UV-induced CD39 expression promotes DNA damage and development of cutaneous squamous cell carcinoma

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Introduction

- Cutaneous squamous cell carcinoma (cSCC) results from UV-induced DNA damage and immunosuppression.
- Regulatory T cells (Tregs), which are present in cSCC, can suppress immune responses via CD39 (ENTPD1), which catalyses the generation of adenosine from extracellular ATP.
- The purpose of the current study was to investigate the role of CD39 in UV-induced epidermal DNA damage and cSCC.



Medicine

Methods

- Flow cytometry was conducted on immune cells isolated from cSCC, peripheral blood and normal skin.
- Immunofluorescence microscopy was performed on frozen cSCC sections.
- Immunohistochemistry was performed on formalin-fixed paraffin embedded primary cSCCs which had not metastasised after 5 years and primary cSCCs which did metastasise.
- Wild type and IL27RA knockout mice were irradiated with 100mJ/cm² UVB and *Entpd1* measured by qPCR.
- Normal human keratinocytes were stimulated with supernatants from anti-CD3 activated skin resident T cells primed with/without 100 ng/ml rhlL-27 and irradiated with 10mJ/cm² UVB prior to immunofluorescence staining for yH2AX as a marker of DNA damage.
- Murine ears were excised, irradiated with 100mJ/cm² UVB in culture and epidermal sheets were obtained for immunofluorescence staining for γ H2AX.
- Keratinocytes were cultured with 10 μ M adenosine and microarray was performed for gene expression analysis.

Results

- CD39 expression is increased in human cSCC compared to normal skin (figure 1).
- Increased CD39 expression in primary cSCCs is associated with metastasis (figure 2).
- CD39 is upregulated on T cells in cSCC and is highly expressed by Tregs (figure 3).
- UV-induced CD39 expression in murine skin is IL-27 dependent and IL-27 is present on CD14+, CD207+ (Langerin) and CD209+ (DCSIGN) antigen presenting cells in human cSCC (figure 4). IL-27 signalling suppresses UV-induced DNA damage repair (shown by yH2AX) staining) in human keratinocytes and murine skin (figure 5). yH2AX is expressed in human cSCCs and perilesional skin infiltrated with CD39+ immune cells (figure 6). Adenosine downregulates *NAP1L2*, a nucleosome assembly protein required for DNA repair (figure 7).



(A) Schematic diagram for experiments performed in (B) and (C). (B) CD39 and FOXP3 expression in healthy donor skin T cells after stimulation with rhIL-27 or vehicle. (C) Human keratinocytes were cultured with medium (n=3), medium from anti-CD3 stimulated skin-resident T cells (n=4), or medium from IL-27 primed and anti-CD3 stimulated skin-resident T cells (n=3) followed by UVB irradiation and measurement of DNA damage by immunofluorescence quantification of γ H2AX after 24 hours. (D) Ears from WT, ENTPD1^{-/-}, and IL27Ra^{-/-} mice were irradiated with UVB and treated with or without IL-27. Immunofluorescence quantification of γ H2AX⁺ in epidermal sheets after 24 hours was performed (n = 3 per group).

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(A) Schematic diagram showing the CD39 axis, leading to the conversion of extracellular ATP to ADP, AMP and adenosine. (B) Flow cytometry showing CD39 expression within CD45+ and CD45- cell populations in normal skin (n=11) and cSCC (n=13). (C) Immunofluorescence microscopy of cSCC demonstrating CD39+ immune cells in the stroma. (D) ADP, AMP and adenosine (ADO) concentrations from supernatants from cSCC and normal skin as measured using liquid chromotography – electrospray tandem mass spectrometry.



Figure 7

(A) Heat map of the top 49 differentially expressed genes in normal human keratinocytes stimulated with vehicle or 10 µM adenosine, showing downregulation of NAP1L2. (B) Measurement of DNA damage (based on cells with $\geq 5.53BP1$ foci) 2 and 24 hours after UVB irradiation of keratinocytes with or without siRNA knockdown of *NAP1L2* (n=4 for all conditions).





(A) Representative immunohistochemistry images of cSCC stained for CD39. (B) Percentages of tumoral immune cells that are CD39⁺ in primary cSCCs that did not metastasise (P-NM, n=51) and primary cSCCs that metastasised (P-M, n=54).

Conclusions

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•••• primary metastatic non-metastatic



(A) Representative FACS plots showing expression of FOXP3 and CD39 on CD3+ T cells from blood, non-lesional skin and cSCC. (B, C) CD39+ frequencies in the (B) total CD3+ T cell and (C) CD4+FOXP3+ Treg populations in blood, non-lesional skin and cSCC (n=13). (D) Representative FACS plot showing expression of cutaneous lymphocyte antigen (CLA, a skin homing marker) within the CD39+ and CD39- T cell populations in cSCC. (E) CD45RO and CD39 coexpression in CD4+FOXP3+ Tregs in blood, non-lesional skin and cSCC (n=7). (F) PD-1 and CD39 co-expression in FOXP3+ Tregs and FOXP3- T cells in blood, non-lesional skin and cSCC (n=8).

- CD39 is highly expressed by Tregs in cSCC and CD39 and adenosine are upregulated in cSCC compared to normal skin.
- Higher CD39 expression in primary cSCCs is associated with metastasis.
- UV radiation induces CD39 in an IL-27 dependent manner.
- Adenosine downregulates the expression of NAP1L2, which is important for repair of UV-induced DNA damage.
- These results indicate a key role for CD39 in UV-induced DNA damage and promoting cSCC development and metastasis.



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