

SINGLE-CELL RNA STUDY OF TRANSCRIPTIONAL ACTIVITIES ASSOCIATED WITH ANAKINRA PROPHYLAXIS AND AXICABTAGENE CILOLEUCEL IN PATIENTS WITH RELAPSED OR REFRACTORY LARGE B-CELL LYMPHOMA

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Background Chimeric antigen receptor T-cell (CAR-T) therapy has shown unprecedented treatment outcomes for B-cell malignancies. The wider utilization of CAR-T, however, is limited by CAR-T-induced adverse events including cytokine release syndrome (CRS). Tocilizumab, a monoclonal antibody blocking Interleukin (IL)-6 receptor, is the only approved treatment for CRS. Recent animal studies suggest that anakinra, a recombinant form of IL-1 receptor antagonist, could have potential benefit in managing toxicity, with or without tocilizumab. We have opened a clinical trial (NCT04150913) testing the use of prophylactic anakinra in patients with relapsed or refractory large B-cell lymphoma eligible to receive axicabtagene ciloleucel (axi-cel) as per the registration study ZUMA-1. Here, we used single-cell RNA sequencing (scRNA-seq) on clinical samples of the first ten subjects on the study¹ to probe the molecular pathways altered by anakinra and to discover potential mediators of breakthrough cases of CRS.

Methods We conducted scRNA-seq on the peripheral blood mononuclear cells (PBMCs) of 10 patients 7 days post infusion (D7) and the infusion products (IP). We compared data from the trial subjects to 4 control samples from the same institution along with our previously generated scRNA-seq IPs and D7 PBMCs from 19 subjects that were treated with axicel without prophylaxis for toxicity.² We utilized pseudo-bulk differential expression analysis of major immune cell types to reveal transcriptional signals associated with CRS and anakinra treatment.

Results Our study revealed that IL-4 and IL-10 anti-inflammatory pathways in IPs of both anakinra and non-anakinra cohorts were negatively associated with breakthrough CRS. We further observed that the same pathways were enriched at D7 in anakinra-treated CAR+ CD4+ T cells populations. Anakinra prophylaxis had little effect on the overall CAR+ T-cell compositions from IPs to D7, but was associated with an increased proportion of CAR+ regulatory T cells (Tregs). Expression of interferon gamma (IFN γ) pathways, cytokine levels of IFN γ and IFN γ -induced protein 10 (CXCL10) in CD14+ monocytes were significantly enriched in patients with breakthrough CRS treated with tocilizumab in the anakinra cohort. Differential cell-cell interaction analysis further showed the association of IFN γ ligand receptor activities with breakthrough toxicity uniquely among anakinra- and tocilizumab-treated patients.

Conclusions We identified key molecular pathways and immune cell populations that were possibly modulated by anakinra, including the upregulation of IL-10 signaling pathway in CAR+ CD4+ T cells and increased abundance of Tregs. IFN γ enrichment in patients with breakthrough CRS further suggests that this pathway is also targetable and not inhibited by anakinra alone.

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Ethics Approval The clinical study complies with IRB-approved protocol (19-348) at the Dana-Farber/Harvard Cancer Center.

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