

Neutralization of interleukin-17 produced by gamma delta T cells constrains inflammation in experimental biliary atresia

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Biliary atresia (BA) is a rare disease of the infant with unknown pathogenesis. It is characterized by inflammatory, progressive destruction of the biliary system leading to liver fibrosis and progressive deterioration of liver function. Interleukin-17a (IL-17) has been identified as a cytokine driving inflammatory and autoimmune processes. We investigated the role of IL-17 and IL-17 producing cell populations in the pathogenesis of experimental and human BA.

In the rotavirus induced BA mouse model, symptomatic animals had a significantly increased hepatic transcription of IL-17. We identified gamma delta ($\gamma\delta$) T cells as the exclusive source of IL-17, while classical Th17 cells were completely absent. The increased number of IL-17⁺ $\gamma\delta$ T cells in BA⁺ animals was associated with an up-regulation of typical markers of the IL-17-axis, such as IL17a, IL17f, ROR γ t, CCR6 and the IL-23-receptor. *In vivo*, blockage of IL-17 by administration of monoclonal antibodies ameliorated the clinical course of disease, improved survival and serum bilirubin, and reduced liver inflammation.

In human infants with BA, hepatic transcription of IL-17 was significantly up-regulated compared to patients with

other neonatal cholestatic diseases, while no differences in IL-17 levels were detected in patient sera.

Taken together, IL-17 released by lymphocytes bearing the $\gamma\delta$ T cell receptor appear to be a causative factor in the inflammatory destruction of the biliary system in experimental BA. Furthermore, our data suggest an important role of the IL-17 axis in human BA. Thus, targeting the IL-17 axis could be a promising approach for therapeutic interventions.