

## PERIPHERAL IMMUNOLOGICAL DYNAMICS OF RESPONSE IN PANCREATIC CANCER TREATED WITH IMMUNE CHECKPOINT INHIBITORS

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**Background** Pancreatic cancer (PC) is the fourth leading cause of cancer death in the United States,<sup>1</sup> as it is highly resistant to conventional treatments,<sup>2–4</sup> and nearly all patients develop metastases and progress. Immunotherapy with immune checkpoint inhibitors (ICI) has shown strong, durable responses in patients with various types of cancer,<sup>5</sup> but these successes have not yet completely translated to PC. Here we analyzed samples from a phase II clinical trial where patients with PC were treated with a combination of radiotherapy and ICI to understand the peripheral immunological dynamics of responding patients.

**Methods** Peripheral blood mononuclear cells (PBMCs) were isolated from patients with pancreatic cancer enrolled in the CheckPAC protocol (NCT02866383<sup>6</sup>) and subjected to multi-omic single-cell analyses. In detail, CITESeq was performed on baseline and on-treatment (+8 weeks) blood samples from responders (n=5) and non-responders (n=6) in the trial to obtain both gene expression and protein expression data. The data was further analyzed with the R package Seurat.<sup>7</sup>

**Results** By exploring gene and surface protein expression on multiple immune cell subsets, we identified increased numbers of effector cells (CD8 T cells, Natural Killer cells, and  $\gamma\delta$  T cells) in the PBMCs of responders, consistent with increased immune activation leading to tumor regression. The expression of Granulysin in effector cells (GNLY) strongly correlated with clinical responses, confirming previous reports where increased levels of GNLY have been associated with clinical responses.<sup>8</sup> Specifically,  $\gamma\delta$  T cells were the most enriched cell type for GNLY expression, with a statistically significant higher expression in responders at baseline, suggesting the role of GNLY+  $\gamma\delta$  T cells as a predictive biomarker population. Furthermore, surface protein analyses confirmed that the specific subset of V $\delta$ 2 $\gamma$ 9 T cells was associated with response and showed high expression of other effector molecules like Granzyme B or Tumor Necrosis Factor, confirming its highly cytotoxic phenotype. Of note, analysis of TCGA datasets showed that PC has a high expression of the genes BTN3A1 and BTN2A1, which are necessary for phosphoantigen presentation and tumor recognition by V $\delta$ 2 $\gamma$ 9 cells.<sup>9</sup> Expression of GNLY, as well as other cytotoxic markers, will further be validated by flow cytometry on the responding patients.

**Conclusions** This data sheds light on the peripheral immunological dynamics of patients with PC who respond to ICI therapy and suggest a highly cytotoxic subset of GNLY+ V $\delta$ 2 $\gamma$ 9 T cells as the potential driver of response.

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**Ethics Approval** This study was conducted using PBMCs from patients enrolled in clinical trials performed at the National Center for Cancer Immune Therapy (CCIT-DK), Department of Oncology, Copenhagen University Hospital, Herlev, Denmark. All procedures were performed in compliance with clinical protocols approved by the Ethics Committee of the Capital Region of Denmark (Reference H-20070020) and national regulations for biomedical research, including appropriate data protection of human participants (Reference P-2021–303). Written informed consent was provided by all patients before obtaining any samples.

<http://dx.doi.org/10.1136/jitc-2023-SITC2023.1030>