milestones

Milestone 11

Thelper cells: a house divided

n 1986, Timothy Mosmann, Robert Coffman and colleagues at DNAX Research Institute stimulated a panel of mouse $CD4^+$ T cell lines and discovered two subpopulations of murine $CD4^+$ T cells with distinct profiles of cytokine secretion that they defined as T helper 1 (T_H1) or T helper 2 (T_H2) subsets. This pivotal finding noted that T_H1 cells secrete the cytokines IL-2, IL-3 and IFN γ , while T_H2 cells produce IL-3, IL-4 and IL-5. This finding led to a paradigm shift in our understanding of CD4⁺ T cell biology and the emergence of the complex and heterogeneous composition of the CD4⁺T cell lineage.

Owing to the emergence of these two distinct T_H cell lineages, one aspect of research focused on understanding and establishing the functional role of these cells. A study by Phillip Scott and colleagues showed that in mice infected with *Leishmania major*, adoptive transfer of IFNy-producing T_H1 cells could confer significant protection against disease, while transfer of IL-4⁺IL-5⁺ T_H2 cells exacerbated parasitic lesions. Consistent with these findings, Hienzel et al. reported a strain-specific response to *L. major* infection, observing clearance of infection in C57BL/6 mice within in a few weeks, whereas in Balb/c mice, the authors documented substantive disease progression. It was also shown that the underlying cause of this differential response was linked to an increased expression of IFN γ mRNA in recovering mice and an increased expression of IL-4 and IL-5 mRNA in mice that displayed progressive infection.

Following these discoveries, William Paul and colleagues uncovered the critical role of the cytokine milieu during the development of CD4⁺ T_H cell responses. They and others found that naive CD4⁺ T cells induced the formation of T_H2 cells and were linked to the phosphorylation of STAT6 when treated with IL-4 and IL-2. By contrast, stimulation with IFN γ resulted in preferential development of T_H1 cells and inhibition of T_H2-related cytokines. Further work also showed a role for IL-12 in the production of IFN γ , which drives the skewing of naive T cells

"a paradigm shift in our understanding of CD4⁺T cell biology and the emergence of the complex and heterogeneous composition of the CD4⁺T cell lineage"



toward the $T_H I$ lineage. Subsequent work also showed that $T_H 2$ cells secreted a regulatory cytokine identified as IL-10, which effectively led to the suppression of T cell proliferation and cytokine secretion.

Evidence for T_H subsets in humans was obtained in 1991 when the team of Sergio Romagnani reported distinct T_H1 and T_H2 populations in humans, which was consistent with the initial reports in the murine setting. Further studies suggested that bacterial and viral antigens result in T_H1-type responses, and fungal or helminth-derived antigens resulted in the elicitation of T_H2 responses. Taken together, these and other studies built a consensus in the association of T_H1 responses with intracellular pathogens and a requirement for support for cytolytic-type responses, whereas T_H2 responses were linked to extracellular pathogens and allergens owing to their ability to support strong antibody responses.

Along with characteristic cytokines, the $T_H I/T_H 2$ subsets are also associated with characteristic lineage-specific transcription factors. In 1997, Zheng and Flavell identified GATA3 as the master transcription factor that regulates the $T_H 2$ subset. The same study also reported that silencing of GATA3 in cell lines resulted in the loss of $T_H 2$ cytokines, and showed that naive T cells committed to the $T_H 1$ pathway also downregulate GATA3 expression. Subsequent work from Laurie Glimcher and colleagues further identified T-bet as the master transcription factor for $T_H 1$ cells.

Enabled by these seminal discoveries and others, we now have a clearer view of the composition, diversity and immunological function of the CD4⁺ T cell compartment. $T_H 1$ and $T_H 2$ cells, and the subsequent other T_H cell lineages (see Milestones 15, 18 and 19), now have well-defined roles in critical aspects of immunological homeostasis, immunopathology and immune regulation, and this positions them as key therapeutic targets in the modulation of the immune response to combat disease.

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Milestone study

Mosmann, T. R. et al. Two types of murine helper T cell clone. I. Definition according to profiles of lymphokine activities and secreted proteins. J. Immunol. **136**, 2348–2357 (1986)

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