

resulted in an increased production of CD4⁺ T cells, in contrast to the transplantation of *Cd83*^{-/-} epithelial cells.

These results imply that the developmental defect that is seen in *Cd83*^{-/-} mice is due to an impaired thymic microenvironment in these mice and that the expression of CD83 by non-lymphoid cells within the thymus is required for CD4⁺ T-cell development. This work is the first to show the importance of CD83 engagement for the regulation of CD4⁺ T-cell differentiation in the thymus.

Jenny Buckland

References and links

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AUTOIMMUNITY

Different cuts

Local variations in proteasome composition might regulate organ-specific autoimmunity, according to a recent report in *The Journal of Experimental Medicine*.

Ulrike Kuckelkorn and colleagues set out to investigate why self-antigens that are expressed ubiquitously can drive organ-specific autoimmune responses. They used a mouse model in which T-cell-receptor-transgenic CD8⁺ T cells that cross-react with the ubiquitous self-antigen heat-shock protein 60 (Hsp60) are transferred into T-cell-deficient mice. The resulting autoimmune response is restricted to the small intestine, despite similar migration of transgenic T cells to the colon. The authors wondered if tissue-specific differences in the proteolytic processing of Hsp60 epitopes might account for this.

A key player in the generation of peptide epitopes that are recognized by CD8⁺ T cells is the proteasome, which consists of a cylindrical active core with a regulatory complex at each end. The core — which is known as the 20S proteasome — consists of two α -rings and two β -rings, each of which is formed by seven subunits. The basic 20S proteasome is expressed constitutively. However, interferon- γ induces the expression of components that replace the constitutive β -subunits; this forms the ‘immunoproteasome’, which has altered activities.

The authors examined the composition of 20S proteasomes from various tissues by two-dimensional gel electrophoresis. Comparing the small intestine with the colon, they found organ-specific expression patterns of the inducible β -subunits. Differentially expressed variants of $\alpha 4$ and $\alpha 6$ subunits (which interact with the regulatory complexes) were identified also.

Proteasomes from different organs were shown to have distinct cleavage-site preferences for the processing of Hsp60 *in vitro*. Notably, the Hsp60-derived peptide that is recognized by the pathogenic T cells was produced most efficiently by the proteasomes of the small intestine.

But, are the quantitative and qualitative differences in peptide production relevant to CD8⁺ T-cell recognition? This was tested in a chromium-release assay, in which target cells were pulsed with peptide products that had been generated *in vitro* by proteasomes isolated from various tissues. Only target cells that had been pulsed with peptides processed by small-intestine-derived proteasomes were lysed by the transgenic T cells.

The authors propose that this organ-specific generation of epitopes, “represents an important mechanism to control the reactivity of autoreactive CD8⁺ T cells that escape thymic deletion”.

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References and links

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