutaneous blood flow (which shows considerable site-dependent variability), 4 areas distant from the transfection sites (2 on the abdomen and 1 on each thigh) were analyzed in a standardized manner (Supplemental Figure 3), and the mean blood flow from all 4 areas was calculated for each mouse. Again, this standardized functional approach demonstrated significantly increased vascularization and blood flow in K5.TGF-β1 transgenic mice compared with that in wild-type mice (14.3% of the total dermal volume $\pm$ 4.1% in wild-type and 26.9% $\pm$ 5.4% in K5.TGF-β1 transgenic mice; $P < 0.001$; Figure 3B).

To determine the optimal concentration of RDD plasmid for transfections, we injected anesthetized K5.TGF-β1 transgenic mice intramuscularly with 3 different doses of the pVAX-RDD plasmid (12.5, 25, or 50 μg per hind leg, the latter dose being the highest that was tolerable without injection-related side effects) and performed electroporation. Given that 50 μg pVAX-RDD plasmid yielded the strongest diminution of the cutaneous blood flow (Figure 3C), all experiments were performed using 50 μg plasmid in each calf muscle.

When K5.TGF-β1 transgenic mice ($n = 8$ mice per group) were subjected to electroporation only (PBS control), pVAX electroporation (mock transfection control), or pVAX-RDD electroporation...
and monitored for 12 days, some remarkable effects of RDD on the psoriasiform phenotype became apparent. While the disease severity score showed the expected worsening in both control groups (scores 1.15 [± 0.25] and 1.20 [± 0.17], respectively, by the end of the observation period), disease progression was completely stopped in the group treated by RDD gene therapy (score 0.56 [± 0.25]; \( P < 0.01 \) compared with the pVAX-mock control group; Figure 3D). End-point analysis by immunohistochemistry and subsequent morphometric analysis confirmed the in vivo observations: RDD gene therapy resulted in significantly reduced cutaneous vascularization (overall reduction of CD31-positive blood vessels of 25% compared with that of either of the controls, as determined by morphometric analyses; \( P < 0.01 \)). Thus, morphometric analyses (performed as a second independent method) confirmed the results obtained by Doppler ultrasound. Likewise, gene therapy with RDD led to significantly reduced epidermal thickness (reduction of 32%; \( P < 0.01 \); Figure 3, E and F). The number of cutaneous T lymphocytes was not significantly affected by RDD gene therapy (epidermal CD8+ T cells, 14 [± 7] per mm in control mice and 10 [± 8] per mm in mice treated with pVAX-RDD; CD4+ epidermal cells, 21 [± 12] per mm in control mice and 16 [± 10] per mm in mice treated with pVAX-RDD).
RDD gene therapy prevents the development of psoriasis in the hu/SCID xenotransplantation model. (A) Uninvolved skin from psoriasis patients was transplanted onto immunodeficient Prkdc<sup>−/−</sup> mice, and recipient mice were injected intraperitoneally with PBS (left panel) or $10^7$ activated PBMCs from the same donors to induce psoriasis (middle and right panels). Two days prior to injection of PBMCs, the mice were subjected to gene therapy with pVAX-mock or pVAX-RDD. The macroscopic aspect of the blood vessels on the underside of the skin transplants demonstrates that gene therapy with RDD markedly reduced cutaneous vascularization. Original magnification, ×3. (B) Prkdc<sup>−/−</sup> mice were treated as described in A, and the transplanted human skin was analyzed by immunohistochemistry using the indicated reagents detecting blood vessels. Somatic gene therapy with pVAX-RDD, but not that with pVAX-mock or pVAX-RFP, reduces number and size of cutaneous blood vessels of both human (CD31 and αSMA) and murine (BS-1) origin. Scale bar: 50 μm. (C) Human skin transplants from Prkdc<sup>−/−</sup> mice were further analyzed by H&E staining (left panels) and by immunohistochemistry using the indicated antibodies detecting proliferating cells (Ki67) or T cells (CD3). Both the epidermal thickness and the number of proliferating epidermal keratinocytes are reduced in transplants from mice treated by RDD gene therapy. Scale bar: 50 μm. (D) Quantitative morphometric analysis of transplants from Prkdc<sup>−/−</sup> mice. The top graph depicts epidermal thickness as determined by blinded measurements. The bottom graph shows the number of proliferating epidermal cells in mice subjected to RDD gene therapy compared with that of the controls. ***P < 0.001. Values represent mean ± SD.
Antiangiogenic RDD gene therapy prevents psoriasis in an inducible xenotransplantation model. Having shown that RDD gene therapy was efficacious in K5.TGF-β1 transgenic mice, we next assessed whether this antiangiogenic gene therapy would affect the pathogenesis of human psoriasis in the psoriasis/SCID mouse xenograft model, in which human skin was transplanted onto Prkdc<sup>scid</sup> mice (38) and which allows the study of psoriasis development in connection with angiogenesis (44). After several preliminary experiments in which it was established that in vivo RDD expression can be used in the xenotransplantation models, noninvolvved skin from psoriasis patients (4 patients, each biopsied piece of skin yielding >6 transplants) was transplanted, and psoriasis was induced by injection of staphylococcal enterotoxin B (SEB) and IL-2-stimulated autologous PBMCs (38, 44). The DNA electrotransfer using 50 μg pVAX-RDD or pVAX-RFP (into both tibialis cranialis muscles) was performed 2 days prior to the induction of psoriasis. Disease development was monitored for 14 days.

Psoriasis induction in the transplants was accompanied by profoundly increased vascularization, both macroscopically (Figure 4A) and histopathologically (Figure 4B). Given that the grafts may harbor blood vessels of both human and murine origin (45), we evaluated the effect of RDD using several staining methods. Human-derived vessels were detected using antibodies directed against human CD31 (also known as PECAM-1) and α-SMA. Murine endothelial cells were stained using anti-mouse BS-1-lectin. Of note, gene therapy with RDD resulted in marked reduction of cutaneous blood vessels compared with those of control-treated mice (Figure 4, A and B). Morphometric analysis revealed reduction of 42% (P < 0.001, comparing vascularization of control [pVAX] and RDD-treated mice).

When other features of psoriasis were assessed morphometrically, highly significant effects were observed regarding epidermal thickness and numbers of proliferating cells detected by reactivity with the Ki67 mAb. The epidermal thickness (measured in 8 to 10 randomly chosen sites in the middle of each graft in a blinded fashion) was 162.5 μm (SD = 9.7) in the control pVAX-RFP–treated mice (n = 9) and 89.7 μm (SD = 9.8) in the pVAX-RDD–treated mice (n = 15; reduction of 45%; P < 0.001; Figure 4, C and D). In addition, the numbers of proliferating epidermal cells showed highly significant differences between controls and RDD-treated mice (121.5 cells/mm [SD = 15.9] in pVAX-RFP–treated mice and 42.9 cells/mm [SD = 6.0] in pVAX-RDD–treated control mice; reduction of 65%; P < 0.01; Figure 4, C and D), indicating that the primary effect of RDD gene therapy on cutaneous vascularization indeed alleviated typical epidermal features of psoriasis. Numbers of cutaneous T cells were not significantly different in RDD-treated versus control-treated mice (12 ± 4) CD3<sup>+</sup> epidermal cells/mm in control mice vs. 9 ± 5) CD3<sup>+</sup> epidermal cells/mm in pVAX-RDD–treated animals; Figure 4C). There were slightly reduced numbers of Mac-1–expressing cells (predominantly macrophages and dendritic cells; 35 ± 21) cells/mm in control mice vs. 24 ± 14 cells/mm in mice treated with pVAX-RDD gene therapy). Likewise, when skin sections were stained for CD1a, detecting epidermal Langerhans cells in human skin transplants, 39 (± 22) cells/mm epidermis were seen in control animals, whereas the respective number was 23 ± 16) cells/mm in transplants from mice treated with pVAX-RDD gene therapy (see below). A similar tendency was seen when expression of CD68, a marker for dermal macrophages, was assessed (data not shown). However, those differences did not reach statistical significance.

Therapeutic efficacy of RDD gene therapy on fully developed psoriatic phenotype. We next assessed the impact of halting angiogenesis by RDD gene therapy on full-fledged human psoriatic skin (4 patients, each biopsy specimen yielding >8 transplants). The treatment was administered 14 days after transplantation as intramuscular injection of 50 μg pVAX-RDD or the control pVAX-RFP in both m. tibialis cranialis and subsequent electroporation. The conspicuously prominent vascularization visible on the undersides of the transplanted skin flaps in the control mice was markedly reduced in mice treated by RDD gene therapy (Figure 5A). This macroscopic feature was paralleled by markedly reduced vascularization on the microscopic level (Figure 5B), indicating that in vivo expression of the RDD fragment caused diminution of number and size of the hypertrophic cutaneous microvessels. Morphometric analysis revealed reduction of 46% (P < 0.001 comparing vascularization of control [pVAX] and RDD-treated mice).

Of note, RDD transfection also led to significantly reduced epidermal thickness and numbers of proliferating cells compared with those of the controls. The epidermal thickness was 166.85 μm (SD = 12.1) in pVAX-RFP–treated mice (n = 14 animals) and 88.29 μm (SD = 7.7) in pVAX-RDD–treated mice (n = 13 animals; P < 0.001; Figure 5, C and D). Likewise, the numbers of proliferating epidermal cells were significantly reduced in RDD-treated mice (312.0 cells/mm [SD = 48.3] in pVAX-RFP–treated mice and 42.5 cells/mm [SD = 14.2] in pVAX-RDD–treated mice; P < 0.001; Figure 5, C and D). Finally, somatic gene therapy using RDD led to reduced expression of HIF1α and FLT1 (expressed by endothelial cells) in the dermal compartment of the skin, whereas expression of angiopoietin-1 and VEGFα remained unchanged compared with that of control animals (Figure 6).

Thus, antiangiogenic gene therapy was highly efficacious both in the prevention and treatment of psoriasis lesions in vivo.

Discussion

Following early reports suggesting that angiogenesis plays a role in psoriasis (13, 46–49), specific modulation of dysregulated vascularization is currently evolving into a hot research topic (17, 26, 50–54). However, direct evidence to corroborate this hypothesis is still scant. A VEGF-directed antibody led to complete remission of psoriasis in 1 patient (55). In addition, several transgenic mouse models have been generated to clarify whether angiogenesis-related factors can initiate psoriasiform pathologies (40). For example, targeting VEGF overexpression to the epidermis (56, 57) or transgenic overexpression of Tie-2 resulted in psoriasis-like features (58). In addition, several mouse models generated by manipulations unrelated to angiogenesis, such as adoptive transfer of minor histocompatibility-mismatched CD4<sup>+</sup>/CD45RB<sup>hi</sup> T cells into immunodeficient recipients (59) or K5.TGF-β1 transgenic mice (39), feature increased size and density of cutaneous blood vessels. Based on this briefly delineated circumstantial evidence, it is reasonable to assume that direct targeting of cutaneous angiogenesis will be efficacious to treat psoriasis. In this context, we provide experimental in vivo proof of concept that halting angiogenesis can indeed alleviate psoriasis.

Using K5.TGF-β1 transgenic mice and 2 independent human/mouse transplantation models, we demonstrate the principle that psoriasis and related skin disorders can be alleviated by an innovative non-viral RDD-based antiangiogenic gene therapy that results in transient inhibition of angiogenesis. It is conceivable that this approach will be used in the future to express other therapeutic
Figure 5
RDD gene therapy reduces cutaneous vascularization and alleviates psoriasis in human skin transplanted onto immunodeficient mice. (A) Psoriatic skin with a full-fledged phenotype was transplanted onto Prkdc<sup>−/−</sup> mice, and recipient mice were either left untreated (representative of n = 12 mice) or subjected to gene therapy with pVAX-RFP (representative of n = 14 mice) or pVAX-RDD (representative of n = 13 mice). The macroscopic aspect of the blood vessels on the underside of the skin transplants demonstrates that gene therapy with RDD reduces cutaneous vascularization. Original magnification, ×3. (B) Transplanted psoriatic skin was analyzed by immunohistochemistry using the indicated reagents detecting blood vessels. Somatic gene therapy with pVAX-RDD results in reduction of both number and size of cutaneous blood vessels as compared with that of the controls. Blood vessels of human (CD31 and SMA) as well as murine (BS-1) origin are affected. Scale bar: 50 μm. (C) Transplants of fully developed psoriasis transplanted onto Prkdc<sup>−/−</sup> mice were further analyzed by staining with H&E (left panels) or by immunohistochemistry using the indicated antibodies detecting proliferating cells (Ki67) or T cells (CD3). Both the epidermal thickness and the number of proliferating epidermal keratinocytes are reduced in transplants from mice treated by RDD gene therapy. Scale bar: 50 μm. (D) Quantitative morphometric analysis of transplants from Prkdc<sup>−/−</sup> mice. The top graph depicts the epidermal thickness as determined by blinded measurements. The bottom graph shows the number of proliferating epidermal cells in mice subjected to RDD gene therapy compared with that of the controls. Values represent mean ± SD. ***P < 0.001.
molecules interfering with angiogenesis, such as components of the VEGF signaling pathway. Alternative or improved methods of gene delivery, such as microneedles in the skin, might also be promising future directions (34, 60). While it is possible in principle that RDD expression has non-angiogenesis–related effects (because the target proteins may be expressed by several cell types), no such effects have been observed in our experiments. An exception may be the observed tendency (which was not statistically significant) toward reduced macrophages and dendritic cells in RDD-treated mice, an effect that might have contributed to the overall therapeutic efficacy of RDD gene therapy. When the RDD-RFP fusion protein was expressed in vivo, deposition was only observed within the dermal connective tissue, i.e., the compartment in which angiogenesis occurs. In addition, it is known that ADAM-15 regulates angiogenesis through its disintegrin-like domain that binds to the $\alpha_v\beta_3$ integrin (41). Given that $\alpha_v\beta_3$ is upregulated in lesional dermal vessels in psoriasis as well as in the active psoriasis xenograft model, this receptor presumably contributes to dermal angiogenesis (44, 61). It is thought that RDD interferes with signal transduction by its target receptors, a notion that is supported by decreased expression or activity of ILK, FAK, MMP-2, and ALCAM on cultured endothelial cells upon exposure to RDD. On the cellular level, binding of the RGD motif of ADAM-15 to the $\alpha_v\beta_3$ and $\alpha_5\beta_1$ integrins led to enhanced cell-cell interactions in a recent study, indicating that ADAM-15 plays a role in cell signaling events (32, 62–64). As suggested by our experiments, RDD-based gene therapy interfered transiently with important endothelial cell functions in vivo, thus resulting in reduced cutaneous vascularization in conditions with hypertovascularization. Non-viral gene delivery methods have attracted considerable interest in recent years (34). However, a drawback is the low efficiency of naked DNA gene transfer compared with that of viral delivery methods. In vivo electroporation to facilitate gene transfer

**Figure 6**
RDD gene therapy and mediators of endothelial functions in psoriatic skin transplanted onto immunodeficient mice. (A) Uninvolved skin from psoriasis patients was transplanted onto immunodeficient Prkdc<sup>scid</sup> mice, and the recipient mice were treated as outlined in Methods to induce psoriasis (PN, top row), or psoriatic skin with a full-fledged phenotype (PP, bottom row) was transplanted onto Prkdc<sup>scid</sup> mice. The recipient mice were either subjected to control treatment using pVAX-mock or pVAX-RFP or pVAX-RDD. Expression of HIF1α was detected by immunohistochemistry. Scale bar: 50 μm. (B) Expression of angioptoin-1 (ANG1) was detected by immunohistochemistry in transplants from mice treated as described in A. Scale bar: 50 μm. (C) Expression of VEGF-1 was detected by immunohistochemistry in transplants from mice treated as described in A. Scale bar: 50 μm. (D) Expression of FLT1 was detected by immunohistochemistry in transplants from mice treated as described in A. Scale bar: 50 μm. (E) Expression of CD1a was detected by immunohistochemistry in psoriatic transplants from mice treated as described in A. Scale bar: 50 μm.
shows high rates of transfection in several tissue types and avoids the problems associated with viral vectors (34, 35). Indeed, electrotransfer of a plasmid coding for IL-12 to metastatic melanomas resulted in increased IL-12 levels, tumor necrosis, and a lymphocytic infiltrate. In 8 out of 19 patients, disease stabilization or partial remission occurred, and in 2 patients, even non-treated distant lesions showed complete regression (65). In a preclinical study using collagen-induced arthritis in mice, intraarticular DNA electrotransfer of plasmids encoding for soluble TNF receptor variants led to decreased ankle joint destruction (66), underscoring the potential of DNA electrotransfer in inflammatory diseases.

Vector-directed immune responses are a major challenge with all types of gene therapy. Plasmid DNA vectors rapidly stimulate the innate immunity via TLR activation on leukocytes, and specific cellular and humoral responses may occur within a few days (67). Thus far, we have not observed abnormal immune responses in our models. However, since we have used mice with altered immune-related phenotypes, potential immune reactions need to be investigated in further studies.

Plasmid DNA electrotransfer is an attractive approach for systemic transgene delivery to patients with psoriasis. However, in this proof-of-concept study, we used a constitutive promoter. For future clinical applications, it may be advantageous to regulate gene expression or to implement tissue/cell-specific expression. Exogenous compounds, such as the doxycycline-dependent Tet-on system (68), the GeneSwitch system, which uses progerosterone antagonists (69), or tissue-specific promoters (70), can control transgene expression through inducible ligand-dependent promoter complexes. Recently, it was shown that it is possible to restrict the expression of a DNA plasmid to keratinocytes using a keratin 14 promoter and to mature dendritic cells using a facin promoter (71). DNA electrotransfer to the readily accessible skin is an attractive way of administering transgene constructs. Keratinocytes are interesting targets for transfection from a safety point of view, because they show high cell turnover and no need for exogenous controls, as the transgene will only be expressed for the short period of the turnover time. Thus, transfection of keratinocytes holds promise for future treatments of skin diseases. However, the skin barrier function involves challenges to penetration, electrical field distribution, and electrode design (60).

In any case, our study provides the first experimental evidence to our knowledge that targeting angiogenesis by non-viral somatic gene therapy is a promising approach for future treatments of psoriasis and, possibly, other inflammatory disorders.

**Methods**

**Plasmid preparation.** The plasmids pVAX1 (Invitrogen), pVAX-RDD, pVAX-RFP, and pVAX-RDD-RFP (Supplemental Figure 1A and ref. 33) were used. pVAX-RFP contained a rPFP-Monomer-N1 derived from the tetrameric *Discosoma* RFP (Clontech). RFP and/or RDD were cloned into the pGA4 (ampR) site of pVAX1. The plasmids were amplified in One Shot TOP10F- E. coli Competent Cells (Invitrogen) and prepared using the EndoFree Plasmid Mega Kit or Giga Kit (Qiagen) or the NucleoBond XtraMaxi Kit (Macherey-Nagel). Plasmid concentration and quality were controlled by spectrophotometry and gel electrophoresis and confirmed by sequencing.

**Cells and functional in vitro assays.** HUVECs (Cambrex) or C6PA endothelial cells (33) were cultured in Endothelial Cell Growth Medium (PromoCell). Human fibroblasts were cultured in MEM, and murine LL/2 cells (LGCT Standards) were cultured in RPMI1640. The Nucleofector NHDF Kit (Amaxa) was used for fibroblast transfections, and HiPerFect (Qiagen) was used for LL/2 cells. Transfection efficiencies were determined by FACS.

**HUVEC migratory activity was assessed for up to 29 hours in a standardized scratch-wound assay in 6-well plates, using conditioned medium from different fibroblast transfectants.**

**Mouse models.** Experiments using human skin were approved by the Danish National Committee on Biomedical Research Ethics and informed consent was obtained from the patients. Animal experiments were approved by the governmental authorities (Regierung von Unterfranken or Danish Experimental Animal Inspectorate). Mice were housed under SPF conditions.

**Mice with epidermal expression of latent human TGF-β1 (K5.TGF-β1) (39) were genotyped by PCR. All experiments were performed with offspring from wild-type females and transgenic males, and wild-type littermates served as controls. Transgenic and wild-type mice were used at 4–8 weeks of age. All mice were monitored in a blinded and standardized manner.**

For xenotransplantation, psoriasis patients without systemic treatment or phototherapy for 2 months and without topical treatment for at least 14 days were biopsied using a keratome (0.2 mm; 4 biopsies of involved psoriatic skin from 4 patients [for the active psoriasis model] and 4 biopsies of noninvolved skin from 4 patients [for the inductive model]). Mice were anesthetized by intraperitoneal injection with 100 mg/kg ketamine (Parke-Davis) and 20 mg/kg xylazine (Bayer) in 0.5 ml PBS. Skin pieces were transplanted onto the backs of Prkdc<sup>−/−</sup> mice (Taconic), and the wounds were closed using Histoacryl (TissueSeal). To induce psoriasis in nonlesional xenografts, 10<sup>5</sup> autologous PBMCs, activated by 20 U/ml IL-2 and 1 μg/ml SEB (Sigma-Aldrich), were injected intraperitoneally after 14 days.

**Intramuscular DNA electrotransfer.** Gene therapy was performed in 6- to 12-week-old K5.TGF-β1 transgenic mice and in Prkdc<sup>−/−</sup> mice on day 12 after transplantation of nonlesional skin (2 days before PBMC injection) or on day 14 after transplantation of involved psoriatic skin. The mice were anesthetized, and 50 μg plasmid DNA in 30 μl PBS was injected into each tibialis cranialis muscle. Plate electrodes (4 mm) and a specially modified Cliniporator (IGEA) were used to apply 1 high-voltage pulse (700 V/cm, 100 μs) and 8 low-voltage pulses (100 V/cm, 50 μs).

Persistence of the transgene was determined in various tissues (including untreated and treated muscles as well as the gonads) using quantitative RT-PCR (level of quantification, 100 plasmid copies) at various times for more than 4 months after gene electrotransfer.

**Western blot analysis.** To detect secreted protein within the serum of treated mice, 150 μg total serum was loaded in each lane in a 12% polyacrylamide gel with 1X MOPS. The gel was blotted using the dry iBlot system (Invitrogen). Membranes were blocked in 5% skimmed milk, washed in TBS, and incubated with rabbit polyclonal antibody against RFP (Ab34771; Abcam). The signals were visualized using an enhanced chemiluminescence Western blotting detection system according to the manufacturer’s instructions (Amersham, GE Healthcare).

**Histology and immunohistochemistry.** Paraffin-embedded tissues were stained using H&E according to standard protocols. For immunohistochemistry of murine tissues, cryostat-cut sections (4 μm) were stained as described previously (59). For immunohistochemistry of human tissue, 4-μm paraffin-embedded sections were stained. Antibodies against the following antigens were used: mCD31 (clone MEC13.3; BD Biosciences), mCD51 (RMV-7; BD Biosciences), mCD49e (HMα5; BD Biosciences), RFP (Ab34771; Abcam), hANGPT1 (HPA018816; Sigma-Aldrich), hCD1a (M3571; Dako) hCD3 (UCHT1; Dako), hCD31 (JC70A; Dako), hCD68 (M0876; Dako), hFLT1 (HPA011740; Sigma-Aldrich), hHIF1A (HPA001275; Sigma-Aldrich), hKi67 (MIB-1; Dako), hCD31 (Jc70A; Dako), hCD68 (M0876; Dako), and hVEGFA (M7273; Dako). Furthermore, Bandeiraea simplicifolia-1-agglutinin (BS-1-lectin; Sigma-Aldrich), staining vascular endothelium in murine tissues, cryostat-cut sections (4 μm). Primary antibodies were replaced with appropriate control antibodies. As a positive control, a colon adenocarcinoma was used for the Ki67 stainings, normal...
human and murine skin samples were used for the BS-1-stainings, and a multilobed of liver, kidney, and tonsil were used for the other stainings. Samples were analyzed in a blinded fashion using light and fluorescence stereomicroscopy (Leica or Zeiss). Morphometric analyses to quantitate cutaneous vascularization were performed using ImageJ freeware (http://rsbweb.nih.gov/ij) as described previously (73, 74). The epidermal thickness was measured at 10 randomly chosen sites in each sample (LAS EZI1.6.0, Leica), and averages of the measurements were calculated. Keratinocyte proliferation was determined by counting Ki67+ cells per mm epidermis.

Flow cytometry. Antigen expression was determined using antibodies against the following antigens: CD49d (SAM1); Beckmann-Couller), CDS1 (13C2; Chemikon), and CDS1/CDD1 (LM609; Chemikon). Cells were analyzed using a FACSCan FII and the DIVA software (BD Biosciences).

RT-PCR. Expression of RDD-encoding mRNA was analyzed using semiquantitative RT-PCR. Total RNA was extracted using the E.Z.N.A. RNA Kit (Omega). Thereafter, cDNA was generated using the First Strand cDNA Synthesis Kit (Fermentas) and an oligo(dT) primer (Fermentas). For amplification, Sawady-Taq (PeqLab) and sequence-specific primers (forward: 5′-GCGCCCTTGGTGGTGAACAC-3′; reverse: 5′-AGTCATCCAGGAAGCCA-3′) were used. PCR was performed in a Mastercycler (Eppendorf) as follows: 5 minutes at 94°C, followed by 30 cycles of 30 seconds at 94°C, 30 seconds at 60°C, and 5 minutes at 72°C.

In vivo blood flow. The Vevo770 Imaging System, equipped with a RMV708 Scanhead and a 3D motor (VisualSonics), was used for high-resolution Doppler ultrasound in 4 areas in each mouse in a standardized fashion (Supplemental Figure 4).


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