Proteasome inhibition reduces superantigen-mediated T cell activation and the severity of psoriasis in a SCID-hu model

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There is increasing evidence that bacterial superantigens contribute to inflammation and T cell responses in psoriasis. Psoriatic inflammation entails a complex series of inductive and effector processes that require the regulated expression of various proinflammatory genes, many of which require NF-κB for maximal trans-activation. PS-519 is a potent and selective proteasome inhibitor based upon the naturally occurring compound lactacystin, which inhibits NF-κB activation by blocking the degradation of its inhibitory protein IκB. We report that proteasome inhibition by PS-519 reduces superantigen-mediated T cell–activation in vitro and in vivo. Proliferation was inhibited along with the expression of very early (CD69), early (CD25), and late T cell (HLA-DR) activation molecules. Moreover, expression of E-selectin ligands relevant to dermal T cell homing was reduced, as was E-selectin binding in vitro. Finally, PS-519 proved to be therapeutically effective in a SCID-hu xenogeneic psoriasis transplantation model. We conclude that inhibition of the proteasome, e.g., by PS-519, is a promising means to treat T cell–mediated disorders such as psoriasis.


Introduction

Bacterial superantigens are characterized by their ability to interact with and activate T cells that share defined T cell receptor Vβ segments. Therefore, up to 20% of T cell receptor α/β+ T cells are stimulated by any given superantigen, and this proportion is several orders of magnitude higher than that of activation by conventional antigens (1). Superantigen-mediated T cell activation results in increased proliferation along with secretion of proinflammatory cytokines and upregulated expression of activation markers such as adhesion molecules, including the skin-homing receptor cutaneous lymphocyte-associated antigen (CLA) (2–4). There is increasing evidence that superantigens are involved in the pathogenesis of several autoimmune diseases, e.g., rheumatoid arthritis and diabetes mellitus (5, 6). Another T cell–mediated autoimmune disease is psoriasis. Clinically, there is an association of this disease with bacterial infections, as colonization and infection with Staphylococcus and Streptococcus have been reported to exacerbate psoriasis (7, 8). In this regard, Staphylococcus aureus has been found on the skin of more than half of the patients with chronic plaque psoriasis, and more than half of the clinically isolated strains produce superantigens (8, 9). Exacerbation of chronic plaque psoriasis by superantigens has been observed (9), although the most convincing clinical association between bacterial infection and psoriasis is in patients with acute guttate psoriasis. The observation that streptococcal M proteins and keratins share common epitopes led to the hypothesis of psoriasis being triggered by superantigen-activated T cells cross-reacting with keratins (10, 11).

Induction of psoriasis has been demonstrated experimentally in the SCID-hu xenogeneic transplantation model by injecting bacterial superantigens into nonlesional psoriatic skin transplanted onto mice lacking functional B and T cells (12). This phenomenon was found to be T cell–dependent (13, 14). Lesional T cells are characterized by the expression of numerous activation markers such as the very-early-appearing CD69 (15), early IL-2 receptor α-chain CD25, the later-appearing HLA-DR (16), and adhesion molecules such as CLA and CD15s (17–19). Regulation of these molecules involves NF-κB (20).

The major intracellular pathway for protein degradation is the ubiquitin-proteasome pathway (UPP) (21, 22). The proteasome is a large multimeric protease present in all eukaryotic cells that exhibits a highly conserved so-called 20S core structure (23). Proteasomes are responsible for the degradation of protein substrates after they have been “tagged” by a poly-ubiquitin chain (24). The inflammatory events in psoriasis are composed of a complex series of inductive and effector...
purified T cells was performed using magnetic cell separation according to standard procedures using the Pan T cell Isolation Kit (Miltenyi Biotec GmbH, Bergisch Gladbach, Germany). Selected cells were ≥ 97% CD3+. PBMCs or highly purified T cells were cultured in AIM-V medium (Life Technologies Inc., Grand Island, New York, USA) supplemented with 5% FCS (Life Technologies Inc.). Phytohemagglutinin (Biochrom AG, Berlin, Germany) (1 µg/ml), N-acetylcysteine (25 mM), and toxic shock syndrome toxin-1 (TSST-1) (100 ng/ml) were obtained from Sigma-Aldrich Co., and TGFB-β (50 ng/ml) from R&D Systems Inc. (Minneapolis, Minnesota, USA). IL-2 (100 U/ml) was a kind gift of Chiron Behring GmbH and Co. (Marburg, Germany). PS-519 was produced by Millennium Pharmaceuticals Inc. (28).

Monoclonal antibodies were purchased from BD Biosciences (Heidelberg, Germany) (CD3, HLA-DR, CLA, CD11s) or Immunotech GmbH (Hamburg, Germany) (CD25, CD69). Recombinant human E- and P-selectin IgG fusion proteins were obtained from R&D Systems Inc. and BD Biosciences, respectively.

Electrophoretic mobility shift assay. Total cell extracts from 5 × 10^6 to 1 × 10^7 PBMCs or T cells were prepared by resuspending PBS-washed cell pellets with a buffer containing the detergent Igepal CA-630 (Sigma-Aldrich Co.), as described previously (36). Protein concentrations of supernatants were determined by Bio-Rad protein assay (Bio-Rad Laboratories, Hercules, California, USA). DNA-binding conditions for NF-κB have been described in detail previously (37). Briefly, 10 µg of protein was used for binding reaction, which contained 10,000 cpm of 32P-labeled, double-stranded oligonucleotide with a high-affinity NF-κB–binding motif from the κ light chain enhancer (Promega Corp., Madison, Wisconsin, USA). DNA-binding reactions were analyzed by electrophoresis on native 4% polyacrylamide gels. Dried gels were exposed to Kodak XR films (Eastman Kodak Co., Rochester, New York, USA). The radioactivity in the NF-κB DNA complexes was quantitated by scanning and NIH Image analysis (Scion Image for Windows; Scion Corp., Frederick, Maryland, USA).

Proliferation analysis. PBMCs (2 × 10^6/ml) were stimulated with TSST-1 (100 ng/ml) in 96-well flat-bottom microplates in AIM-V medium supplemented with 5% FCS for 4 days in the presence or absence of PS-519 (0–5 µg/ml) or vehicle (propylene glycol/0.9% saline [1:1]). At the culture initiation, PS-519 was added 1 hour prior to TSST-1 and supplemented daily. After 4 days at 37°C in 5% CO₂/air, cells were labeled for 18 hours with 0.5 µCi [3H]-thymidine (NEN Life Science Products Inc., Boston, Massachusetts, USA), and proliferative response was determined using a liquid scintillation counter. Control experiments were performed with vehicle only (propylene glycol/0.9% saline [1:1]) and showed no change in [3H]-thymidine uptake.

Phenotype analysis. Freshly isolated PBMCs (1 × 10^6/ml) were stimulated for 1–7 days with TSST-1 (100 ng/ml)
in the presence or absence of PS-519 (0.25–2.5 µg/ml). PS-519 was added 1 hour before culture initiation and supplemented daily. Control experiments were performed with vehicle only (propylene glycol/0.9% saline [1:1]) and showed no change in the expression of the analyzed surface markers. Staining was performed at the indicated time points by standard direct or indirect immunofluorescence labeling techniques with goat-anti-rat IgM-F(ab)_2-phycocyanin. In each sample, irrelevant mAb’s of the appropriate isotype were used as controls. Fluorocytometer analysis was performed on a FACSCalibur (BD Biosciences) using standard procedures with CellQuest research software (BD Biosciences). Lymphocytes were detected in a gate set on lymphocyte-sized cells.

**In vitro E-selectin and P-selectin binding assay.** Recombinant human E- and P-selectin IgG fusion proteins were obtained from R&D Systems Inc. and from PharMingen (Hamburg, Germany), respectively. We incubated 5 × 10^6 T cells for 30 minutes with 1 µg fusion protein in Ca^++/Mg^++-containing PBS at room temperature. Thereafter, cells were washed, labeled with anti-human IgG-phycocyanin (Dianova GmbH, Hamburg, Germany) for 15 minutes, and analyzed on a FACSCalibur using the CellQuest software (BD Biosciences) as outlined above. Staining in the absence of fusion proteins and in the presence of anti-CD3–FITC together with secondary anti-human IgG-phycocyanin demonstrated absence of unspecific binding reactivity.

**Skin donors.** This study was approved by the ethics committee of the faculty of medicine of the Johann Wolfgang Goethe-University Frankfurt. Written informed consent was obtained from patients with chronic plaque-stage psoriasis, and lesional skin was excised from the upper arm in local anesthesia.

**Transplantation procedure.** The animal experiments were approved by the Regierungspraesidium Darmstadt. Transplantations were done as described previously (38, 39). Human full-thickness xenografts were transplanted onto the backs of 6- to 8-week-old C.B17 SCID mice (Charles River Deutschland GmbH, Sulzfeld, Germany). For the surgical procedure, mice were anesthetized by intraperitoneal injection of 100 mg/kg ketamine and 5 mg/kg xylazine. Spindle-shaped pieces of full-thickness skin measuring 1 cm in diameter were grafted onto corresponding excisional full-thickness defects of the shaved central dorsum of the mice and fixed by 6-0 atrumatic monofilament sutures. After applying a sterile Vaseline-impregnated gauze, the grafts were protected from injury by suturing a skin pouch over the transplanted area using the adjacent lateral skin. The sutures and over-tied pouches were left in place until they resolved spontaneously after 2–3 weeks.

**Treatment protocol.** Grafts were allowed 2 weeks for acceptance and healing. Thereafter, daily intraperitoneal injections were performed between days 15 and 42 after transplantation. Mice received either vehicle (propylene glycol/0.9% saline [1:1]), dexamethasone (0.2 mg/kg body weight), or PS-519 (1 mg/kg body weight) in a final volume of 200 µl. The grafts of each of the donors underwent an identical treatment protocol. The data on histological analysis of the skin sections are based on findings of 12 different grafts from four psoriasis patients (four control, four PS-519–treated, and four dexamethasone-treated animals), whereas data on 20S proteasome activity are based on 16 different grafts (four control, four dexamethasone-treated, and eight PS-519–treated animals).

**Histological analysis of the grafts.** Mice were sacrificed at day 42, and after excision with surrounding mouse skin the grafts were formalin-embedded. Subsequently, routine hematoxylin-and-eosin stainings were performed, and the grafts were analyzed with regard to their pathological changes both qualitatively (epidermal differentiation, inflammatory infiltrate) and quantitatively (epidermal thickness) by a blinded investigator as described previously (38, 39). Briefly, maximal epidermal thickness was measured from the tip of the rete ridges to the border of the viable epidermis. The values were determined using an ocular micrometer, taking the mean of ten consecutively measured rete ridges. The number of infiltrating leukocytes was evaluated by counting stained leukocytes in five adjacent high-power fields. For immunohistochemical analysis, skin samples were immediately snap-frozen in liquid nitrogen and stainings were performed as described previously (38).

**Assay of 20S proteasome activity.** Inhibition of 20S proteasome activity was measured in blood samples. Blood (500 µl) was collected in sodium heparin-containing tubes and centrifuged at 2200 g for 10 minutes at 4°C. Plasma was discarded and the pellet washed with cold 1× PBS. The blood cells were lysed with 5 mM EDTA (pH 8.0) for 1 hour and then centrifuged at 6600 g for 10 minutes at 4°C. The whole-blood lysate samples were then used in the 20S proteasome assay as previously described (40). Briefly, samples (10 µl) were added to 2 ml of substrate buffer (20 mM HEPES, 0.5 mM EDTA, 0.05% SDS, and 60 mM Ys substrate Suc-Leu-Leu-Val-Tyr-AMC [Bachem Biochemica GmbH, Heidelberg, Germany]). The reaction was carried out at 37°C for 5 minutes, and the rate of substrate cleavage per 20S proteasome activity was determined. The protein content of the samples was determined using a Coomassie protein assay (Pierce Chemical Co., Rockford, Illinois, USA).

**Statistical analysis.** Data are presented as mean ± SD. All data satisfied the normality test and were analyzed by either one-way ANOVA with Dunnett’s multiple comparison post-test or Student t test using InStat version 3.00 for Windows 95 (GraphPad Software for Science Inc., San Diego, California, USA). P values less than 0.05 were considered statistically significant.

**Results.**

*Cell activation–induced NF-κB DNA binding in human T cells is suppressed by the proteasome inhibitor PS-519. Since activated T cells are involved in maintaining psoriatic...*
skin lesions and many T cell activating signals are transduced via activation of the nuclear transcription factor κB, we assessed the effects of PS-519 on superantigen- and cytokine-stimulated cells. PBMCs were stimulated with TSST-1 (100 ng/ml), and highly purified human T cells were stimulated with PHA (1 µg/ml), IL-2 (100 U/ml), and TGF-β (50 ng/ml). Total cell extracts were isolated 4 hours after culture initiation as outlined in Methods and analyzed for NF-κB–binding activity using a 32P-labeled oligonucleotide probe. As we reported earlier, we found, in a time-dependent manner, the appearance of a protein-DNA complex, which was first observable after 30 minutes and reached a maximum after 4 hours (41). The PHA/IL-2/TGF-β-induced protein-DNA complex was characterized by using subunit-specific antibodies against p50 and p65 subunits of NF-κB. These subunits form prototypic heterodimers that are most frequently observed upon stimulation of various cell types with diverse conditions. Almost all of the protein-DNA complex newly induced by PHA/IL-2/TGF-β treatment was supershifted upon the addition of specific antibodies against the p50 and p65 subunits of NF-κB to the DNA-binding reaction. A nonspecific antibody was ineffective (41). This suggests that PHA/IL-2/TGF-β treatment induced complexes predominantly containing p50 and p65 subunits.

The appearance of the NF-κB DNA-binding complex was reduced in a dose-dependent manner using PS-519. T cells or PBMCs were pretreated for 1 hour with PS-519 (1–10 µg/ml). Thereafter, cells were stimulated with PHA/IL-2/TGF-β or TSST-1 for 4 hours. As expected, PS-519 suppressed NF-κB DNA-binding activity in stimulated T cells and PBMCs (Figure 1).

**PS-519 inhibits TSST-1–induced T cell proliferation.** Superantigens such as TSST-1 belong to the strongest T cell–activating compounds because they stimulate a large proportion of T cells, as compared with nominal antigens, via their interaction with the T cell receptor Vβ-chain. Additionally, they have been shown to be involved in the induction of psoriasis in animal models of the disease and in human studies (12, 13, 42, 43). Therefore, we analyzed the effects of PS-519 on T cell proliferation and activation in superantigen-stimulated cells. TSST-1 induced a 28-fold increase in 3H-thymidine incorporation, which was inhibited in a dose-dependent manner by PS-519 and reached a maximum at 2.5 µg/ml. Stimulation indexes were: unstimulated cells, 1; TSST-1–stimulated cells, 28.13 ± 3.30; cells stimulated with TSST-1 and PS-519 (2.5 µg/ml), 1.65 ± 1.81 (Figure 2). Proliferation of unstimulated cells cultured in the presence of PS-519 (0.25–2.5 µg/ml) remained unchanged (Figure 2). Using trypan blue exclusion tests and propidium iodide uptake, we excluded toxic effects of PS-519 as a possible cause for reduced T cell proliferation (data not shown).

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**Figure 1**
The PHA/IL-2/TGF-β–induced NF-κB DNA complex is suppressed by the proteasome inhibitor PS-519. Human T cells were stimulated with PHA/IL-2/TGF-β for 4 hours. PS-519 (1–10 µg/ml) suppressed NF-κB DNA-binding activity. The NF-κB DNA complex is indicated by a filled arrowhead. The open arrowhead shows the position of the unbound DNA probe. Binding mixture without cell extract was applied on lanes 1 and 10. The antioxidant N-acetylcysteine (NAC; 25 mM), which is known to inhibit IkB kinase, served as control (lane 4). The results are representative of three electrophoretic mobility shift assays from three independent donors.

**Figure 2**
PS-519 inhibits TSST-1–induced T cell proliferation. PBMCs (2 × 10⁶/ml) obtained from five healthy volunteers were stimulated with TSST-1 (100 ng/ml) in the absence or presence of PS-519 (0.25–2.5 µg/ml) for 4 days and thereafter pulsed with ³H-thymidine. Incorporation of ³H-thymidine into DNA was calculated using a liquid scintillation counter. Stimulation index was calculated by the ratio: decays per minute of experimental group/decays per minute of control group. Open symbols represent resting PBMCs, filled symbols TSST-1–stimulated PBMCs. A significant reduction in proliferation was observed starting at 0.25 µg/ml PS-519 (P < 0.001). Values represent mean ± SD of five healthy donors.
PS-519 inhibits TSST-1–induced expression of T cell activation molecules. Since superantigens are potent activators of T cells and activated T cells are involved in maintaining the psoriatic skin lesions, we investigated the effects of the proteasome inhibitor PS-519 on superantigen-mediated T cell activation. For this purpose we chose the very early activation marker CD69, the early activation marker CD25, and the late activation marker HLA-DR, which are all under the transcriptional control of NF-κB (15, 25, 44, 45). In TSST-1–stimulated cells, CD69 expression increased about fourfold at day 1 (unstimulated cells 9.62% ± 5.24% versus TSST-1–stimulated cells 43.89% ± 10.14%) and remained unchanged for 5 days. PS-519 significantly and dose-dependently reduced CD69 expression by T cells up to 40% (Figure 3a). CD25 expression by T cells continuously increased, reaching a plateau at day 5 of TSST-1 stimulation (day 5: unstimulated cells 8.32% ± 2.29% versus TSST-1–stimulated cells 53.34% ± 2.99%) and remained stable until day 9 (data not shown). Again, PS-519 induced a dose-dependent inhibition of this activation marker with a complete suppression of CD25 expression at 2.5 µg/ml PS-519 (P < 0.001). For HLA-DR expression, significant reduction was observed on day 1 at 2.5 µg/ml PS-519 (P < 0.001), on day 3 starting at 1.0 µg/ml PS-519 (P < 0.001), and on day 5 starting at 0.25 µg/ml PS-519 (P < 0.001). Data represent means of five experiments ± SD.

Figure 3
PS-519 inhibits TSST-1–induced expression of T cell adhesion molecules. PBMCs were stimulated with TSST-1 (100 ng/ml) in the presence or absence of PS-519 (0.25–2.5 µg/ml). CLA+CD3+ (a), CD15s+CD3+ (b), and HLA-DR+CD3+ (c) surface expression and binding of CD3+ cells to E-selectin (c) was measured at days 1, 3, 5, 7, and 9 (days 7 and 9 not shown) by flow cytometry. Appropriate isotype Ig’s served as controls to set gates for positive and negative staining. For CLA expression, significant reduction was observed on day 3 at 2.5 µg/ml PS-519 (P < 0.05), and on days 5 and 7 starting at 0.25 µg/ml PS-519 (P < 0.001). For CD15s expression, significant reduction was observed on days 5 and 7 starting at 0.25 µg/ml PS-519 (P < 0.001). For E-selectin binding, significant reduction was observed on day 5 starting at 0.25 µg/ml PS-519 (P < 0.001), and on day 7 starting at 0.5 µg/ml PS-519 (P < 0.05). Data represent means of five independent experiments ± SD.

Figure 4
PS-519 inhibits TSST-1–induced expression of T cell activation molecules. PBMCs were stimulated with TSST-1 (100 ng/ml) in the presence or absence of PS-519 (0.25–2.5 µg/ml). CD69+CD3+ (a), CD25+CD3+ (b), and HLA-DR+CD3+ (c) surface expression was measured at days 1, 3, 5, 7, and 9 (days 7 and 9 not shown) by flow cytometry. Appropriate isotype Ig’s served as controls to set gates for positive and negative staining. For CD69 expression, significant reduction was observed on day 1 starting at 1.0 µg/ml (P < 0.05), and on days 3 and 5 starting at 0.5 µg/ml PS-519 (P < 0.001 and P < 0.05, respectively). For CD25 expression, significant reduction was observed on day 3 starting at 1.0 µg/ml (P < 0.001), and on day 5 starting at 0.25 µg/ml PS-519 (P < 0.001). For HLA-DR expression, significant reduction was observed on day 1 at 2.5 µg/ml (P < 0.05), on day 3 starting at 1.0 µg/ml (P < 0.05), and on day 5 starting at 0.25 µg/ml PS-519 (P < 0.001). Data represent means of five experiments ± SD.
CD25 upregulation at 2.5 µg/ml (Figure 3b). HLA-DR expression by TSST-1–stimulated T cells peaked at day 5 (unstimulated cells 21.79% ± 3.41% versus TSST-1–stimulated cells 51.79% ± 12.26%) with a decline at day 9 (data not shown). HLA-DR expression by T cells treated in the presence of PS-519 was significantly reduced even as compared with unstimulated cells (5.81% ± 4.40%; P < 0.01; Figure 3c).

**PS-519 inhibits TSST-1–induced expression of T cell homing molecules.** Recirculation of activated T cells into the skin is of key importance for both immunosurveillance and T cell–mediated inflammatory disorders such as psoriasis. We therefore examined the effects of PS-519 on TSST-1–induced upregulation of molecules known to be expressed by skin-seeking T cells. Expression of CLA and the closely related moiety sialyl LewisX (sLeX = CD15s) is induced by superantigens (2–4, 46). As described, we observed a maximum induction of both antigens at day 5 of TSST-1 stimulation (CLA: control cells 14.57% ± 2.52% versus TSST-1–stimulated cells 35.67% ± 5.63%; CD15s: control cells 15.31% ± 1.99% versus TSST-1–stimulated cells 50.19% ± 2.51%). In PS-519–treated cells, the TSST-1–induced upregulation of both skin-homing molecules was completely inhibited in a dose-dependent manner, and, even in concentrations as low as 0.5 µg/ml, CLA and CD15s expression remained below control levels (CLA: PS-519–treated [2.5 µg/ml] and TSST-1–treated cells 9.61% ± 1.72%; CD15s: 9.61% ± 1.72%; Figure 4, a and b). CLA and CD15s expression correlated with in vitro E-selectin binding, which was also completely inhibited by PS-519 treatment in concentrations above 1 µg/ml (Figure 4c). In vitro binding of T cells to P-selectin remained unchanged in TSST-1– and TSST-1+PS-519–treated cells as compared with control cells (data not shown). CD54 expression was also inhibited by PS-519, but to a lesser extent as compared with suppression of CLA or CD15s (data not shown).

Interestingly, lower concentrations of PS-519 were necessary for inhibition of T cell proliferation, of T cell activation markers, and of cell adhesion molecules as compared with inhibition of NF-κB DNA binding (Figures 1–4). This can be explained by the fact that threshold levels of active NF-κB are required for T cell activation and optimal cellular adhesion molecule expression. Therefore, a small decrease in the level of activated NF-κB can lead to a profound change in the level of transcription (27, 30).

**PS-519 is effective in the treatment of psoriasis in a xenogeneic transplantation model.** Having demonstrated reduced superantigen-mediated T cell activation in vitro, we employed PS-519 as a therapeutic tool in a SCID-hu xenogeneic transplantation model that proved to be a useful tool in the evaluation of antipsoriatic drugs (47, 48). Only grafts from lesional psoriatic skin treated with the vehicle retained their increased epidermal thickness (414 ± 54 µm; Figures 5a and 6) and papillomatosis. The percentage of proliferating basal keratinocytes expressing the proliferation marker Ki-67 was 10% ± 4% (Figures 5b and 6). A dense leukocytic infiltrate was present in the upper dermis (75 ± 12 leukocytes per high power field; Figure 5c). In contrast, grafts treated with PS-519 exhibited a markedly reduced epidermal thickness (110 ± 43 µm) along with fewer Ki-67–positive basal keratinocytes (1% ± 1%). The remaining inflammatory infiltrate localized in the upper dermis was reduced (41 ± 7 leukocytes per high power field; Figure 5c). These changes paralleled the findings in grafts of mice that received dexamethasone treatment (Figure 5, a–c, and Figure 6).

**20S proteasome activity is reduced in PS-519–treated animals.** To determine the pharmacodynamics of PS-519 from these transplanted mice, blood was collected 2 hours after treatment at days 4, 8, 14, and 28 from PS-519– and vehicle-treated animals. Results indicate an 85.7% ± 8.6% (mean ± SD) reduction in proteasome activity in the PS-519–treated animals as compared with the control (Figure 7). Already after 4 days of treatment,
20S proteasome activity was reduced by 81.6% ± 1.7%, indicating that the effects were achieved quite early after the beginning of treatment. In the vehicle-treated mice, 20S proteasome activity remained constant during the whole experimental period (data not shown). No effect on 20S proteasome activity was seen in the dexamethasone-treated group (data not shown). These data support the critical role the proteasome plays to suppress the activation of NF-κB, thereby decreasing the transcription of many genes encoding proinflammatory proteins, including the T cell surface markers.

**Discussion**

Our results demonstrate that the selective proteasome inhibitor PS-519 can significantly inhibit numerous parameters in the process of superantigen-mediated T cell activation in vitro and act as a potent therapeutic agent for psoriasis in a xenogeneic transplantation model.

The relevance of superantigen-mediated T cell activation in the triggering process of psoriasis is documented by findings of numerous research groups and is thought to be greater for patients suffering from acute guttate psoriasis than for those with chronic plaque-stage psoriasis (9, 10, 12, 13, 43). Moreover, activated T cells are involved in maintaining the psoriatic skin lesions (49, 50). This is underscored by the observation that injection of IL-2 and superantigen-stimulated T lymphocytes into SCID-hu mice is sufficient to induce a full-fledged psoriatic plaque (13, 14). Therefore, at least in this model, the contribution of other cell types seems to be of minor relevance. Based on these observations, the capability of any given compound to interfere with superantigen-mediated T cell activation makes it a putative agent to treat at least certain subsets of psoriasis. Indeed, PS-519 interfered with T cell activation. The activation markers analyzed here represent molecules that are expressed during various phases of T cell activation: CD69 appears very early, CD25 early, and HLA-DR late after in vitro activation.

Besides these phenomenological findings, demonstration of proliferation inhibition and even more interference with adhesive interactions may be of direct functional importance. Lymphocyte migration is an essential requirement for efficient surveillance of tissues for infectious pathogens and for the recruitment of effector cells at sites of inflammation. The extravasation of leukocytes at sites of inflammation is a multiple-step process that involves the interaction of selectins with their carbohydrate ligands, thus mediating the initial tethering of blood leukocytes to the vessel wall.

**Figure 6**

PS-519 suppresses hallmarks of psoriasis in a xenogeneic transplantation model. Grafted skin in PS-519-treated mice (1 mg/kg body weight) showed normalization of epidermal architecture, loss of papillomatosis, and marked reduction of acanthosis (c, hematoxylin-and-eosin stain) as compared with vehicle-treated mice (a, hematoxylin-and-eosin stain). In PS-519-treated (d, Ki-67 stain) as compared with vehicle-treated mice (b, Ki-67 stain), proliferation of basal keratinocytes was markedly reduced. Treatment with dexamethasone (0.2 mg/kg body weight; e and f) was as effective as PS-519 treatment. The sections show one representative of four experiments. x200.

**Figure 7**

20S proteasome activity is reduced in PS-519 as compared with vehicle treated mice. Peripheral blood from vehicle- and PS-519-treated mice was drawn 2 hours after the final injection at days 4, 8, 14, and 28. Thereafter, 20S proteasome activity was determined as described above. PS-519–treated animals showed an 85.7% ± 8.6% (mean ± SD) inhibition of the 20S proteasome activity as compared with vehicle-treated mice at day 28 (***P < 0.0001). Already after 4 days, 81.7% ± 1.6% inhibition as compared with controls was achieved (**P < 0.01). The 20S proteasome activity in vehicle-treated mice remained unchanged when measured at 4, 8, and 14 days as compared with day 28 (data not shown). The 20S proteasome activity is given in pmol/s/mg protein. Data represent mean ± SD; n = 6, days 4–14; n = 8, day 28.
sel wall (51, 52). With respect to the initiation and perpetuation of psoriasis, we and others have recently suggested a role for inappropriate T cell skin homing based on the upregulation of CLA (2–4, 12, 53, 54). Of note, bacterial superantigens are among the most potent inducers of CLA upregulation (2–4).

Expression of the various T cell surface markers investigated in this study is under the control of NF-κB (refs. 25, 41, 45; and M. Podda et al., manuscript submitted for publication). This transcription factor can be activated by a large number of signals, including antigens, cytokines, and infectious agents, in particular bacterial superantigens (25, 45, 55, 56). We therefore investigated the effects of PS-519 on T cell activation–induced NF-κB binding to DNA. That PS-519 exhibits its effects via inhibition of NF-κB is suggested by our finding of dose-dependent suppression of DNA binding in electrophoretic mobility shift assays. Additionally, in vivo data from PS-519–treated mice showed significant reduction in the activity of its biochemical target, the 20S proteasome.

The SCID-hu xenogeneic transplantation model employed in our in vivo studies is a well-established tool for the investigation of various aspects of the pathogenesis of psoriasis as well as for the screening of putative antipsoriatic drugs (14, 48, 57, 58). Using this model, we were able to show normalization of grafts from lesional skin in PS-519–treated mice. The effects were at least equivalent to those obtained with dexamethasone, a drug with known antipsoriatic potential. Given the critical role of NF-κB in the immune response and other defensive responses, it is noteworthy that the mice treated with PS-519 did not show signs of infection or wasting. Our observations, therefore, clearly are in line with reports from other groups on the beneficial effects of PS-519 in animal models of T cell–mediated immune disorders (29–31).

The SCID-hu xenogeneic disease model of psoriasis is complex. In this model, several aspects of proteasome inhibition may contribute to the observed beneficial effects: (a) inhibition of T cell activation within the transplanted psoriatic lesion, since suppression of T cell activation was demonstrated in vitro; (b) inhibition of adhesion molecule expression by T cells; and (c) inhibition of other cell types such as neutrophils, macrophages, and keratinocytes, since inhibition of the proteasome by PS-519 is not T cell–specific.

In summary, the selective proteasome inhibitor PS-519 is capable of interfering with superantigen–mediated T cell activation, presumably by blocking NF-κB. Its effectiveness in an animal model for psoriasis suggests its potential as a novel therapeutic regimen in this disease.

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