Autoimmunity and cancer immunosurveillance in the biliary tree

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ntroduction Primary biliary cholangitis (PBC)¹ and primary sclerosing cholangitis (PSC)² are two chronic inflammatory diseases of the biliary tract. Chronic inflammation is known to be one of the main risk factors for cancer onset. In this line, PSC is the first etiology of cholangiocarcinoma (CCA)³ in Western countries. But surprisingly, patients with PBC close to never develop CCA. Since PBC harbors an autoimmune component, we hypothesised that PBC-associated autoimmunity could fuel cancer immunosurveillance and thus prevent CCA appearance.

1. Validation of the murine models of cholangitis



means of ANOVA test Tukey's pairwise multiple comparison.

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2. PBC protects against CCA in a specific & T-cell dependent manner

Figure 2. A. Experimental scheme. Growth of subcutaneous syngeneic CCA tumors in control, PSC or PBC mice (B-D) or PBC mice without or with injections of antibodies targeting either CD4 and/or CD8 or CD20, to deplete CD4⁺ and/or CD8⁺ T or B cells, respectively (H-M). Growth of subcutaneous syngeneic tumors of hepatocellular carcinoma (HCC) (E), non-small-cell lung cancer (NSCLC) (F), and fibroscarcoma (G) in control and PBC mice. Graphs show mean (± SEM) (B, E-H, K) and individual (C-D, I-J, L-M) tumor growth curves. P-values were calculated by means of a linear mixed-effect model.

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3. PBC protection against CCA relies on type-1/2 immune responses



4. Hepatic and CCA tumor tissues share similar enriched TCR clonotypes upon PBC

Conclusion Our data demonstrated that PBC protects against CCA outgrowth through an active mechanism mainly relying on both CD4⁺ and CD8⁺T-cells, and in a lesser extent on B-cells. PBC-associated antitumor activity was specific to CCA. Type-1 and -2 immune responses, relying on the cytokines IFNy and IL4 respectively, were mediating CCA immunosurveillance upon PBC. Finally, single-cell TCR/RNA-sequencing analyses revealed an enrichment of T cells with identical TCR in both liver and ectopic CCA tumor of a PBC mice. Altogether, our data provide mechanistic insights into an overlap between autoimmunity and cancer immunosurveillance in the biliary tree.

Figure 3. A. Experimental scheme. Relative expression of Ifng (B) and II4 (E) genes measured by RT-qPCR within CCA tumors from control, PBC and PSC mice at day 56. Graphs show individual and mean (± SD) values. P-values were calculated by ANOVA with Tukey's pairwise multiple ^b comparison. **C-D, F-G.** Growth of subcutaneous syngeneic CCA tumors in control and PBC mice, or PBC mice injected with antibodies targeting either IFNy or IL4. Graphs show mean (± SEM) (C, F) and individual (D, G) tumor growth curves. P-values were calculated by means of a linear mixed-effect model.

Figure 4. T cell clones infiltrating CCA, hepatic and blood tissues of a PBC mouse were characterized by single-cell TCR sequencing coupled with single-cell RNAsequencing. A. Venn diagram showing the number of TCR clonotypes overlapping between tumor, liver and blood. **B.** Number of TCR clonotypes enriched (i.e. >0,1% of T cells) only in tumor (n=103), or liver (n=58), or both tissues (n=25). C. Proportion in the liver versus CCA tumor of each of the 25 TCR clonotypes enriched and shared between both tissues. **D.** Phenotypic clustering of total T lymphocytes sorted out from CCA and hepatic tissues of a PBC mouse. E. Phenotypic clustering of the 25 TCR clonotypes enriched in both CCA tumor and liver upon PBC.

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