Psoriasiform skin disease in transgenic pigs with high-copy ectopic expression of human integrins α2 and β1

ABSTRACT

Psoriasis is a complex human-specific disease characterized by perturbed keratinocyte proliferation and a pro-inflammatory environment in the skin. Porcine skin architecture and immunity are very similar to that in humans, rendering the pig a suitable animal model for studying the biology and treatment of psoriasis. Expression of integrins, which is normally confined to the basal layer of the epidermis, is maintained in suprabasal keratinocytes in psoriatic skin, modulating proliferation and differentiation as well as leukocyte infiltration. Here, we generated minipigs co-expressing integrins α2 and β1 in suprabasal epidermal layers. Integrin-transgenic minipigs born into the project displayed skin phenotypes that correlated with the number of inserted transgenes. Molecular analyses were in good concordance with histological observations of psoriatic hallmarks, including hypogranulosis and T-lymphocyte infiltration. These findings mark the first creation of minipigs with a psoriasiform phenotype resembling human psoriasis and demonstrate that integrin signaling plays a key role in psoriasis pathology.

KEY WORDS: DNA transposition, Integrin, Pig cloning, Psoriasis, Skin inflammation, Sleeping Beauty

INTRODUCTION

Psoriasis vulgaris is an inflammatory and chronic cutaneous disorder affecting an estimated 2–3% of the Caucasian population worldwide. Main clinical hallmarks include well-demarcated thick, red and scaly skin plaques, whereas key histological hallmarks include (i) angiogenesis and dilation of blood vessels causing macroscopically visible erythema; (ii) dermal and epidermal infiltration of activated leukocytes (T-lymphocytes, dendritic macrophages, and neutrophils); (iii) hyperproliferation of keratinocytes leading to keratinocyte hyperplasia (acanthosis) and the formation of rete ridges; (iv) retention of nuclei in corneocytes (parakeratosis) related to the atypical keratinocyte proliferation and differentiation; and (v) reduction or absence of the granular layer (hypogranulosis) (Nograles et al., 2010; Bowcock and Krueger, 2005; Lowes et al., 2007; Svensson et al., 2012). The lack of naturally occurring psoriasis in laboratory animals has encouraged the generation of genetically modified animal models with phenotypes reminiscent of psoriasis. Such models have helped demonstrate that vascular changes, epidermal hyperplasia and cell-mediated immunity are inter-related prerequisites for disease development (Schon, 2008; Wagner et al., 2010; Gudjonsson et al., 2007), but no model has been able to capture all histological and clinical traits of psoriasis. The immense concordance between human and porcine skin architecture makes pig models for skin disorders highly attractive for the study of disease development and treatment.

Psoriasis is a multifactorial disorder and a variety of genetically and environmental factors are involved in the etiology of the disease. However, it is evident that both dysregulation of the immune system as well as keratinocyte hyperproliferation are prerequisites for disease development. Hence, psoriasis has been associated with increased levels of inflammatory markers including interleukin (IL)-23, IL-17A and nuclear factor (NF)-κB, but also with markers of keratinocyte homeostasis such as signal transducer and activator of transcription 3 (STAT3) and keratinocyte growth factor (KGF) (Sano et al., 2005; Finch et al., 1997; McKenzie and Sabin, 2003; Nair et al., 2008).

In the epidermis, proliferation is normally confined to the stratum basale, where keratinocyte stem cells are attached to the basal membrane through integrins (Hotchin et al., 1995). Transplanting cells leave the stratum basale and undergo a limited number of cell divisions before they commit to spatiotemporal differentiation as they ascend. Acting as a negative regulator of differentiation, integrin expression is lost in the process (Brakebusch et al., 2000). Hence, cells strongly positive for β1 integrin are indicative of stem cells, whereas cells expressing low levels of β1 integrin are transient-amplifying cells, and β1 integrin-negative cells are post-mitotic, differentiating cells. Notably, transgenic suprabasal expression of integrins promotes oncogenesis, emphasizing their pro-proliferative function (Carroll et al., 1995).

Integrins constitute a superfamily of glycosylated, heterodimeric transmembrane receptors composed of an α- and a β-subunit, exercising functions in cell adhesion as well as inside-out and outside-in signaling. The presence of active α2β1 integrin is normally confined to the stem cell-containing stratum basale, where it binds collagen as part of focal adhesions. In wound healing and psoriatic lesions, however, expression continues into the suprabasal...
layers. On this basis, transgenic mice have been generated that express integrin subunits driven by the involucrin promoter, which is specifically active in suprabasal keratinocytes. Induced skin irritation of such α2β1 transgenic mice led to a chronic condition resembling psoriasis with persistent hyperplasia, cutaneous influx of leukocytes and development of microabscesses, as well as secretion of pro-inflammatory cytokines (Carroll et al., 1995; Haase et al., 2001; Teige et al., 2010).

Suprabasal presence of the β1 integrin subunit has been shown to induce phosphorylation of the epidermal growth factor receptor (EGFR), even in the absence of EGFR ligands, which leads to activation of the cell cycle-stimulating MAPK/ERK pathway (Yu et al., 2000; Moro et al., 1998). Furthermore, the engineered suprabasal expression of β1 integrin triggers release of the potent pro-inflammatory cytokine interleukin 1α (IL-1α) in a ligation-independent manner, which further stimulates the MAPK/ERK pathway (Haase et al., 2001; Hobbs and Watt, 2003).

We have previously described the generation of transgenic pigs featuring ectopic expression of either human integrin α2 or integrin β1 (Staunstrup et al., 2012). These animals presented with an aberrant inflammatory profile in the skin but no visually apparent skin abnormality. Here, we describe the generation and evaluation of double-transgenic Göttingen minipigs (Sus scrofa domestica) featuring suprabasal expression of both human α2 and β1 integrin. Our findings document the development of a severe skin phenotype with evidence of psoriasiform epidermal hyperplasia and inflammation in minipigs engineered with multiple copies of the expression cassette encoding both α2 and β1 integrins.

RESULTS
Production of double-transgenic pigs expressing human ITGA2 and ITGB1

To co-express human integrins α2 and β1, we produced Sleeping Beauty (SB) DNA transposon-based vectors carrying human ITGA2 and ITGB1 genes in a bicistronic expression cassette. Using a CMV promoter to drive expression, production and correct translational processing of α2 and β1 integrin from genomically integrated transposons was verified in mouse fibroblasts (Fig. S1). To generate transgenic pigs with ectopic expression of the two integrins, we exchanged the promoter with the involucrin (INV) promoter, resulting in the plasmid pT2/INV-A2B1 carrying the integrins, we exchanged the promoter with the involucrin (INV) promoter, resulting in the plasmid pT2/INV-A2B1 carrying the

In vitro

animals and, notably, with higher copy numbers in pig TG-410 resulting in massive PCR amplification (Fig. 1C). An estimation of the number of transposon copies in each piglet was achieved by Southern blot analysis (Fig. 1D, Fig. S4). Interestingly, TG-410 contained multiple transposon inserts, presumably over 50 copies. TG-456 harbored only 5 copies, of which one most likely originated from random insertion of the plasmid. Intriguingly, the remaining three piglets displayed identical band patterns with an estimated 8 copies, of which a single insert appeared to be a transposase-independent insertion, suggesting that these three animals arose from the same donor cell clone.

Keratinocytes were expanded in vitro from skin explants for at least 3 weeks, allowing keratinocytes to differentiate and, thus, to switch on the INV promoter. Total RNA was extracted from pooled outgrowths from each of the five transgenic pigs and two age-matched wild-type controls (WT C1 and C2) and subsequently used for RT-qPCR (Fig. 1E). A cycle threshold (Ct) value above background could not be detected in the non-transgenic animals, whereas a marked signal of integrin-encoding mRNA, using both ITGA2 and ITGB1 primer sets, was recorded in all transgenic animals, indicating that the complete polycistronic transcript was being produced. Notably, we observed an apparent coherence between copy numbers and expression levels. Hence, pig TG-456 with the fewest insertions also had the lowest level of transcripts, whereas TG-410 exhibited levels of RNA at least 400-fold higher than TG-456. The three genetically identical animals displayed comparable levels of expression, but at a much lower level than TG-410. In summary, these findings confirmed that all five cloned animals were transgenic and that one multi-copy transgenic pig showed a severe skin phenotype.

Membrane localization of human α2 and β1 integrin in keratinocytes derived from transgenic pigs

Only heterodimeric integrin complexes can accomplish the canonical function of integrins as transmembrane receptors. To track membrane localization of heterologous human α2 and β1 proteins, primary keratinocytes from the transgenic pigs TG-410 and TG-458, as well as from the non-transgenic control WT-C3 were expanded in vitro and then immunostained using primary antibodies against human α2 (P1E6) and β1 (P5D2) integrin and a green fluorophore-conjugated secondary antibody (Fig. 2A). The presence of both exogenous integrins was evident in keratinocytes from both transgenic animals. Furthermore, staining was not observed in P1E6- or P5D2-stained keratinocytes derived from WT-C3. Notably, the more intense staining in TG410-derived keratinocytes compared with cells from TG-458 correlated with the higher mRNA level seen in Fig. 1E. A merged picture of sequentially stained cells derived from the transgenic animals employing a green and red fluorophore-conjugated secondary antibody demonstrated co-localization of the two integrins at the cell membrane (Fig. 2A, right column). Furthermore, it was evident that the fluorescence intensity increased in ascending cells, which is in concordance with the spatiotemporal activity of the INV promoter being limited to differentiation-committed cells. Confocal microscopy of dual-stained keratinocytes derived from TG-459 further supported the notion of membrane anchorage of the ectopically expressed integrins (Fig. 2B). A nearly complete overlap in the localization of α2 and β1 integrin was visible in the cultured keratinocytes. In most cells, transport vesicles that stained positive for both α2 and β1 integrin subunits were visible, further emphasizing a correct cellular function and distribution of exogenous integrin complexes (Fig. S5). Of note, there was no bleed-through of the fluorescent signals.
A third evaluation of the human $\alpha_2$ and $\beta_1$ integrin distribution was made by flow cytometry. A subpopulation of the cultured porcine keratinocytes were harvested and stained with P5D2 or P1E6 without permeabilizing the cell membrane. Given that both antibodies recognize an extracellular epitope, this ensured that only membrane-protruding integrin subunits were marked. The percentage of $\alpha_2$ and $\beta_1$ integrin-presenting cells in the heterogeneous pool of keratinocytes, all at variable stages of differentiation, was significantly higher in cells originating from the transgenic animals compared with cells from the control pig (WT C3), although the $\beta_1$ antibody showed some cross-specificity (Fig. 2C). In accordance, quantitative PCR on cDNA revealed a significant increase in the relative level of the pro-proliferative markers antigen Ki-67 ($Ki-67$) and proliferation cell nuclear antigen ($PCNA$) in the three transgenic pigs compared with the control (P<0.0001) (Fig. 2D). This is in line with previous findings showing that suprabasal epidermal expression of human integrins in mice leads to acanthosis (Carroll et al., 1995) and suggested that homing of transgenic integrins to the membrane was active. In summary, these data support the notion that exogenous integrins were correctly processed and displayed at the cell membrane, allowing interaction with extracellular ligands and endogenously expressed membrane receptors.

**Generation of piglets with severe acanthosis by re-cloning of TG-410**

Owing to the lack of visual phenotypic changes in the skin of low-copy transgenic pigs, we investigated whether chronic skin inflammation could be induced in pigs TG-456 and TG-459 using potent chemical inducers of inflammation. Although epidermal...
Fig. 2. Translation and correct distribution of heterologous α2 and β1 integrin. (A) Representative immunocytochemistry images using human-specific antibodies against α2 (P1E6) and β1 (P5D2) integrin performed on permeabilized cultured primary keratinocytes from two transgenic pigs (TG-410 and TG-458) and a non-transgenic control (WT-C3). The cells were either stained singly or in combination. DAPI was used to stain the nucleus. Detection of integrin was restricted to transgenic cells. (B) Representative images from confocal fluorescence microscopy of saponin permeabilized keratinocytes from pig TG-459 stained with the nucleus marker DAPI along with anti-human antibodies against α2 (P1E6) and β1 (P5D2) integrin. Integrin accumulates at the plasma membrane and especially at cell-cell adhesion sites, suggesting correct processing of the peptide. (C) Flow cytometry analysis of unpermeabilized keratinocytes from transgenic pigs, TG-457 and TG-459, and control pig, WT-C3, stained with α2 (P1E6) or β1 (P5D2) integrin-specific antibodies. Staining solely with the secondary antibody served as a control. Representative dot plots comparing fluorescence from pig WT-C3 and TG-459 are depicted. (D) Quantitative RT-PCR on total RNA extracted from skin biopsies obtained from transgenic pigs TG-456, TG-457 and TG-459, and the non-transgenic age-matched control WT-C2. Relative mRNA levels of Ki-67 and PCNA are depicted normalized to endogenous β-actin. Data presented as mean values±s.d. of biological triplicates where the difference is calculated using the Student’s t-test.
thickness and the transcription level of inflammation markers were increased by the treatment, they reverted to near-baseline levels post treatment (Figs S6,S7). Blood profiling revealed increased numbers of eosinophils and neutrophils and decreased numbers of lymphocytes in the transgenic pigs (Table S1). Interestingly, a moderately higher neutrophil load in peripheral blood was consistently found in psoriatic patients (Zanolli et al., 1989). Also, the lower number of lymphocytes is in agreement with findings in moderate-to-severe psoriasis patients and in α2/β1 transgenic mice (Teige et al., 2010; Lowe et al., 1981; Xia et al., 2003). Together, these findings suggested that low-copy number transgenic pigs were indeed in a state of inflammatory acuteness but at a level inadequate to surpass disease threshold. We therefore chose to focus on pig TG-410, which according to the autopsy had slightly immature but normally positioned internal organs. The lungs of TG-410 were atelectatic in about 70% of the tissue, and agenesis of the gall bladder was evident. The skin was discolored by meconium and showed signs of hyperkeratosis and acanthosis. Based on these findings, we decided to re-clone TG-410. Skin fibroblasts isolated from an ear clip of TG-410 were propagated and used for HMC. Achieving a blastocyst rate of 24%, a total of 92 reconstituted embryos were surgically transferred to a single recipient sow, resulting in five piglets (TG-S1 to TG-S5) delivered at term (Fig. 3A, Table S2, Fig. S3). Unfortunately, the five pigs were either stillborn or perished shortly after birth. They all presented the same phenotypic traits as TG-410 with whole-body thickening of the skin, even on the tongue, primarily attributed to an extensive stratum corneum layer (Fig. 3B). As with TG-410, autopsy of TG-S1 and TG-S2 revealed that the internal organs were normally developed and positioned, but also that the lungs were partially atelectatic. For one of the pigs, TG-S1, serohemorrhagic fluid could be observed in the abdominal and thoracic cavity.

Histological changes in the re-cloned transgenic pigs with psoriasiform characteristics

To investigate the skin pathology in the multi-copy transgenic pigs, paraffin sections of skin biopsies from six transgenic pigs (TG-457, TG-S1 to TG-S5) and three non-transgenic controls (WT-C3, WT-C7 and WT-C8) (Fig. S3) were stained with Hematoxylin and Eosin (H&E; Fig. 4, Fig. S8). The overall architecture of the skin in TG-457 was intact when judged against WT-C3 (Fig. 4A-D), and signs of keratinocyte dysregulation or influx of inflammatory cells were not evident. By contrast, the skin architecture of the multi-copy transgenic animals was grossly altered (Fig. 4E,F, Fig. S8), and the skin of these animals was clearly thickened as a result of hyperkeratosis and acanthosis. The stratum corneum was several folds thicker in the transgenic animals, displaying a seemingly chaotic flaking. The epidermis was also denser in TG-S1 and TG-S4, and peculiarly, the stratum corneum contained hair follicle structures, possibly due to a rapid keratinocyte turnover (Fig. S9). Furthermore, rete ridges (downwards extending epidermal ‘fingers’) were identifiable, often fused to adjacent ridges, whereas no signs of papillomatosis (papillary projections of the epidermis forming an undulating surface) were evident. Magnification of the epidermis (Fig. 4G,H, Fig. S8) revealed a high number of mitotic cells in the
Non-terminally differentiated nucleated cells were also seen in the suprabasal layer (stratum spinosum), supporting the notion of a hyperproliferative state, presumably instigated by the high expression of the heterologous integrins. Interestingly, the stratum granulosum was largely absent (hypogranulosis) in the transgenic pigs, which could be ascribed to the inhibited terminal differentiation of the keratinocytes as a classical hallmark of psoriatic skin. Equally notable was the dilation of vessels in the papillary dermis and the presence of granulocytes, mostly in the form of neutrophils. Multiple non-terminally differentiated nucleated cells were also seen in the stratum corneum (Fig. 4J, Fig. S8), which revealed clear signs of parakeratosis.

Infiltration of immune cells in the skin of multi-copy transgenic pigs

To analyze cutaneous influx of immune cells, skin sections from six transgenic pigs (TG-457 and TG-S1 to TG-S5) plus three non-transgenic age-matched controls (WT-C3, WT-C7 and WT-C8) were stained for cellular markers for T-lymphocytes (CD3), macrophages (CD163) and granulocytes (CD66b), as well as for proliferation (Ki-67) (Fig. 5, Fig. S10). Whereas only a few scattered CD3+ T-lymphocytes were visible in the controls (Fig. 5A), a slightly increased number was evident in the skin of TG-457 (Fig. 5E). In the skin of multi-copy transgenic animals, however, we observed a markedly increased influx of T-lymphocytes along the papillary epidermis and throughout the suprabasal layer (Fig. 5I, Fig. S10). Moreover, staining for macrophages showed a massive infiltrate mostly confined to the dermal layer in all multi-copy transgenic pigs (Fig. 5J, Fig. S10), whereas the control animals and TG-457 showed a baseline presence of macrophages (Fig. 5B,F, Fig. S10). Staining for granulocytes (CD66b) was negative in wild-type animals, as well as in TG-457 (Fig. 5C,G, Fig. S10). However, aggregates of neutrophils were easily detectable in the stratum corneum of TG-S1 to TG-S5 (Fig. 5K, Fig. S10), providing evidence for the chemotactic recruitment of neutrophils characteristic of an ‘acute’ inflammatory state of psoriasis. We also stained skin sections for Ki67 and found scattered proliferating keratinocytes in the stratum basale of the control pigs and TG-457, with only a limited number of transient-amplifying cells present (Fig. 5D,H, Fig. S10). In contrast, basically all cells in the entire stratum basale in sections derived from all five multi-copy transgenic pigs were actively proliferating (Fig. 5L, Fig. S10). Moreover, large sections of the suprabasal layer were Ki67 positive (Fig. 5L), stressing the impact of integrin expression as a stimulator of cell division and a negative regulator of differentiation.

Psoriasiform skin disease, and not dermatitis, in multi-copy ITGB1/ITGA2 transgenic minipigs

Based on Hematoxylin and Eosin-stained and immunostained tissue sections, a cumulative score for characteristic traits of psoriasis and dermatitis was given to three control pigs (WT-C3, WT-C7 and WT-C8) as well as six transgenic pigs (TG-457 and TG-S1 to TG-S5). The sections were blinded, after which they were independently assessed and granted a score for each trait from 0 to 2 (0, absent; 1, present; 2, severe). Seven psoriasis-specific identifiers and four dermatitis-specific markers, as well as two markers common for the two diseases were assessed (Fig. S11). A modest non-significant difference in the psoriasis-specific scoring between the controls and TG-457 was attributed to the partial loss of the stratum granulosum and mild exocytosis in the latter (Fig. S11). Nevertheless, these findings indicate that the expression of the heterologous integrins and not genetic disruption by the transgenic cassette itself was responsible for the observed phenotype. In contrast, all transgenic pigs derived from TG-410 obtained a psoriasis score close to the theoretically highest score, reaching a mean of 13.2 (s.d.=1.7) (Fig. 6). Notably, the deficit to the highest score could be almost exclusively ascribed to the lack of perivascular-located neutrophils in the transgenic pigs (Fig. S11). For comparison, the total score in control pigs was 0.3 (s.d.=0.7). These findings showed that a very consistent psoriasiform phenotype was present in piglets carrying multiple copies of the transgene cassette. In accordance, none of the pigs showed signs of dermatitis, and significant dermatitis-specific...
dissimilarities between transgenic and control animals were not evident. Taken together, a combined list of the interrogated pathological traits characteristic of psoriasis revealed a compelling concordance between the observed changes in the skin of the multi-copy transgenic pigs and expected changes found in human psoriasis (Table 1). In summary, our findings demonstrate that transgenic minipigs expressing high levels of $\alpha_2$ and $\beta_1$ integrin in the skin develop severe skin inflammation showing disease characteristics consistent with a severe psoriasiform condition both at the macroscopic and cellular level.

**DISCUSSION**

With two reported exceptions in monkeys, psoriasis has not been observed in animals other than humans (Zanolli et al., 1989; Lowe et al., 1981). Several transgenic and knockout murine models of psoriasis have been created (Sano et al., 2005; Xia et al., 2003; Groves et al., 1995; Bullard et al., 1996; Pasparakis et al., 2002), but common to all these models is that they fail to completely recapitulate all aspects of the disease in humans. Failure to model the disease in mouse skin may largely be due to crucial differences in skin morphology and immune response between humans and rodents, including keratinocyte turnover rate and differences in the subtypes of skin-homing T-lymphocytes (Berking et al., 2002; Khavari, 2006; Elbe et al., 1996; Ardavín, 2003). Xenotransplantation of human psoriatic skin onto SCID mice, therefore, still ranks as one of the most faithful models of the disease (Haftek et al., 1981; Kundu-Raychaudhuri et al., 2014).

The human-like skin architecture of pigs and the establishment of pig cloning from genetically modified somatic cells have made porcine models for human skin conditions attractive. However, besides the incidental finding of some psoriasiform characteristics in a transgenic pig with constitutive cutaneous expression of the hedgehog transcriptional activator, Gli2 (McCalla-Martin et al., 2010), porcine models of psoriasis-like conditions have not, to our knowledge, been described. We have previously described the generation of transgenic pigs with ectopic expression of either human integrin $\alpha_2$ or integrin $\beta_1$ in the skin (Staunstrup et al., 2012). These minipigs displayed altered cytokine profiles in the skin, but did not develop a visible phenotype reminiscent of psoriasis. Here, we generate double-transgenic pigs featuring...
cell migration, survival and proliferation. Hence, ligated integrins in multi-component complexes at the cell membrane of adherent cells function as negative regulators of apoptosis and convey pro-proliferative signals. Under homeostasis, epidermal expression of integrins is confined to stem cell compartments of the stratum basale, suggesting that keratinocytes ascending through the suprabasal layer rapidly lose their proliferative potential and start to differentiate. Deregulation of epidermal homeostasis and chronic inflammation are characteristics of psoriasis. Notably, keratinocytes constitute a key node in this regard as they both belong to the innate immune system and are the primary cells of the epidermis. In line with this, faulty regulation of keratinocyte fate can lead to augmented release of leukocyte chemoattractants, such as IL-1α. Recruited active leukocytes release additional cytokines, possibly leading to a state of self-sustainable inflammation. We and others have previously shown that suprabasal expression of α2 and β1 integrins in transgenic animals leads to the augmented release of IL-1α (Haase et al., 2001; Staunstrup et al., 2012).

The first round of cloning produced four low copy-number transgenic pigs (TG-456 to TG-459) and a single piglet with >50 copies of the transgene cassette (TG-410), all featuring transgene transcription levels in skin correlating to the number of copies. Whereas the transgenic pigs harboring only 4–8 copies at first sight were indistinguishable from control animals, TG-410 presented with severe skin abnormalities. The correlation between transcription level and phenotypic appearance was a compelling indicator of heterologous integrin functionality. Whereas the nature of the high permissibility of the primary fibroblast that eventually led to TG-410 is unknown, only some of the insertions were mediated by SB transposition. As evaluated by Southern blotting, most of the inserted transgenes were organized in concatamers, probably through transposase-independent mechanisms (Chen et al., 2001; Dai et al., 2010). Expression and colocalization of membrane-homing human α2 and β1 integrins indicated correct processing of the peptides, including maturation of pre-β1 integrin and dimerization in the endoplasmic reticulum (Carroll et al., 1995; Koivisto et al., 1994). Along with the demonstrated expressional regulation of proliferation marker, this suggested that functional integrin dimers were indeed present in the cell membrane of keratinocytes in transgenic pigs.

Re-cloning of TG-410 resulted in piglets that were phenotypically and molecularly identical to TG-410 at birth, suggesting that the phenotypic traits directly reflected the genetic composition of the pigs. Intriguingly, histological assessment revealed traits consistent with psoriasis vulgaris. In all animals, hyperproliferation of basal and suprabasal keratinocytes was evident, leading to the formation of rete ridges, thinning of inter rete and acanthosis. It is important to note that the suprabasal integrin expression is not a corollary of inflammation as intradermal injection of cytokines into normal human skin has been shown not to induce such expression (Hertle et al., 1995). Previous studies have found that ectopic expression of integrins on suprabasal keratinocytes does not directly prompt cells in the stratum basale to enter mitosis (Romero et al., 1999). Rather, the augmented release of IL-1α from keratinocytes stimulates the MAPK/ERK and NF-κB pathways, leading to growth homeostasis and inflammatory responses (Murphy et al., 2000; Kaufman and Fuchs, 2000). We and others have previously reported a higher production of IL-1α from human ITGB1 transgenic primary keratinocytes (Haase et al., 2001; Staunstrup et al., 2012).

The rapid turnover and faulty differentiation of ascending keratinocytes gave rise to parakeratosis, hyperkeratosis and hypogranulosis. An exacerbated release of cytokines and chemokines from the vast number of proliferating keratinocytes

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**Table 1. Concordance between phenotypic traits of human psoriasis and the double integrin transgenic pigs**

<table>
<thead>
<tr>
<th>Psoriasis-observed changes</th>
<th>Phenotype</th>
<th>Human psoriasis</th>
<th>TG-S1 to TG-S5 pigs</th>
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<tr>
<td>Epidermal changes</td>
<td>Acanthosis</td>
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<td>Parakeratosis</td>
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<td>Hyperkeratosis</td>
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<td>Spongiosis</td>
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<td>Loss of stratum</td>
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<td>Thinning inter rete</td>
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<td>Vascular changes</td>
<td>Dilation of vessels</td>
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<td>Perivascular neutrophils</td>
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<td>Inflammatory changes</td>
<td>Cutaneous T-cell infiltrate</td>
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<td>Granulocyte infiltrate</td>
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<td>Eosinophil infiltrate</td>
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<td>Macrophage infiltrate</td>
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<td>Intra-epidermal micro-abscesses</td>
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<td>Injury response</td>
<td>‘Koebner’ phenomena</td>
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Phenotypic changes observed in the transgenic pigs are all in agreement with findings in human psoriasis except for changes in perivascular neutrophils (indicated by asterisk). Question mark indicates that the feature in question was not examined.
Interestingly, as one of several hallmarks of psoriatic skin (Lowes et al., 2007), dermal CD3+ cells clustered in the upper dermis of the transgenic animals. Overall, the release of recruitment factors ensures cutaneous influx of lymphocytes and macrophages, as well as stratum corneum-homing granulocytes forming Munro’s microabscesses (Kim et al., 2014; Mitsui et al., 2012). Expansion of the microvascular system is another hallmark of psoriatic lesions that helps to nourish the hyperproliferating skin and permits massive extravasation of leukocytes. A marked dilatation of tortuous dermal capillaries was palpable in the multi-copy transgenic pigs.

The only psoriasis-specific parameter absent in the skin of the transgenic pigs was the anticipated presence of perivascular neutrophils. However, as opposed to observations commonly found in dermatitis, no spongiosis (intercellular edema between keratinocytes) or orthokeratosis (hyperkeratosis without parakeratosis) could be observed in the skin of the transgenic pigs. Collectively, this argues in favor of a psoriasiform condition. Furthermore, the accumulation of cutaneous neutrophils and macrophages points to a self-perpetuating chronic/acute state of inflammation, as these cells potentiate both the T-cell response and keratinocyte deregulation. This, on the other hand, facilitates activation of the complement system by activated keratinocytes (Nagashima et al., 2004; Matsunari et al., 2013). Our work demonstrates that deregulated keratinocytes can instigate psoriasis (Staunstrup et al., 2012). A cassette encompassing human ITGA2 and 2AP was PCR-amplified from pT2/CMV- hITGA2-SV40-neo using a reverse primer containing the 2AP sequence. Subsequently, ITGA2 was released from pT2/CMV-hITGA2-SV40-neo by BglII and CiaI digestion and substituted with ITGA2-2AP. PCR amplification of ITGA2 was undertaken with pT2/CMV-hITGA2-SV40-neo, subsequently the ampiclon was inserted in CiaI-digested pT2/CMV-hITGA2-2AP SV40-neo producing the pT2/CMV-hITGA2-2AP-hITGB1 SV40-neo construct. The CMV promoter was released by Ascl digestion and replaced with the INV promoter and the first intron amplified from pH-3700-pL2 (kindly provided by Rikke Christensen, Dept of Biomedicine, Aarhus University, Denmark), generating pT2/INV-hITGA2-2AP-hITGB1-bGHPA SV40-neo. pCMV-SBI00X has been previously described (Mátés et al., 2009).

Transposition and generation of donor cells for HMC For stable transfection of NIH3T3 with pT2/INV-hITGA2-2AP-hITGB1-bGHPA SV40-neo or pT2/CMV-hITGA2-2AP-hITGB1-bGHPA SV40-neo, cells were seeded in 6-well plates at a density of 2×105 cells/well. On the following day, the cells were transfected using 0.3 μg SB100X transposase plasmid with 1.2 μg transposon vector mixed with Fugene 6 (Roche, Basel, Switzerland) according to the manufacturer’s protocol using a 3 μl/2 μg ratio. After 24 h, the cells were trypsinized and transferred to a 10 cm dish (Greiner Bio-one GmbH, Frickenhause, Germany). After a further 24 h, the cells underwent G418 (Invitrogen) selection (1 mg/ml) for 14 days. Subsequently, the resistant colonies were isolated and pooled for DNA or protein extraction. Primary male Göttingen fibroblasts were seeded, transfected and selected as described above, using 20 ng SB100X plasmid and 1 μg transposon vector along with Fugene 6 at a 3 μl/2 μg ratio. Pooled transgenic cells were re-seeded in one well of a 6-well dish and allowed to expand until confluency. At confluency, the cells were re-seeded into 4 wells of a 12-well dish at a density of 3×104 cells/well and allowed to expand for 3 days to reach confluency without change of medium, ensuring G1 phase synchronization.

Handmade cloning (HMC) and transfers HMC was performed as described before (Du et al., 2007; Schmidt et al., 2010). Briefly, cumulus cells were removed from matured cumulus-oocyte complexes (COCs) with 1 mg/ml hyaluronidase. After partial digestion of zona pellucida with 3.3 mg/ml Pronase, oriented bisection of oocytes was performed to remove nuclei. Each cytoplast (without polar body) was attached to a single transgenic fibroblast and fused in fusion medium (0.3 M mannitol, 0.1 mM MgSO4 and 0.1 mg/ml PVA) in a fusion chamber (BTX microslide 0.5 mm fusion chamber, model 450; BTX, San Diego, CA, USA) with a single direct current (DC) impulse of 2.0 kV/cm for 9 μs. One hour later, each cytoplast-somatic cell pair was fused with another cytoplast in activation medium (fusion medium with 0.1 mM CaCl2) by a single DC pulse of 0.86 kV/cm for 80 μs. After incubation in porcine zygote medium 3 (PZM-3) supplemented with 5 μg/ml cytochalasin B, 10 μg/ml cycloheximide for 4 h, the reconstructed embryos were cultured in PZM-3 medium for another 5-6 days to develop into transgenic blastocysts. Blastocysts and morulae, at day 5 and 6, were surgically transferred into recipient sows (Schmidt et al., 2010).

Cells The murine fibroblast cell line NIH3T3 and human keratinocyte cell line HaCaT were cultured in Dulbecco’s modified Eagle’s medium, DMEM

et al., 2001). Thus, pig models of skin inflammation should be valuable tools for development and refinement of topical therapies.

**MATERIALS AND METHODS**

**Ethics statement**

All animal procedures were performed in accordance with relevant guidelines and regulations and approved by the Danish Animal Experiments Inspectorate (license no. 2006/561-1230).

**Plasmids and vector construction**

SB transposon-based vector constructs were generated based on the plasmid vector pT2/CMV-hITGA2-SV40-neo and pT2/CMV-hITGB1-SV40-neo, which have been previously described (Staunstrup et al., 2012). A cassette encompassing human ITGA2 and 2AP was PCR-amplified from pT2/CMV-hITGA2-SV40-neo producing a reverse primer containing the 2AP sequence. Subsequently, ITGA2 was released from pT2/CMV-hITGA2-SV40-neo by BglII and CiaI digestion and substituted with ITGA2-2AP. PCR amplification of ITGA2 was undertaken with pT2/CMV-hITGA2-SV40-neo, subsequently the ampiclon was inserted in CiaI-digested pT2/CMV-hITGA2-2AP SV40-neo producing the pT2/CMV-hITGA2-2AP-hITGB1 SV40-neo construct. The CMV promoter was released by Ascl digestion and replaced with the INV promoter and the first intron amplified from pH-3700-pL2 (kindly provided by Rikke Christensen, Dept of Biomedicine, Aarhus University, Denmark), generating pT2/INV-hITGA2-2AP-hITGB1-bGHPA SV40-neo. pCMV-SBI00X has been previously described (Mátés et al., 2009).
Authentication of cell type and monitoring of cell culture contamination was selected against as a result of the low calcium concentration of the medium.

Invitrogen, Paisley, UK). Outgrowth of melanocytes and fibroblasts is (BPE) containing serum-free keratinocyte medium, K-SFM (Gibco, keratinocytes to migrate from the explants. Thereafter, the medium was hydrocortisone (Sigma-Aldrich) at 37°C, 5% CO2 for 7-10 days, allowing keratinocytes to migrate from the explants. Thereafter, the medium was substituted with epidermal growth factor (EGF) and bovine pituitary extract (BPE) containing serum-free keratinocyte medium, K-SFM (Gibco, Invitrogen, Paisley, UK). Outgrowth of melanocytes and fibroblasts is selected against as a result of the low calcium concentration of the medium. Authentication of cell type and monitoring of cell culture contamination was performed by visual inspection.

**Western blot analysis**

NIH3T3 cell lysates were prepared in lysis buffer 9803 (Cell Signaling Technology) containing a cocktail of protease inhibitors (Roche, Basel, Switzerland). Keratinocyte cell lysates were prepared in a lysis buffer containing 50 mM Tris-HCl pH 7.6, 2.5% SDS, 10% glycerol, 10 mM β-glycerophosphate, 10 mM NaF, 95 µM sodium orthovanadate and a cocktail of protease inhibitors (Roche). Total protein content was determined using a BCA Protein Assay Kit (Pierce). Lysate cocktail of protease inhibitors (Roche, Basel, Switzerland) and incubated bottom-up overnight at 37°C, 5% CO2.

**PCR and RT-qPCR**

PCR was performed on 100 ng gDNA extracted from primary fibroblasts originating from the transgenic pigs or non-transgenic pig controls. Primers specific for the neomycin selection marker (500 bp product), ITGB1 (280 bp product) and ITGA2 (175 bp product) transgenes, and for the SB100X transposase (390 bp product) were used in a standard Taq polymerase (Ampliqon, Skovlunde, Denmark) PCR assay. Reactions were run for 35 cycles and visualized on a 1.5% agarose ethidium bromide stained gel. For qRT-PCR, 150-300 ng total RNA (RNaseasy kit, Qiagen, Hilden, Germany) from transfected NIH3T3, Göttingen primary keratinocytes and from skin biopsies was DNase treated (DNA-free, Ambion, TX, USA) and used for standard 20 µl cDNA synthesis (AffinityScript cDNA synthesis kit, Stratagene). Of total cDNA, 1:20 to 1:10 was used per qPCR reaction. Q-PCR analyses were performed according to the manufacturer’s instructions with a LightCycler 400 (Roche). The following TaqMan gene expression assays (Applied Biosystems) were deployed; human ITGA2 detection (Hs00158127), human ITGB1 (Hs00559595).

**Southern blot analysis**

Between 11 and 15 µg of column-purified genomic DNA was digested overnight at 37°C in a total volume of 200 µl with the following restriction enzyme combinations; EcoRV+PvuII or EcoRV+Sall (New England Biolabs). The digested DNA was precipitated with sodium acetate and resuspended in 20 µl TE. The DNA fragments were resolved on a 0.8% agarose gel overnight at 1 V/cm and transferred to a nylon membrane (Amersham Hybond-N+, GE Healthcare) by vacuum suction. During transfer, the DNA was nicked, denatured and neutralized in 0.25 M HCl, 1.5 M NaCl+0.5 M NaOH and 1.5 M NaCl+1 M Tris-HCl pH 7.6, respectively. Subsequently the membrane was washed with 20×SSC. The DNA was UV-crosslinked to the membrane and pre-hybridized with 0.4 mg/ml salmon sperm ssDNA (D1626, Sigma-Aldrich) in 5×SSPE (0.75 M NaCl, 0.05 M NaH2PO4, 0.005 M EDTA), 5×Denhart’s (0.1% Ficoll 400, 0.1% polyvinylpyrrolidone, 0.1% BSA), 1% SDS, 50% formamide, 5% dextran-sulfate for 4 h at 42°C. A neomycin probe of 700 bp was CTP 32P-labeled (Prime-It II random primer labeling kit, Stratagene) and mixed with salmon sperm ssDNA (1 mg/ml) and hybridized buffer and finally incubated with the membrane at 42°C overnight. The membrane was washed in 2×SSPE for 2×5 min at 26°C and in 2×SSPE/0.5% SDS for 2×15 min at 53°C and finally in 0.2×SSPE/5% SDS for 1×5-15 min at 53°C. The membrane was air-dried and enclosed in a cassette for exposure of an X-ray film (Konica Minolta Medical and Graphic).

**Immunocytochemistry**

Skin explants from α2β1 transgenic and non-transgenic pigs were placed in slideflasks from which primary keratinocytes were expanded. After 2 weeks, the explants were removed and the cells fixed in 4% formalin for 5 min, washed in TBS+0.05% Tween 20 (TBS-T) and blocked in 1% BSA, 0.3% Triton X-100 in PBS for 30 min at ambient temperature. Subsequently, the cells were incubated overnight at 4°C with mouse monoclonal antibodies against human β1 integrin (PSD2, kindly provided by Uffe Birk Jensen, Dept of Biomedicine, Aarhus University, Denmark) at a 1:100 dilution or mouse monoclonal antibodies against human α2 integrin (Santa Cruz Biotechnology, P1E6) at a 1:50 dilution with mouse monoclonal antibodies against human α2 integrin (Santa Cruz Biotechnology, P1E6) at a 1:50 dilution or mouse monoclonal antibodies against human α2 integrin (Santa Cruz Biotechnology, P1E6) at a 1:50 dilution or mouse monoclonal antibodies against human α2 integrin (Santa Cruz Biotechnology, P1E6) at a 1:50 dilution or mouse monoclonal antibodies against human α2 integrin (Santa Cruz Biotechnology, P1E6) at a 1:50 dilution.

**H&E staining**

Formalin-fixed paraffin-embedded (FFPE) blocks of skin from transgenic and non-transgenic pigs were sectioned in 4 µm slices deparaffinized in xylene and rehydrated. The skin samples were placed in 0.1% Mayer’s Hematoxylin (Sigma-Aldrich) for 15 min and rinsed in copious amounts of tap water. The slides were then successively dipped in 0.5% Eosin for 2 min and in ddH2O until the Eosin stopped streaking. The samples were dehydrated in increasing concentrations of ethanol and lastly in xylene before being mounted with Clarion Mounting Medium (Sigma-Aldrich).

**Immunohistochemistry**

Immunohistochemical staining was performed on Benchmark XT (Ventana Medical Systems, Tucson, AZ, USA) by an indirect sequential immunoenzymatic technique. Sections of 4 mm were cut from each paraffin-embedded tissue block, mounted on Superfrost Plus slides (Thermo Fisher Scientific), and dried for 5 min at 60°C. Standard settings and reagent kits of Benchmark XT (Ventana Medical Systems) were used in deparaffinization, rehydration, antigen retrieval and endogenous peroxidase blocking steps. Rabbit anti-CD3 monoclonal antibodies (ready to use, Ventana Medical Systems, 790-4341), mouse anti-CD66b monoclonal antibodies (1:600, BD Biosciences, 555723), mouse anti-CD163 monoclonal antibodies (1:450, Serotec, MCA 1853) and rabbit anti-Ki-67 monoclonal antibodies (1:200, Cell Marque, 275R-16) were incubated for 20 min (anti-CD163) or 32 min (anti-CD3, anti-CD66b and Ki-67) at room temperature followed by Ventana ultraView Universal 3,3-diaminobenzidine (DAB) Detection Kit that includes HRP-conjugated.
anti-mouse and anti-rabbit antibodies. Slides were counterstained with Mayer’s Hematoxylin and bluing reagent and later manually dehydrated and mounted.

Confocal microscopy
Transgenic or control pig keratinocytes were seeded at a density of 10^{5} chamber in an 8-well poly-L-lysine coated 1 µm slide plate (Ibidi, Munich, Germany). After incubation for 24 h, the cells were fixed, permeabilized and stained with mouse anti-human monoclonal antibodies P5D2 or mouse anti-human monoclonal antibodies P1E6, followed by incubation with the secondary Alexa Fluor 488 goat anti-mouse IgG Ab as described above. The cells were visualized using a 488 nm line of a multiline argon laser (detection of Alexa Fluor 488) and the 405 nm line of a 405-430 nm diode laser (detection of DAPI) in a confocal laser scanning microscope (LSM 710, Zeiss) using 63× oil-immersion objective with a numerical aperture of 1.4.

Flow cytometry
Transgenic and non-transgenic primary porcine keratinocytes expanding from skin explants were harvested by trypsinization and fixed in 4% formalin for 10 min. After washing with cold 0.5% TBS-T, cells were permeabilized and blocked in PBS containing 0.2% saponin (Sigma-Aldrich) and 0.5% BSA at 4°C, 30 min. All samples were washed in 0.5% TBS-T and stained for αβ1 integrin (P1E6, 1:50) or β1 integrin (P5D2, 1:100) for 4°C, 60 min. Hereafter, the cells were washed in 0.5% TBS-T and subsequently incubated with the highly cross-adsorbed Alexa Fluor 488 goat anti-mouse IgG (Invitrogen, A11029) at 1:400 and 4°C for 60 min. The cells were washed and resuspended in 300 µl PBS (no MgCl2, no CaCl2). The permeabilized and blocked in PBS containing 0.2% saponin (Sigma-Aldrich) and 0.5% BSA at 4°C, 30 min. All samples were washed in 0.5% TBS-T and stained for αβ1 integrin (P1E6, 1:50) or β1 integrin (P5D2, 1:100) for 4°C, 60 min. Hereafter, the cells were washed in 0.5% TBS-T and subsequently incubated with the highly cross-adsorbed Alexa Fluor 488 goat anti-mouse IgG (Invitrogen, A11029) at 1:400 and 4°C for 60 min. The cells were washed and resuspended in 300 µl PBS (no MgCl2, no CaCl2). Finally, 10,000 events were analyzed on a BD FACS diversion machine using BD FACSDiva and FlowJo software.

Epidermal thickness
The mean epidermal thickness (epidermal area divided by epidermal length) was calculated using the Visiopharm Integrator System (VIS) (Visiopharm, Hørsholm, Denmark).

Sensitization
A shaved area on the left dorsal flank of the pigs was sensitized with 200 µl of 10% fluoro-2,4-dinitrobenzene (DNFB; Sigma) on day 0. Following sensitization, tissues were treated three times a week with 200 µl of 1% DNFB until day 43. Five-mm punch biopsies were collected from untreated skin at day 0 and from the sensitized area on day 37, 44, 51, and, in the first round of experiments, on day 58. One half of the biopsy (cut longitudinally) was kept in 4% formalin and the other half was kept in 1 ml RNAlater (Sigma). Gene expression analyses were performed on skin biopsies from the area on the back exposed to DNFB. RNA was purified using Qiagen RNaseq Lipid Tissue Mini Kit. cDNA was synthesized using an Applied Biosystems High-Capacity cDNA Reverse Transcription Kit. Gene expression analyses were performed using pre-designed Applied Biosystems TaqMan assays according to the manufacturer’s instructions; porcine CXCL10 (S03391846), porcine PCNA (S03377029), porcine STAT1 (S03392291), and porcine STAT3 (S03388439).

Histological score
H&E-stained tissue sections were blinded and evaluated by two observers, of which one was a trained dermatopathologist, to compare the histological features of psoriasis and dermatitis. The following characteristic psoriatic parameters were assessed: pannkeratosis, neutrophil infiltration in epidermis, increased epidermal thickness (acanthosis), absence of stratum granulosum, psoriasis-like papillary pattern (rete ridges), thinning of the epidermis over the dermal papilla, dilatation of vessels in the dermal papilla and presence of neutrophils outside the vessels in the dermal papilla. The following characteristic dermatitis parameters were assessed: orthokeratosis, retained stratum granulosum, acanthosis, spongiosis, exocytosis and occurrence of eosinophil leukocytes. The histological score for each pattern was graded on a scale from 0 to 2 (0, absent; 1, present; 2, severe) (Nagashima et al., 2004).

Statistics
P-values were calculated by a two-tailed Student’s t-test or by Fisher’s exact test, where appropriate, to test the null hypothesis of no difference between the compared groups. The assumption of equal variances was tested by the F-test. In all statistical analyses, P<0.05 was considered significant.

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Competing interests
The authors declare no competing or financial interests.

Author contributions

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Supplementary information
Supplementary information available online at http://dmm.biologists.org/lookup/doi/10.1242/dmm.028662.supplemental

References
Supplementary Figure S1: Effective transcription from genome integrated integrin vectors. (A) Relative expression level of human α2 and β1 integrin in T2/CMV-A2B1 stable transfected compared to un-transfected NIH3T3 cells. (B) Western blot with antibodies against human α2 or β1 integrin on total cellular protein extracts from T2/CMV-B1 or T2/CMV-A2B1 harboring NIH3T3 cells compared to un-transfected cells.
Supplementary Figure S2: Cellular expression of heterologous transposon inserted vectors harboring a bicistronic α2 and β1 integrin cassette. Inter-correlation of the relative expression level of endogenous and transgenic α2 and β1 integrin in T2/INV- A2B1 stable transfected or un-transfected NIH3T3 cells. mITGA2 and mITGB1, murine integrin α2 and β1, respectively; hITGB1 and hITGA2, human integrin α2 and β1, respectively. Asterisk (*) indicates statistical significance with p<0.05.
Supplementary Figure S3: Overview of the animals interrogated.
Supplementary Figure S4: Genotyping of the five cloned pigs TG-410, TG-456, TG-457, TG-458, and TG-459 by Southern blot. Black arrows indicate transposase-mediated genomic insertions and blue arrows random integration, presumably as concatemers.
Supplementary Figure S5: Intracellular co-localization of exogenous integrin α2 and β1 in primary keratinocytes from transgenic pigs. Representative confocal fluorescence microscopy images of keratinocytes cultured from the transgenic pig TG-459 stained with nucleus staining-dye DAPI and with anti-human β1 (P5D2) and α2 (P1E6) integrin mAb. Single channel images and a merged image are shown.
Supplementary Figure S6: Increase in epidermal thickness after DNFB provocation. (A) Variation in epidermal thickness measured in three non-transgenic pigs (WT-C4, WT-C5, and WT-C6) and in two transgenic TG pigs (TG-456 and TG-459) and in the hiTGA2 transgenic pig (TG-555) upon application of the irritant 2,4-dinitro-1-fluorobenzene (DNFB) three times weekly. Baseline epidermal thickness is measured at day 0 prior to sensitization with DNFB. Provocation with DNFB is undertaken up to day 43 and epidermal thickness is measured in biopsies taken at day 37, 44, 51 and 58. Enlargement (about 2-fold) of the epidermis was evident in all animals following sensitization with DNFB but reversed to baseline within 14 days post treatment (day 58). Gray line marks the termination point for DNFB provocation. (B) Repeat of (A) using same conditions and application area. Pig WT-C4 was not included and measurement of epidermal thickness was performed only at day 0, 37 and 51. (C) Baseline epidermal thickness in the six interrogated animals measured at day 0 prior to provocation with DNFB. Notably, the transgenic pigs had thinner epidermis than the wildtype pigs. Data shown are mean values ± standard deviation. (D) Histological sections (MT staining) of skin biopsies from non-transgenic pig WT-C6 and the transgenic pigs TG-555 and TG-459. Top panel: Sections from unprovoked skin (day 0). Lower panel: Sections from skin collected at day 37. Magnification: x4.
Supplementary Figure S7: The relative expression level of transgenes, inflammatory and proliferative markers is influenced by the treatment with DNFB. Total RNA was extracted from skin biopsies taken from non-transgenic pigs (WT-C4, WT-C5, and WT-C6), transgenic TG pigs (TG-456 and TG-459) and hITGA2 transgenic pig (TG-555) at the indicated time-points. Quantitative RT-PCR was performed and normalized to endogenous beta-actin. Relative expression of (A) signal transducers and activators of transcription 1 (STAT1), (B) STAT3, (C) chemokine ligand 10 (CXCL10), (D) proliferation cell nuclear antigen (PCNA) is depicted as a function of time. (A-D) mRNA level indicated is relative to day 0 for pig WT-C4. Gray bars mark the termination point for DNFB provocation. All data points are shown as mean values ± standard deviation.
Supplementary Figure S8: Skin from multi-copy transgenic pigs exhibits parakeratosis, leukocyte infiltration and hyperkeratosis
Supplementary Figure S9: Hair follicle structures are driven out into the stratum corneum in the muli-copy transgenic animals. H&E staining of cross-sectioned skin biopsies from the re-cloned transgenic pigs TG-S1 and TG-S4. At hair dense skin areas, follicle structures are erroneously pushed out into the stratum corneum (arrow 1), likely attributed to the rapid turnover rate of keratinocytes.
Supplementary Figure S10: Cutaneous infiltration of T-lymphocytes, macrophages and granulocytes in addition to epidermal hyperproliferation as demonstrated by immunohistopathology.
Supplementary Figure S11: Histological score across psoriasis and dermatitis specific markers portrayed a psoriasis pattern in the null-copy transgenic pigs. Scores between 0 and 2 (0 = absence, 1 = mild, 2 = severe) were assigned to 2-4 cross-sectional slides of skin from each animal.
**Supplementary Table S1: Increased leukocyte numbers after DNFB provocation.**

<table>
<thead>
<tr>
<th>Unit (bill./L)</th>
<th>Control animals</th>
<th>hiTGA2</th>
<th>hITGA2-hITGB1</th>
<th>TG animals</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Day 0</td>
<td>Day 44</td>
<td>Day 0</td>
<td>Day 44</td>
</tr>
<tr>
<td>Basophils</td>
<td>0.02</td>
<td>0.03</td>
<td>0.01</td>
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<tr>
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<td>0.09</td>
<td>0.05</td>
<td>0.10</td>
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<tr>
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<tr>
<td>Monocytes</td>
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<tr>
<td>Lymphocytes</td>
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<td>4.48</td>
<td>5.22</td>
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**Reference intervals (bill./L)**

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<tr>
<th></th>
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<th>Day 44</th>
</tr>
</thead>
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<tr>
<td>Leukocytes</td>
<td>6.12-8.94</td>
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<tr>
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<tr>
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<tr>
<td>Lymphocytes</td>
<td>3.6-5.46</td>
<td>4.36-5.8</td>
</tr>
</tbody>
</table>

**Supplementary Table S1: Increased leukocyte numbers after DNFB provocation.** Cell counts of leukocyte subtypes from the three non-transgenic pigs (WT-C4, WT-C5, and WT-C6), the hITGA2 transgenic pig (TG-555) and the two hITGA2 and hITGB1 transgenic pigs (TG-456 and TG-459) at day -6 prior to DNFB provocation and at day 38. A reference table indicates the range of normal cell densities based on the control animals. Yellow text marks values below the reference index, green text within the reference index, and red text above the reference index.
### Supplementary Table S2: Summary of cloning efficiencies obtained with INV-A2B1 transgenic Göttingen fibroblasts.

<table>
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<tr>
<th>Donor cell ID</th>
<th>Recipient sow ID</th>
<th>Blastocysts-rate (%)</th>
<th>No, of blastocysts / morulae transferred</th>
<th>Developmental stage (no, / day)</th>
<th>Born/Alive</th>
<th>Piglet ID</th>
</tr>
</thead>
<tbody>
<tr>
<td>Pool of stable INV-hITG2-2AP-hITGB1 Göt PF cells</td>
<td>2175</td>
<td>36%</td>
<td>70</td>
<td>Blastocysts: 60 / 6 10 / 5</td>
<td>5/0</td>
<td>410, 456, 457, 458, 459</td>
</tr>
<tr>
<td>Primary fibroblasts derived from pig 410</td>
<td>2342</td>
<td>24%</td>
<td>92</td>
<td>Blastocysts: Morulae: 39 / 6 26 / 5 2 / 6 25 / 5</td>
<td>5/0</td>
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