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## **$\gamma\delta$ T cells turn the tables on immune-evasive colon cancer**

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### **Summary**

Why is checkpoint blockade immunotherapy still effective in tumors that are unrecognizable to CD8<sup>+</sup> T cells? In a recent study published in *Nature*, de Vries *et al.* provide evidence that a lesser-known T cell population, called  $\gamma\delta$  T cells, may mediate beneficial responses to immune checkpoint blockade when cancer cells lose expression of HLA molecules.

### **Main body**

Across all cancer types, those tumors with high mutational burden are particularly responsive to immune checkpoint blockade (ICB). These cancer types include melanoma, lung cancer, and DNA mismatch repair-deficient (MMR-d) colon cancer, among others. In these mutationally high cancer types, boosting (neo)antigen-specific CD8<sup>+</sup> T cell activity with anti-PD-1 prevents immunosuppression and increases T cell-mediated killing. However, these tumors frequently exhibit mutations in  $\beta$ 2-microglobulin (*B2M*), a component of the major histocompatibility (MHC)-I complex, and other proteins involved in antigen processing and presentation, making them invisible to CD8<sup>+</sup> T cells. Fortunately, other lymphocytes, like  $\gamma\delta$  T cells, can recognize mutated cells lacking MHC molecules.  $\gamma\delta$  T cells are distinct from conventional T cells in many ways, including the genes that encode their T cell receptor chains and their inability to bind MHC molecules, but  $\gamma\delta$  T cells also share functions with conventional T cells, such as inducing death of cancer cells (1). What we know about the mechanisms by which  $\gamma\delta$  T cells counteract cancer progression lags far behind what is known about CD8<sup>+</sup> T cells. Therefore, the recent study by de Vries *et al.* showing that  $\gamma\delta$  T cells may participate in ICB response in MMR-d colon cancer patients is a major advance for the cancer immunotherapy field (2).

From a small cohort of 21 patients, de Vries *et al.* found that 95% of *B2M*-mutated, MMR-d tumors are still responsive to ICB despite their inability to engage CD8<sup>+</sup> T cells (2). To gain insight into this seemingly contradictory observation, the authors used gene expression data from MMR-d colon, gastric, and endometrial tumors to compare differences between *B2M* wild-type and *B2M*-mutated tumors. This analysis showed that *B2M*-mutated tumors contain increased numbers of  $\gamma\delta$  T cells. Further phenotypic examination of  $\gamma\delta$  T cells from MMR-d colon cancers revealed that some subsets of  $\gamma\delta$  T cells express PD-1 and other checkpoint molecules. The PD-1-expressing  $\gamma\delta$  T cell subsets were superior at killing *B2M*-deficient colon cancers than  $\gamma\delta$  T cell subsets

lacking PD-1 expression, highlighting the heterogeneity of  $\gamma\delta$  T cell capabilities. In addition, de Vries *et al.* investigated colon cancer tissue from the NICHE trial that consisted of pre- and post-ICB MMR-d samples. Those tumors with *B2M* mutations displayed increased  $\gamma\delta$  T cell numbers after ICB, whereas  $\gamma\delta$  T cell numbers remained the same before and after ICB in *B2M* wild-type tumors. Taken together, these data provide compelling evidence that ICB boosts  $\gamma\delta$  T cell activity in *B2M*-mutated tumors where they participate in tumor control.

One important aspect of the study by de Vries *et al.* is that not all tumor-associated  $\gamma\delta$  T cells are the same. Human  $\gamma\delta$  T cells are defined by the  $\delta$  chain they express: V $\delta$ 1 and V $\delta$ 3 cells are mostly tissue resident, whereas V $\delta$ 2 cells circulate in blood. The majority of studies on the anti-tumor functions of  $\gamma\delta$  T cells are focused on circulating V $\gamma$ 9V $\delta$ 2 cells (3); although, the killing ability of V $\delta$ 1 cells has been established for some time (4). de Vries *et al.* demonstrate that tumor-infiltrating V $\delta$ 1 and V $\delta$ 3 cells can express PD-1 (to a limited degree) and these populations are endowed with killing ability of *B2M*-mutated cancer cells, while PD-1 is mostly absent from V $\delta$ 2 cells that are inefficient at triggering cancer cell death (2).

The authors also show that PD-1-expressing V $\delta$ 1 and V $\delta$ 3 cells can use NKG2D to recognize cancer cells, but whether other molecules are involved in recognition remain unknown. Similarly, the data raise questions regarding the role (if any) of the  $\gamma\delta$ TCR in cancer cell recognition and killing. Gut-resident V $\delta$ 1 cells that dimerize with the V $\gamma$ 4 chain bind members of the butyrophilin family, BTNL3 and BTNL8, which regulates their development and phenotype (5, 6); however, our lab has found that expression of BTNL3 and BTNL8 is lost during cancer progression (7). Whether BTNL3 and BTNL8 are required for V $\delta$ 1 cell killing of colon cancer cells or whether expression of BTNL3 and BTNL8 are maintained specifically in *B2M*-mutated colon cancer is not clear. As BTNL3 and BTNL8 expression is restricted to gut tissue, a TCR-independent role would be advantageous to engage V $\delta$ 1 cells in other *B2M*-mutated cancer types beyond colon cancer. The TCR on V $\delta$ 3 cells can bind the MHC class I-related protein, MR1, dimerized to B2M that presents vitamin B metabolites, so cancer cells would need to express these metabolites (and B2M) for V $\delta$ 3 cell engagement (8). These obscurities on  $\gamma\delta$ TCR importance further confuse the role of PD-1 in these cells. PD-1 signaling suppresses TCR activation in conventional T cells, but whether the same mechanisms occur downstream of V $\delta$ 1 and V $\delta$ 3 TCR signaling is unknown. Recently, our lab discovered that PD-1 signaling inhibits mouse IL-17A-producing  $\gamma\delta$  T cells independently of TCR (9), highlighting the possibility that PD-1 suppression of human V $\delta$ 1 and V $\delta$ 3 cells may work differently than in conventional T cells.

There are multiple implications for the de Vries *et al.* study, such as using  $\gamma\delta$  T cells as biomarkers of ICB responsiveness, expanding this work beyond colon cancer to other *B2M*-mutated cancer types, and testing cellular-based  $\gamma\delta$  T cell therapies together with ICB in patients with *B2M*-mutated tumors. In the meantime, larger cohorts of patients are necessary to verify the reported observations, especially in the context of pre- and post-ICB. To gain further insight into these mechanisms and test new immunotherapy approaches, mouse

models of cancer will be highly advantageous, given the conserved biology between humans and mice in gut-resident  $\gamma\delta$  T cell and BTNL interactions (5, 6). The interest in  $\gamma\delta$  T cells for cancer immunotherapy has increased exponentially in recent years (10), and the findings by de Vries *et al.* expand the potential of ICB beyond CD8<sup>+</sup> T cells to the underappreciated population of  $\gamma\delta$  T cells, lending credence to the notion that  $\gamma\delta$  T cells are viable and effective tools for cancer immunotherapy.

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### **Declaration of Interest**

The authors declare no competing interests.