

# **Resistance mechanisms to PD-1 inhibition in melanoma**

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Immune checkpoint inhibition using antibodies against cytotoxic T-lymphocyte-associated protein 4 (CTLA-4) and programmed death 1 (PD-1) have substantially improved treatment outcomes for patients with advanced melanoma. (1-3). However, despite the remarkable durability of objective responses induced by these treatments, progressive disease occurs in about a quarter of the patients within the first 3 years (4). Understanding the molecular mechanisms of acquired resistance by focused comparison of biopsy samples from paired baseline and relapsing lesions should open strategies for the rational design of salvage therapies and may guide mechanistic biomarker studies for the identification of patients, before the initiation of treatment, who are unlikely to have a response.

A recent study (5) has identified two likely mechanisms for the development of acquired resistance to PD-1 blockade in patients with melanoma. Clonal evolution of tumors to silence effector signaling pathways involved gamma interferon activation by mutations in the  $\mathcal{J}AK1$  and  $\mathcal{J}AK2$  genes, or by affecting antigen presentation through the B2M gene.

The report focuses on 4 advanced melanoma patients treated with pembrolizumab who had tumor progression after an initial confirmed objective response that lasted for at least 6 months; these patients were selected from 15 initial responders who later progressed and from whom serial tumor biopsies were available prior to treatment and at the time of progression. In order to identify resistance mechanisms to PD-1 blockade, analyses of the serial tumor samples were performed and included pathological analysis as well as genomic studies of DNA and RNA. Immunohistochemical studies in pre-treatment and relapse tumor samples of patients 1, 2, and 3 demonstrated CD8 T-cell infiltration at the invasive margin that colocalized with PD-L1 expression on surrounding macrophages and melanoma cells. In two of the 4 patients studied, resistanceassociated loss-of-function mutations in genes for interferon receptor-associated JAK1 or JAK2 were identified, concurrent with suppression of the wild-type allele. In a third patient, a truncating mutation in the gene for the antigen-presenting protein beta-2-microglobulin (B2M) was found. The JAK1 and JAK2 truncating mutations resulted in a lack of response to interferon-gamma and loss of its antiproliferative effects, and the B2M truncating mutation led to loss of surface expression of major histocompatibility complex (MHC) class I (6).

Using whole genome sequencing, the authors identified loss-of-function mutations in kinases linked to the interferon receptor pathway. Comparing the response of the primary cell lines that were derived from the tumor of the patients-at baseline and at the time of relapsethe authors found an absence of JAK1 and JAK2 protein expression, respectively, following relapse, and a consequent lack of response to interferon gamma. The third patient's relapsed tumor showed a mutation in B2M, the gene for B2M, which has been previously implicated in resistance to immunotherapy (7-10). Without this protein, tumor cells are no longer able to present antigens to T cells, a critical function of immune recognition and an effective anti-tumor T cell response. Therefore, while mutations in JAK1 and JAK2 suggest that the tumor is insensitive to T cell mediated cytolysis, a mutation in B2M means that the T-cells cannot identify the tumor in the first place. The fourth patient did not exhibit any of these mutations,

suggesting that other genetic variations may be involved in the acquired resistance to immunotherapy.

Two pathways were identified in the functional inactivation of CD8 T cells during the development of acquired immune resistance. The first pathway-identified in relapsed patients 1 and 2-leads to a decrease of tumor cell response to interferon gamma that was mediated by loss of function mutations in the genes encoding JAK1 or JAK2. Previous in vitro studies had already shown that the tumor cells could still be identified by appropriate T cells. Those produce interferon- $\gamma$ , but the JAK mutations resulted in a complete lack of tumor cell sensitivity to interferon- $\gamma$ evidenced by a lack of phosphorylation of signal transducer and activator of transcription 1 (STAT1) in the interferon- $\gamma$ signaling pathway, lack of interferon gamma-induced expression of MHC class I and PD-L1, and insensitivity to the antiproliferative effects of interferon- $\gamma$  (5,11-13). Using the clustered regularly interspaced short palindromic repeats (CRISPR)-Cas9 approach, melanoma cell lines expressing the cancer testis antigen NY-ESO-1 that lacked JAK1 or JAK2 were created. T cells expressing an NY-ESO-1 transgenic T cell receptor were able to produce IFN-y when exposed to the JAK1 and JAK2 deficient melanoma cell lines however the cell lines were insensitive to IFN-y mediated growth arrest. This finding indicates that the resistance to IFN- $\gamma$  induced cytolysis in the JAK deficient cell lines was due to deficient IFN- $\gamma$  pathway signaling and not a lack of recognition by the T cells. The second pathway-identified in patient 3-linked to a previously identified mechanism for cytotoxic T-cell escape-loss of expression of MHC class I on the tumor cell surface, that in this case was due to a truncating mutation in the gene encoding B2M (8). Consequently, whereas mutations in JAK1 and JAK2 suggest that the tumor is insensitive to T cell inhibition, a mutation in B2M means the T-cells cannot recognize the tumor in the first place.

Similar to targeted therapy, Zaretsky *et al.* show that immunotherapy with anti-PD-1 directed therapy can be subject to acquired resistance that is related to mutations in tumor cells (5). The increase in expression of T-cell immunoglobulin mucin 3 (TIM-3) (14), the presence of somatic aberrations in tumor cells (loss of PTEN) inhibiting the recruitment of T cells (15), and characteristic transcriptional signatures such as regulation of mesenchymal transition, cell adhesion, extracellular matrix remodeling, and angiogenesis (16) in the tumor microenvironment are other recently identified mechanisms for acquired resistance to anti-PD1 therapies.

The study demonstrates impressively how the rapidly emerging tools of precision medicine can be employed to dissect immunotherapy-resistance mechanisms in cancer patients. The elimination of IFN- $\gamma$  sensitivity through JAK1 or JAK2 inactivation is a particularly compelling resistance mechanism in the context of PD-1 blockade given the adaptive upregulation of PD-L1 by tumors in response IFN- $\gamma$  secreted by tumor infiltrating T cells. The B2M mutation identified in a 3<sup>rd</sup> patient is an additional, independent persuasive resistance mechanism. Other mechanisms surely are at play as the lack of a genomic alteration explaining the resistance to PD-1 inhibition in the 4<sup>th</sup> patient illustrates. The by nature correlative analyses can also not rule out additional, non-genomic mechanisms for resistance, even in the patients with JAK inactivation and B2M mutation. Larger cohorts of patients should be studied to confirm these results and will almost certainly reveal additional mechanisms. This research shows that residual cancer cells continue to fight for survival, even after prolonged inhibition by immunotherapy, through evolution to induce resistance, suggesting that we need new ways to determining and estimating residual disease.

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