

Animal models of psoriasis: a critical appraisal

Michael P. Schön

Department of Dermatology, University of Göttingen, Göttingen, Germany

Correspondence: Prof. Dr. Michael P. Schön, Department of Dermatology and Venereology, University of Göttingen, Von-Siebold-Str. 3, 37075 Göttingen, Germany, Tel.: +49551396401, Fax: +49551396841, e-mail: michael.schoen@med.uni-goettingen.de

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Abstract: Although there is no naturally occurring disorder in laboratory animals that mimics the complex phenotype of psoriasis, a large number of spontaneous or genetically engineered mutations in rodents, immunological reconstitution approaches or xenotransplantation models have shed light on specific aspects implicated in the pathophysiology and therapy of psoriasis. Animal models have helped to elucidate functions of inflammatory mediators or to unravel the contribution of innate or adaptive immune mechanisms, keratinocytes or endothelial cells to chronic hyperproliferative inflammatory skin disorders. However, given that several distinct manipulations of molecular pathways, resident cutaneous cell types or immigrating immunocytes result in remarkably similar phenotypes in experimental animals, it appears that interfering with cutaneous

homeostasis in general may ultimately initiate a rather uniform reaction pattern that mirrors some features of psoriasis. This limitation of animal models generated without the use of human material may, at least in part, be overcome by xenotransplantation of human skin onto immunocompromised animals. The latter approach has been employed in preclinical investigations to study the role of immune cells and/or to predict the efficacy of some therapeutic compounds. This brief review delineates approaches to generate animal models of psoriasis and discusses their strengths and limitations for psoriasis research.

Key words: animal models – psoriasis – skin inflammation – transgenic mouse

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Setting the stage

Psoriasis is a very popular disease among skin researchers. A quick glance at publications related to the pathogenesis of psoriasis over the past four decades may suffice here to substantiate this notion (more than 11 000 papers published from 1965 to 2007 with an ever-increasing tendency; Fig. 1a). Having stated this, one may be somewhat surprised to find a number of central questions in psoriasis research not conclusively answered yet. For example, we still do not have a comprehensive notion on the relative contributions of resident cutaneous cells versus immigrating immune cells, and even the relative importance of subsets of immunocytes (i.e. innate versus adaptive immunity) in the disease process is not entirely clear yet (1–4). Likewise, although several psoriasis susceptibility loci have been identified in the last decade, the genetic basis of psoriasis is still not completely understood (5–7).

In this situation, experimental animal models that mimic the human disease would come in very handy with respect to both researches into the pathogenesis of psoriasis and drug development. However, despite decades of devoted research an animal model that completely resembles the complex human disease without the use of human material, as opposed to animal models, which are based on

transplantation of human tissue or cells, is not yet available (8,9). At least three aspects generally inherent to complex disorders, which are not mutually exclusive, may be considered as explanations for this situation: the first obvious approach to generate an animal model of psoriasis is to somehow tamper with normal skin through targeting of a key molecule or cell type to induce the desired pathological alterations. However, it appears that there is no single feature that is truly unique to psoriasis, for example, metabolic and biochemical abnormalities are found in psoriasis and other inflammatory skin disorders, dysregulation of any given inflammatory mediator or adhesion molecule may also be found in diseases unrelated to psoriasis, genetic markers show only statistical associations with psoriasis, and established therapies are not exclusively effective in psoriasis. Thus, identifying the right candidate target molecules or cells within a plethora of intertwined factors appears difficult, if possible at all. Second, the alternative or complementary approach would be to specifically eliminate (target) candidate cells or mediators in animal models in order to identify crucial elements of the pathogenic cascade. However, as outlined below, this approach also bears some imponderable challenges, as targeting different molecules or cells in a given model may similarly influence the phenotype and, therefore, may not convincingly allow

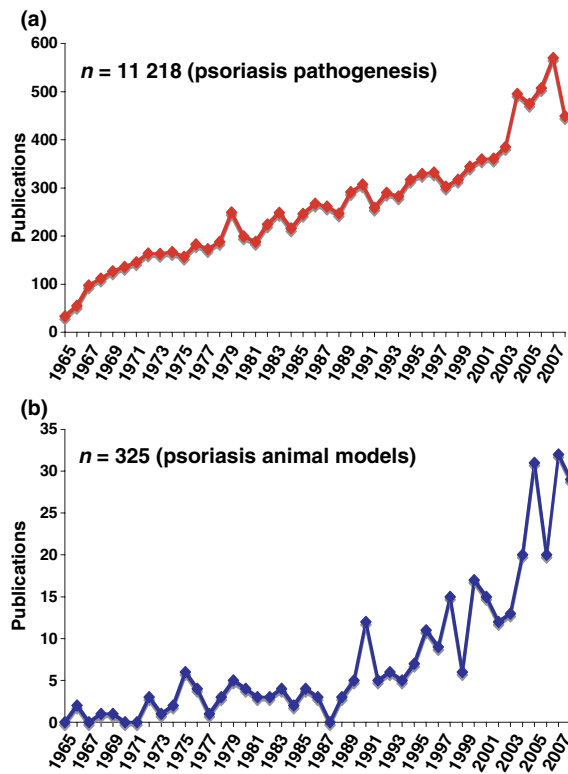


Figure 1. Publications related to the pathogenesis of psoriasis (a) and animal models of psoriasis (b) (resulting from a Medline search in November 2007) show consistently increasing numbers over the past four decades.

identifying the primary pathogenic components. Finally, with the possible exception of a few anecdotal reports on spontaneously occurring conditions with psoriasis-like features in some non-human primates, dogs and pigs (10–13), which occurred sporadically, thus precluding systematic investigations, there is no known naturally occurring disorder in non-human species that exhibits all of the pathologic alterations seen in psoriasis (i.e. chronic inflammatory erythroscaly skin lesions underlied by epidermal hyperproliferation, altered differentiation, angiogenesis and a psoriasis-like infiltrate) and which responds to antipsoriatic therapy. It is, therefore, not easy to predefine the setting in which psoriasis can be studied successfully in an animal model.

Approaches to mimic features of psoriasis in animal models

Despite the aforementioned *a priori* challenges, many publications (whose numbers are steadily increasing; Fig. 1b) have addressed psoriasis-related issues in animal models during the past four decades. On one hand, these approaches have vastly increased our understanding of

various mechanisms of chronic inflammation, which are thought to be relevant for psoriasis. On the other hand, they also highlight some as yet unresolved issues in psoriasis research, e.g. the contribution of the epidermis/keratinocytes, innate and adaptive immunity to the pathogenesis of psoriasis and to drug development. However, as discussed below, such models may ultimately not suffice to resolve those issues. There are several principle approaches to animal models of psoriasis whose specifics have been extensively reviewed (8,9).

Spontaneous mutations

The first psoriasis models described were spontaneous mutations in mice, which are associated with a more or less psoriasis-like phenotype on certain genetic backgrounds and allelic mutations. Spontaneous mutations with a psoriasiform phenotype include mice homozygous for the asebia (*Scd1^{ab}/Scd1^{ab}*) (14), chronic proliferative dermatitis (*Sharpin^{cpdm}/Sharpin^{cpdm}*) (15,16) and the flaky skin (*Ttc7^{sn}/Ttc7^{sn}*) mutation (17,18). The asebia mouse mutation exhibits moderate epidermal acanthosis, increased dermal vascularity and a dermal infiltrate composed of macrophages and mast cells, but neither T cells nor neutrophils. In contrast, both chronic proliferative dermatitis and flaky skin mice exhibit epidermal hyperproliferation, a mixed inflammatory infiltrate with neutrophils accumulating in epidermal microabscesses, and increased dermal vascularity with dilated blood vessels. Backcrosses to different mouse strains suggested that several modifier genes affect the phenotype, including the composition of the inflammatory infiltrate. The flaky skin phenotype is very complex and comprises aspects not present in psoriasis. The *Sharpin^{cpdm}/Sharpin^{cpdm}* mutation results in a similarly complex phenotype, affecting several organ systems. Both phenotypes, chronic proliferative dermatitis and flaky skin, appear to develop independent of T cells, and both disorders do not respond adequately to some antipsoriatic therapies (19–21).

Genetically engineered animals

More specific genotype–phenotype studies can be performed in transgenic animals. This approach has been employed to investigate the role of numerous adhesion molecules, cytokines, transcription factors and other mediators in the pathogenesis of hyperproliferative inflammatory skin disorders. A frequently used approach is epidermal overexpression of molecules of interest under the control of promoters acting in basal (e.g. keratin 14) or suprabasal keratinocytes (e.g. involucrin or keratin 10). Among the target molecules studied in such transgenic approaches are transforming growth factor (TGF)- α (22), interleukin (IL)-6 (23), keratinocyte growth factor (KGF) (24), IL-1 α (25), latent human TGF β (26), vascular endothelial growth

factor (VEGF) (27,28), interferon (IFN)- γ (29), bone morphogenic protein (BMP)-6 (30), the angiogenesis-related Tie2 (31), amphiregulin (32), p40 (the common subunit of the proinflammatory cytokines IL-12 and IL-23) (33,34), IL-20 (35), collagenase (36), MEK1 (a MAP kinase upstream of Erk) (37), or human α_2 , α_5 and β_1 integrin subunits (38). The opposite strategy, i.e. deletion (or diminution) of putative key molecules, was followed to study leukocyte β_2 integrins (CD18) (in a hypomorphic mutation in PL/J mice) (39), IL-1ra (40), IRF-2 (41) or integrin α_E (42). In addition, the role of signal transduction in psoriasis-form skin inflammation was studied by deleting the inhibitor of nuclear factor (NF)- κ B-kinase 2 (IKK2) (43), signal transducer and activator of transcription 3 (Stat3) (44), or Jun proteins (45) within the epidermis.

Immunological approaches

Based on the assumption that immune mechanisms underlie the pathogenesis of human psoriasis, several rodent models have focused on primary immunological manipulations to generate psoriasis-form phenotypes. Along this line, adoptive transfer of CD4⁺ T cells from human HLA-B27/ β 2m-transgenic rats (46) into non-transgenic immunocompromised rats (*Foxn1^{tmu}/Foxn1^{tmu}*) was sufficient to induce psoriasis-like skin alterations in the recipients (47). In addition, a psoriasis-form disorder was observed in *Prkdc^{scid}* mice receiving CD4⁺/CD45RB^{hi} T cells from major histocompatibility complex (MHC)-matched, but minor histocompatibility mismatched donors (48,49). Recombinase activating gene (*Rag*)-2-deficient mice reconstituted with CD4⁺/CD45RB^{hi} T cells developed a similar phenotype (50), indicating that psoriasis-form skin lesions can, in principle, be induced by T cells alone.

Strengths and weaknesses of animal models of psoriasis

A number of important questions in psoriasis research have been addressed by the above mentioned rodent models, and the authors of the respective papers usually conclude that their findings may contribute to the resolution of important issues in the human disease. Few such questions have been outlined briefly.

Epidermal pathogenesis of psoriasis?

The decade-long discussion whether the epidermal alterations observed in psoriasis are cause or consequence of the chronic inflammation (51) has been fuelled prominently by animal models. However, as exemplified by some selected models, the results appear to be somewhat disparate: when mice overexpressing IL-6 (23), KGF (24) or TGF α (22) under the control of the keratin 14 promoter were studied, keratinocyte proliferation and

differentiation, inflammatory cells or angiogenesis within the skin were found to be affected either not at all or only marginally. When IL-1 α was assessed in a similar way, chronic inflammatory lesions were occasionally observed depending on the level of transgene expression (25), while involucrin promoter-driven expression of IFN γ resulted in epidermal hyperproliferation, enlarged dermal capillaries, and induction of MHC class II and intercellular adhesion molecule (ICAM)-1 (29). However, no amelioration of cutaneous inflammation was observed in IFN γ -deficient mice in a more recent study (52). Transgenic mice with low levels and patchy expression of BMP-6 exhibit marked epidermal hyperproliferation along with an inflammatory infiltrate, including epidermal microabscesses (30). The phenotype of mice expressing some human β 1-integrins within the suprabasal epidermis (controlled by the involucrin promoter) resembled psoriasis in several aspects, including epidermal recruitment of CD8⁺ T cells (38). Some recent studies indicate that altered signal transduction within keratinocytes may cause a psoriasis-like phenotype in mice: mice with an epidermis-specific deletion of inhibitor of NF- κ B kinase 2 (IKK2) develop a T cell-independent, psoriasis-like skin disease (43). Another study highlights the importance of JunB, the expression of which is substantially reduced in lesional psoriatic skin. C-Jun, a proposed antagonist of JunB, shows the opposite expression pattern in psoriasis (45) as well as in some hyperproliferative skin disorders in mice, such as the psoriasis-form disease caused by reconstitution of *Prkdc^{scid}* mice with CD4⁺/CD45RB^{hi} T cells (Fig. 2) (49). It was found that mice lacking either gene alone showed a normal phenotype, whereas double knockout mice exhibited a psoriasis-like skin disease. Finally, when Stat3 was constitutively active within the epidermis of mice, a chronic hyperproliferative inflammatory skin disorder developed. However, this psoriasis-like disorder was dependent on T lymphocytes (44).

Pathogenic role of the cutaneous vasculature?

Increased and altered vascularisation and angiogenesis, which is an integral part of the psoriatic phenotype, is featured by several of the spontaneous or genetically engineered mutant rodents as part of their inflammatory phenotype. Again, transgenic approaches were taken to clarify whether angiogenesis-related factors were sufficient to initiate psoriasis-form inflammation. Indeed, targeting expression of VEGF to the epidermis resulted in some psoriasis-like features (27,28). In addition, transgenic mice overexpressing Tie-2 stimulating cutaneous angiogenesis exhibit a chronic hyperproliferative inflammatory skin phenotype (31).

Therefore, some of the animal models mentioned thus far support the hypothesis that under certain conditions,

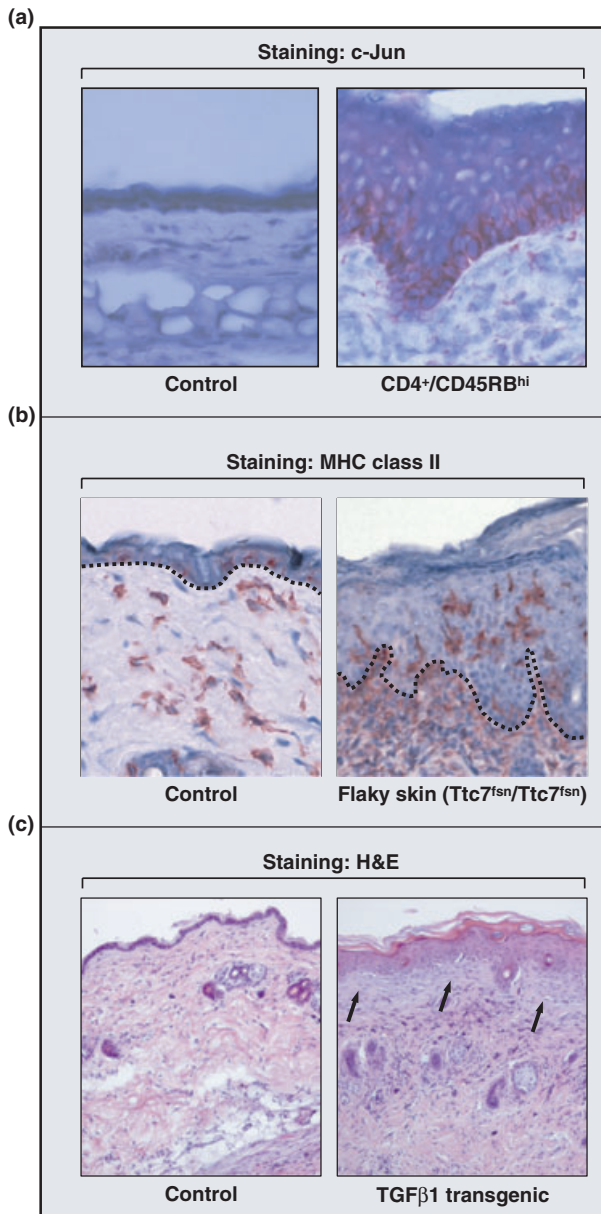


Figure 2. Similarities and differences of animal models compared with human psoriasis. The immunocompromised recipients of CD4⁺/CD45RB^{hi} T lymphocytes (a), flaky skin mice (b) and transgenic mice with epidermal expression of latent TGFβ1 (c) all show a hyperproliferative inflammatory cutaneous phenotype displaying some features of human psoriasis. The examples shown highlight that some features are shared between human psoriasis and certain animal models [e.g. epidermal induction of c-Jun in (a)], while others may differ, such as the lacking induction of MHC class II on keratinocytes [only epidermal Langerhans cells are MHC class II positive in flaky skin mice; the dashed lines indicate the location of the dermo-epidermal junction; (b)] or the development of fibrosis in the TGFβ1-transgenic mice [indicated by arrows in (c)].

primary cutaneous abnormalities (either within the epidermis or the vascular system) are sufficient to initiate a pathogenic cascade, leading to a psoriasis-like phenotype.

However, while important insights into the *in vivo* action of such key molecules within the skin have been gained, tampering with a single cytokine or adhesion molecule apparently did not elicit the complete phenotype of psoriasis, thus supporting the notion of a cytokine network instead (53). In addition, all of the phenotypes mentioned thus far resulted from aberrant expression of gene products within the epidermis, a possibly relevant factor that will be discussed below.

Adaptive immunity and psoriasis?

A substantial body of clinical and experimental evidence suggests that T cells are key effectors in psoriasis (1,2,4). Indeed, adoptive transfer of T cells was sufficient in some animal models to induce a psoriasis-like skin disorder without primary epithelial abnormalities: transgenic expression of human HLA-B27/β2m in rats resulted in psoriasis-form skin changes and arthritis in a subset of animals (46). However, transfer of CD4⁺ T cells from these animals into non-transgenic *Foxn1^{tmu}/Foxn1^{tmu}* rats was sufficient to induce psoriasisform skin alterations in the recipients (47). In addition, induction of a psoriasis-like phenotype was achieved in immunodeficient mice receiving CD4⁺/CD45RB^{hi} T cells from MHC-matched, but minor histocompatibility mismatched donors (49). Similar to psoriasis, IL-12 and IFNγ appear to be important for the pathogenesis of the chronic skin inflammation in this model (48), and the principle has been confirmed later in other immunodeficient recipients, i.e. *Rag-2*-deficient mice (50). This phenotype responded to immunosuppressive therapies and was suppressed by CD4⁺/CD45RB^{lo} T cells, indicating that psoriasis-like skin lesions can, in principle, be evoked through T cell dysregulations in the absence of primary cutaneous abnormalities. However, as detailed below, the penetrance of the chronic inflammatory phenotype is variable in these models and, although T cells are involved, the models have not led to the identification of a putative antigen.

Pathogenic role of innate immunity?

A role of the innate immune system in the pathogenesis of psoriasis is suggested not only by several lines of experimental evidence (54) but also by the clinical efficacy of compounds inhibiting tumor necrosis factor (TNF)α, a primary cytokine linking functions of innate and adaptive immunity (55). Indeed, some animal models appear to corroborate the notion that components of the innate immune system contribute prominently to the pathogenesis of psoriasisform lesions: the skin lesions of CD18 hypomorphic mice (*Itgb2^{tm1Bay}/Itgb2^{tm1Bay}* on the PL/J strain) contain large numbers of TNF-producing macrophages (39). Neutralising TNF or macrophage depletion in these mice resulted in alleviation of the inflammatory skin disorder

(56). T cells, neutrophils, mast cells or Langerhans cells were not affected by macrophage depletion. Given that the psoriasis-like phenotype was only obtained after combined injection of monocyte chemoattractant protein 1 (MCP-1) and TNF α into unaffected skin, but not by either mediator alone, both cutaneous recruitment and subsequent activation of macrophages into the skin appeared to be necessary for disease development (56). Similar results were obtained when the role of macrophages was assessed in mice with epidermis-specific deletion of IKK2 (43,52), suggesting that interference with macrophage functions may be a common denominator in some mouse models of chronic hyperproliferative skin disorders, independent of their way of generation. It has been concluded that dermal macrophages, once activated through various triggers, produce large amounts of TNF, ultimately sustaining a vicious circle resulting in psoriasis-like transformation of the skin. Again, these observations have been extrapolated to human psoriasis (57).

In flaky skin mice, a phenotype that may develop in the absence of a functional adaptive immune system (20), a pathogenic role of neutrophils for the generation of the skin symptoms has been suggested based on antibody-mediated depletion (58). In contrast, neutrophil depletion in CD18 hypomorphic mice (*Itgb2^{tm1Bay}/Itgb2^{tm1Bay}* on the PL/J strain) did not alleviate inflammation (56). It appears, therefore, that at least some aspects regarding the role of innate immune mechanisms are discrepant between certain animal models, and the transferability of the respective mechanisms to the human disease awaits further investigations.

Uniform cutaneous reaction patterns in animal models of psoriasis

All of the above mentioned approaches represent animal models with more or less psoriasis-like cutaneous features whose phenotypes and putative pathogeneses have been reviewed extensively (8,9). However, considering the diversity of targeted structures primarily affected in these spontaneously mutated, genetically engineered or immunologically manipulated rodents, which have been published as models for psoriasis, it appears that very similar hyperproliferative inflammatory phenotypes can be obtained through many diverse approaches (examples depicted in Fig. 2). For example, many genetically engineered animals with targeted mutations within the epidermis, the vasculature or immune cells exhibit chronic hyperproliferative inflammatory skin changes. Similar phenotypes may occur in spontaneous mutations or immunological transfer models, although some aspects may vary (such as dermal fibrosis in TGF β 1-transgenic mice or lacking induction of MHC class II in flaky skin mice, Fig. 2). Thus, the mere generation of such models may not suffice to identify cells or

mediators that are crucial initiators of psoriatic skin lesions in humans, if that is possible at all.

The complementary approach, i.e. specific elimination (targeting) of candidate cells or mediators in animal models to identify crucial elements of the pathogenic cascade may also not be sufficient to conclusively identify initiators of the pathogenesis of psoriasis, as exemplified by CD18 hypomorphic mice (*Itgb2^{tm1Bay}/Itgb2^{tm1Bay}* on the PL/J strain) where either depletion of T cells (59) or macrophages (56) resulted in alleviation of the psoriasiform phenotype.

To some degree, it appears that perturbations of the cutaneous homeostasis at (almost) any point may result in a chronic inflammatory phenotype that may – sales-promotional to some extent – be called psoriasis-like or psoriasiform (some examples are schematically depicted in Fig. 3). As evidenced by several examples, genes whose expression is targeted to the basal layers of the epidermis appear to be particularly prone to lead to an inflammatory hyperproliferative phenotype. Thus, animal models have not yet answered the central question in psoriasis research, namely whether psoriasis is based on an intrinsic defect

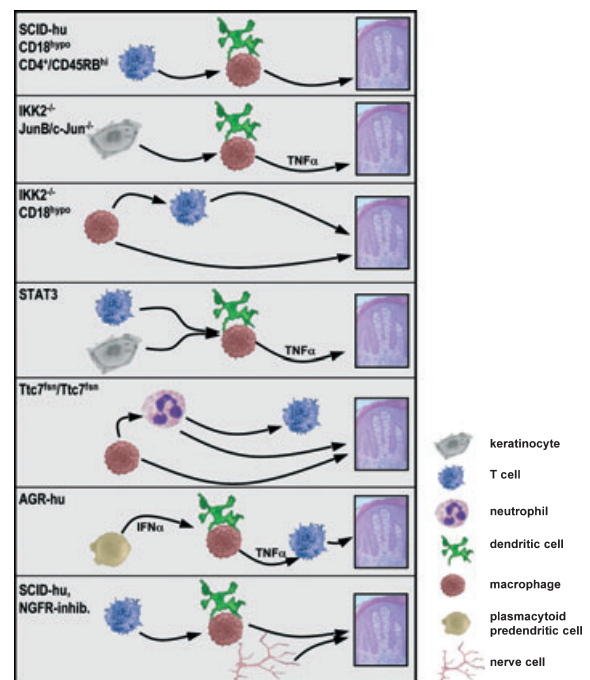


Figure 3. Synopsis of selected pathogenic concepts of psoriasis supported by animal models. As detailed in the text, rather diverse manipulations of the cutaneous homeostasis may result in chronic inflammatory hyperproliferative skin lesions. Depending on the model used – some examples are listed in the panels – tampering with different cell types or molecular functions trigger the development of hyperproliferative inflammatory skin changes, suggesting different pathogenic mechanisms.

within the skin or is a primary autoimmunological disorder. Further along this line, if psoriasis is indeed a disorder with an immunological basis (our currently favoured notion), then there are some experimental systems highlighting the primary importance of the adaptive immune system (i.e. T-cell based), while others point to a crucial role of components of the innate immunity (i.e. some macrophage populations); again, there is evidence for both (and both are not mutually exclusive, of course). However, having stated these – admittedly – somewhat provocative hypotheses, it has also to be said that although animal models have failed thus far to provide the ultimate clue to the pathogenesis of psoriasis, much of our current notion of psoriasis actually stems from those models, and animal models have driven several lines of research in the human disease. Even if no animal model resembles the complete pathogenic network of human psoriasis, we have gained insight into the relevance of many cell types and/or molecules in the pathophysiology of chronic skin inflammation, aspects that presumably are relevant in the setting of psoriasis.

Overall, animal models have undoubtedly shed light on important functions of cells and mediators contributing to chronic cutaneous inflammation, but the uniform psoriasisiform reaction pattern in many of these animals, regardless of the underlying molecular manipulation, appears to limit their suitability as true models of human psoriasis.

Why is it so difficult to interpret animal models of psoriasis?

Homogeneity of experimental animals versus heterogeneity of human patients

First, targeted mutations of single genes in mammals (mice in most cases) are usually introduced in animals of a homogeneous genetic background. In addition, most of these animals are kept under controlled conditions (e.g. specific pathogen-free conditions). These features alone are quite different from the situation in humans, where the clinical spectrum of psoriasis is rather broad (60,61) and where the disease is thought to result from interactions of several genetic susceptibility loci (6). Animal models only begin to reflect these aspects (39).

Imponderable influence of modifying genes

That modifying genes may influence the accrual of a psoriasisiform phenotype is exemplified by several models, including spontaneous mutations [e.g. the flaky skin mutation that results in phenotypes of different severity depending on the genetic background; (20)] and genetically engineered animals [e.g. the CD18 hypomorphic mice (*Itgb2^{tm1Bay}/Itgb2^{tm1Bay}*), which show the psoriasis-like phenotype only when crossed into the PL/J strain; (39)].

In many cases, the potential influence of modifying genes in animal models is not sufficiently appreciated and needs further investigations.

Phenotype penetrance

The expression levels of the transgenes may greatly and unpredictably influence the phenotype in some animal models, as exemplified by IL-1 α transgenic mice, which showed a more severe phenotype in animals with higher expression levels of the transgene (25) or BMP-6-transgenic animals, which showed a phenotype unlike psoriasis (i.e. epidermal atrophy) in a strain with high expression of the transgene (30). It is, therefore, conceivable that genetically engineered animals described as having no or weak hyperproliferative inflammatory phenotypes, such as IL-6 (23) or KGF (24) transgenic mice, could develop inflammatory changes if the transgenes were expressed at different levels (i.e. exceeding a quantitative threshold to initiate the inflammatory cascade), if the genetic background was altered, or if environmental factors (trigger factors) were changed.

Influence of environmental factors

That environmental factors influence the accrual and penetrance of psoriasis-like phenotypes is exemplified by a model based on adoptive transfer of minor histocompatibility mismatched CD4⁺/CD45RB^{hi} T cells into *Prkdc^{scid}* mice, where penetrance and severity of the psoriasisiform phenotype differed between different animal facilities and could be modulated by food and bedding of the animals (42,48,49).

Inherent differences between human and mouse skin

There are several considerable anatomical and physiological differences between human and mouse skin, which may profoundly impact on the phenotype of any disorder claimed to resemble psoriasis. Such inherent differences include, to name but a few, the thickness and architecture of the epidermis (2–3 layers in mice as compared with >6 in humans), the presence or absence of epidermal rete ridges, the density and length of hair follicles, the antigen expression pattern in follicular versus interfollicular epidermis, the constitutive presence of certain types of immune cells (e.g. dendritic epidermal T cells or CD8⁺ dendritic cells), the epidermal turnover time, the thickness of the dermis or the presence of a panniculus carnosus in mice, but not in humans [reviewed in (8)].

'Psoriasis sells'

Sometimes, the prominent publication of a phenotype in laboratory animals may be facilitated if there is a connection to a specific human disease. It is, therefore, possible

that there is an inherent bias in the research community inasmuch as there may be a tendency to (over)-emphasise the similarity to psoriasis when chronic hyperproliferative inflammatory phenotypes are published. Indeed, without going into too much detail here, a glance at the last 25 primary publications of animal models (without the use of human material) exhibiting such a cutaneous phenotype shows that, papers stating the similarity to psoriasis upfront in the title have an average impact factor that is almost twice as high as compared with that of publications without the 'buzzword' psoriasis in the title. The phenotypes described in the respective papers, however, were sometimes not dramatically different. The reasons for that apparent tendency may be manifold, and I will not speculate about them here. However, when interpreting chronic inflammation in animals, one may be well advised to check carefully which criteria may apply to study aspects of psoriasis and which ones do not.

Studying psoriasis in xenotransplantations of human skin

Given that animal models of psoriasis bear the abovementioned limitations, the currently best approach to study psoriasis in animals may be humanisation of mice, although the use of transplanted human skin may not be an animal model in the strict sense of the word. Humanisation of animals to study psoriasis comprises xenotransplantation of human skin and/or immune cells into immunodeficient animals (schematically depicted in Fig. 4) (62).

Nude mice lack functional T cells because of a mutated forkhead box transcription factor N1, resulting in defective thymus development (63). The humoral immune response is also impaired, possibly because of the absence of T helper cells. In contrast, the *Prkdc^{scid}* mutation causes a defect in the antigen receptor gene rearrangement of lymphocytes, resulting in a severe combined immune deficiency (SCID) of the T- and the B-cell system, that is in part reversible (leaky) (64,65). A similar phenotype without the handicap of potential leakiness results from deletion of *Rag-1* or *Rag-2* involved in the development of T and B cells (66,67).

Psoriatic skin retains its phenotype for more than 2 months following transplantation onto nude mice (68). Given that non-lesional skin grafted onto nude mice also adopted features of psoriatic skin (69), *Prkdc^{scid}*, *Rag*-deficient or AGR mice (AGR129 mice are mice deficient in type I (A) and type II (G) IFN receptors in addition to being *Rag-2^{-/-}*; either on a 129Sv/Ev or C57BL/6 background) were later used primarily as recipients in numerous studies [reviewed in (70)]. Depending on the purpose of the study, full-thickness or split-skin can be

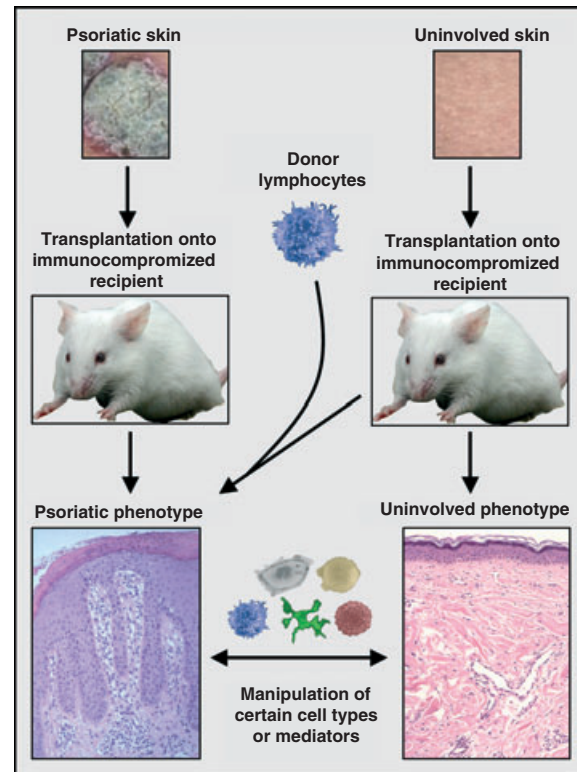


Figure 4. Xenotransplantation of human skin onto immunocompromised recipient mice. Grafts from lesional or non-lesional skin of psoriatic patients preserve their respective phenotypes following transplantation onto immunocompromised mice (e.g. *Prkdc^{scid}* mice, nude mice, AGR mice). Non-lesional skin, however, can be triggered to develop psoriatic features, e.g. by subsets of immune cells (lymphocytes, macrophages, dendritic cells). In addition, the functions of various cell types and/or molecules can be studied in xenotransplantation models.

transplanted from clinically affected or non-affected human skin (71,72). The functional role of immune cells, T cells from the same individual in particular, has been extensively studied in such *Prkdc^{scid}*-hu transplantation models (73,74). Interestingly, this protocol triggered the development of psoriasis in the formerly unaffected skin from psoriasis patients, but not healthy donors (Fig. 4). In addition, the *Prkdc^{scid}*-hu model has helped to understand the role of bacterial superantigens as potential triggers for psoriasis (73). Molecules involved in epidermal recruitment of T cells, such as the $\alpha_1\beta_1$ integrin (75), and the contribution of resident T cells to the pathogenic cascade (76) have also been elucidated recently using xenotransplantation models.

In addition to components of the adaptive immune system, several publications using xenotransplantation of human skin onto mice substantiate a role of macrophages or other cells of the innate immune system for psoriasis-like skin alterations in mice. Infiltration of macrophages, especially in the vicinity of the dermo-epidermal junction,

is a typical feature of psoriasis (77). To interfere with innate immunity, reagents are available to deplete natural killer (NK) cells, neutrophilic granulocytes or macrophages. Alternatively, mice bearing the *Prkdc^{scid}* or the *Rag* mutations can be crossed into other strains affecting the innate immune system: non-obese diabetic (NOD) mice exhibit altered immune functions, including impairment of NK and antigen-presenting cells as well as the complement system (78). The beige mutation (*bg^f/bg^f*) affects NK cell function and granulocyte chemotaxis (79). Tolerance to xenografts is further enhanced in mice with several immune-relevant defects, such as NOD/*Prkdc^{scid}*, mice with additional targeted mutation of $\beta 2$ microglobulin (NOD/*LtSz-scid/scid B2m^{null}*) (80) or a deletion of the common cytokine receptor γ -chain (81,82). Finally, there are so-called BXN mice bearing the beige (*bg^f/bg^f*), nude (*Foxn^{nu}/Foxn^{nu}*) and X-linked immunodeficiency mutation (*Btk^{xid}/Btk^{xid}*) (83).

For drug discovery and development, appropriate animal models should sufficiently mirror human psoriasis, be predictive, inexpensive, reproducible, easy to handle and should allow high throughput experiments. Unfortunately, none of the currently available models fulfils all of these criteria, necessitating the use of different models depending on the specific purpose. Humanised mouse models are important for studying targets, which are only expressed in human tissues, and for testing antibodies or biologics directed against human antigens (84).

What can we do and expect in the future?

Given that phenotypic characterisations are not truly comparable between the various animal models of psoriasis published thus far, it may appear questionable whether synopsis tables depicted in review papers (8,9) accurately reflect strengths and weaknesses of such models with respect to psoriasis features rather than summing up aspects arbitrarily analysed in the respective primary papers. In addition, penetrance and magnitude of phenotypes may critically depend on expression levels of transgenes (and/or copy numbers), environmental factors etc., as outlined above. Therefore, it is not at all precluded that (slight) modifications from the published conditions may impact significantly on the resulting phenotypes. Along this line, the influence of strain-dependent factors, i.e. modifying genes, has not been sufficiently explored yet. Thus, it may be worthwhile to compare animal models under consistent and uniform conditions, which need to be defined in a generally accepted fashion.

So, what can we hope for, short of truly full-fledged psoriasis in laboratory animals? First, I believe that we need to explore if and how different pathogenic aspects are interconnected in given models. This type of research has been

tentatively initiated in some animal models, such as the CD18 hypomorphic mouse (*Itgb2^{tm1Bay}/Itgb2^{tm1Bay}* on the PL/J strain) in which elements of both adaptive (59) and innate immunity (56) have been investigated in the same model system and should now be brought together, possibly including additional aspects such as epidermal or vascular functions. Likewise, xenotransplantation models have been utilised to assess the role of resident cutaneous cells (85) as well as different immigrating cells and inflammatory mediators, indicating the individual relevance of a number of immune cell types (73–76), but failing thus far to sketch the ‘big picture’ of psoriasis. Expanding this line of research, it might be possible to dissect the relative contributions of parts of the immune system to the whole pathogenic mosaic. Hopefully, continuation of such studies and merging of individual aspects will ultimately pave the road to a more complete and unifying picture of the molecular and cellular network in the skin. Furthermore, I think that we need a more reliable means of comparison, preferably in a standardised manner, between human psoriasis and animal models on the one hand, and among the different animal models themselves on the contrary. Towards this end, we need to establish comparability criteria among different models as well as between animal models and human psoriasis. The first scoring system has recently been suggested (86) and now needs to be further developed and elaborated by the community of researchers in the field. Such an algorithm is predicted to be helpful to evaluate clinical, histopathological, immunological, biochemical, therapeutic and other parameters of the various models and to make them directly comparable.

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