Akhuba et al. Cancer Drug Resist 2023;6:327-31 DOI: 10.20517/cdr.2022.142

Commentary



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# Major hurdles of immune-checkpoint inhibitors in pancreatic ductal adenocarcinoma

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How to cite this article: Akhuba L, Tigai Z, Shek D. Major hurdles of immune-checkpoint inhibitors in pancreatic ductal adenocarcinoma. *Cancer Drug Resist* 2023;6:327-31. https://dx.doi.org/10.20517/cdr.2022.142

Received: 15 Dec 2022 First Decision: 28 Feb 2023 Revised: 11 May 2023 Accepted: 29 May 2023 Published: 30 May 2023

Academic Editors: Godefridus J. Peters, Enrico Mini Copy Editor: Dong-Li Li Production Editor: Dong-Li Li

## Abstract

In 2030, pancreatic ductal adenocarcinoma (PDAC) will become the second leading cause of cancer-related mortality in the world. Unfortunately, neither conventional chemotherapy nor novel immunotherapeutic strategies can provide durable responses and the survival prognosis remains very low. PDAC is notorious for its immune-resistant features and unique genomic landscape facilitating tumor escape from immunosurveillance. Novel immune-checkpoint inhibitors (ICI) failed to show promising efficacy and other multi-modal approaches are currently being validated in multiple clinical trials. In this paper, we provide our opinion on the major mechanisms responsible for PDAC resistance to ICI therapy and provide our view on future strategies which may overcome those barriers.

Keywords: Pancreatic cancer, immune-checkpoint inhibitors, tumor resistance, microenvironment

Pancreatic ductal adenocarcinoma (PDAC) represents a major challenge in modern oncology<sup>[1]</sup>. It is predicted that by 2030 PDAC will become the second leading cause of cancer-related death<sup>[2]</sup>. Surgery is curative at earlier stages, whereas advanced or metastatic stages are almost impossible to treat<sup>[3]</sup>.

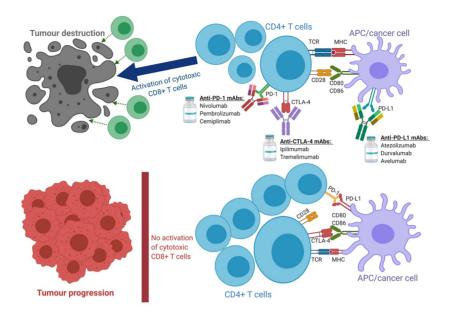


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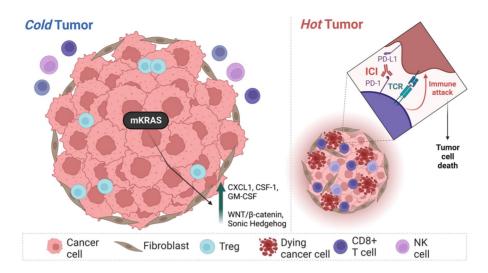




**Figure 1.** Mechanisms of immune-checkpoint inhibitors. ICIs target unique inhibitory checkpoint molecules expressed by T- and antigen-presenting cells. By blocking those receptors, ICIs promote the proper induction and differentiation of T cell-mediated immunity. In contrast, the absence of ICIs results in successful priming of checkpoint receptors with their ligands, thus inhibiting TCR activation overall, leading to cancer escape from immunosurveillance. APC: antigen-presenting cell; CD: cluster of differentiation; CTLA-4: cytotoxic T-lymphocyte associated antigen 4; mAb: monoclonal antibody; MHC: major histocompatibility complex; PD-1: programmed cell death protein 1; PD-L1: programmed cell death protein 1 ligand 1; TCR: T cell receptor.

Conventional chemotherapy can only provide a short partial remission with 5-year overall survival (OS) of less than 9% in patients with advanced PDAC<sup>[4]</sup>. Recent discoveries in cancer immunology have led to the successful use of immune-checkpoint inhibitors (ICIs) in treating advanced solid malignancies. ICIs are monoclonal antibodies that target immune checkpoints such as cytotoxic T-lymphocyte associated antigen 4 (CTLA-4), programmed cell death protein 1 (PD-1) with its ligands PD-L1/L2 and other expressed by antigen-presenting cells (APCs) and T cells [Figure 1]<sup>[5]</sup>. ICIs have shifted treatment paradigms for melanoma, non-small cell lung cancer (NSCLC), and hepatocellular carcinoma<sup>[6,7]</sup>. Unfortunately, PDAC has shown incredible resistance to immunotherapy<sup>[8]</sup>. To date, US Food and Drug Administration (FDA) has only approved PD-1 inhibitor *pembrolizumab*, albeit only for patients with high microsatellite instability (MSI-H)<sup>[9]</sup>. Unfortunately, the majority of patients (~ 97%) with microsatellite stable status (MSS) are not benefited from ICIs and their outcomes remain critically poor<sup>[10]</sup>. Early trials combining chemotherapy with ICIs also fail to show any superior efficacy in MSS patients<sup>[9]</sup>. This paper provides an opinion on factors responsible for PDAC resistance to ICIs and potential strategies to overcome this issue.

Classically, PDAC has an immunologically "cold" tumor microenvironment<sup>[11]</sup> characterized by abundant infiltration of myeloid cells and a small number of infiltrating T- and NK (natural killer) cells [Figure 2]. A few studies suggested that focal adhesion kinases (FAK) can regulate the fibrotic features of cold tumors, including the immunosuppressive microenvironment<sup>[12,13]</sup>. The data from *in vitro* studies on the synergistic efficacy of FAK + PD-1 inhibitors showed promising responses and resulted in further testing of this regimen in clinical trials. Other factors of resistance are low mutational burden and complex immunosuppressive features able to inhibit T cell priming and trafficking, resulting in lower efficacy of immunotherapy<sup>[14]</sup>.



**Figure 2.** Mechanisms of pancreatic cancer resistance to immune-checkpoint inhibitor therapy. Pancreatic ductal adenocarcinoma is known as a tumor with a "cold" microenvironment characterized by a small number of CD8+ T- and NK cells, an abundance of regulatory T (immunosuppressive) cells, and poor response to ICI therapy. Mutation in *KRAS* gene (mKRAS) allows pancreatic cancer cells to induce expression of granulocyte-macrophage colony-stimulating factor (GM-CSF), chemokine C-X-C motif ligand 1 (CXCL1) and C-C motif chemokine ligand 4 (CCL4) playing a crucial role in immunosuppression. Moreover, mKRAS leads to upregulation of WNT/ $\beta$ -catenin pathway and Sonic Hedgehog pro-inflammatory pathways overall, inhibiting the ICI therapy.

Stromnes et al. reported that analysis of tumor samples revealed that PDAC has a lower number of effector T cells and lower clonality of T cell receptors as compared to other solid tumors that can be successfully managed by ICIs<sup>[15]</sup>. Genome studies have established that PDAC almost ubiquitously has activating KRAS (Kirsten ras oncogene) mutations<sup>[16]</sup>. Conventionally, mKRAS is known to be associated with tumor proliferation and metastasis; however, recent results of high-throughput studies have established that mKRAS may orchestrate downstream signaling responsible for immunosuppression<sup>[17]</sup>. A few in vitro studies established that mKRAS inhibits the expression of MHC-I, CD47, and PD-L1<sup>[18,19]</sup>. It is known that PD-L1 is a crucial marker for ICI efficacy in non-small cell lung cancer<sup>[20]</sup>. Perhaps the lower expression of checkpoint proteins (targets) negatively impacts ICI therapy and explains its lower effectiveness in PDAC patients. Secondly, mKRAS can upregulate the expression of GM-CSF and CXCL1, which are involved in the recruitment of myeloid-derived suppressor cells known for their immunosuppressive features<sup>[21,22]</sup>. Furthermore, mKRAS can downregulate the expression of CCL4 via WNT/β -catenin pathway<sup>[23]</sup>. CCL4 is an important factor for recruiting dendritic cells; major APCs require FOR priming T cell response and activating the cytotoxic cascade<sup>[24,25]</sup>. A lower number of APCs impacts the tumor escape from immunosurveillance. Additionally, mKRAS promotes signaling via the Sonic Hedgehog pathway and can induce expression of matrix metalloproteinase 7 (MMP-7)<sup>[26]</sup> as well as selectively target lysosomal degradation of MHC-I molecules through an autophagy-dependent mechanism, thus negatively impacting ICI therapy<sup>[19]</sup>. Overall, it results in chronic inflammation and proliferation of the fibrotic stroma, thus complicating T cell trafficking<sup>[27]</sup>. The development of mKRAS-directed strategies may one day overcome this critical resistance mechanism and result in higher effectiveness of ICIs in PDAC.

In summary, PDAC is among the most immune-resistant tumors. Recent discoveries in understanding key elements of PDAC resistance to ICI therapy, including FAK<sup>[28]</sup>, mKRAS and other novel molecules<sup>[29]</sup>, have reshaped our view on future approaches for PDAC treatment. To effectively treat PDAC, it is crucial to elucidate the rational combinatorial approach(es) targeting both checkpoint proteins and non-redundant mechanisms of PDAC resistance, such as mKRAS. Moreover, novel therapeutic strategies should be selected based on patient's individual genotype, which is responsible for high phenotypic heterogeneity observed

across PDAC patients. Finally, mKRAS remains the bull's eye for PDAC immunologic resistance; thus, the combinatorial approach of ICI + MEK (mitogen-activated protein kinase) inhibitors should be thoroughly studied in randomized trials. The synergistic effect of both drugs may improve clinical outcomes for PDAC patients in the near future.

#### DECLARATIONS

#### Authors' contributions

Wrote and reviewed this manuscript: Akhuba L, Tigai Z, Shek D All authors equally contributed to this work.

#### Availability of data and materials

Not applicable.

# Financial support and sponsorship

None.

#### **Conflicts of interest**

All authors declared that there are no conflicts of interest.

# Ethical approval and consent to participate

Not applicable.

#### **Consent for publication**

Not applicable.

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