

IL-34 and Immune Checkpoint Inhibitors Resistant through Macrophages: Pan-Cancer Overview

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Utilizing immune checkpoint inhibitors (ICIs) to induce therapeutic anti-tumor activity is a promising strategy. ¹ However, malignancies often develop immunological resistance, limiting the clinical benefits of these approaches. ² Several studies have indicated several molecular and cellular processes of immunological resistance. Immunosuppression in the local tumor microenvironment (TME) and epigenetic alterations in tumor cells that reduce interferon-gamma (IFN- γ) signaling constitute two major mechanisms involved in tumor progression. ³ In the context of TME, macrophages are the most common kind of immune cell found in solid tumors, accounting for as much as fifty percent of the total mass of the tumor. ⁴ In response to chemotactic chemicals made in the cancer microenvironment, macrophages (monocyte-derived tumor-associated macrophages, or TAMs) are sent from the blood to the tumor tissue. Infiltration of macrophages into and around the tumor nest is a characteristic that is considered to be one of the most crucial. These interactions between macrophages, tumor cells, and the stromal milieu contribute to the phenotypical polarization of macrophages linked with tumors. The process through which macrophages generate different functional phenotypes in response to certain stimuli and signals arising from their local environment is referred to as “polarization.” “polarization”. Macrophages are capable of polarization into conventionally activated (M1) and alternatively activated (M2) subpopulations. M1-phenotype macrophages are tumor-resistant, whereas M2-phenotype macrophages possess various tumor-promoting activities, including immunosuppression, angiogenesis, and neovascularization. The tumor microenvironment (TME) is frequently generated and continuously maintained by biomarkers secreted by the tumor tissues, and enrichment with immunosuppressive cells, like M2-biased tumor-associated macrophages (TAMs), has been suggested to play crucial roles in innate and acquired ICI resistance. ⁶

Recently, a large humanized model study revealed that interleukin IL-34 dysregulation could lead to resistance to the recently approved immune checkpoint inhibitors ICI, and IL-34 has come to be recognized as a pivotal molecule in regulating the maintenance, expansion, and activity of M2-biased TAMs that exhibit heightened immunosuppressive capabilities. ⁷ **Figure 1**. By single-cell analysis of the IL-34 expressions in 208 single-cell datasets (<https://ngdc.cncb.ac.cn/cancerscem/index>) with 638,341 cells, we showed that the highest expressions were associated with glioblastoma (GBM), lung adenocarcinoma (LUAD), and pancreatic ductal adenocarcinoma (PDAC). **Figure 2A**; **Supplementary Figure 1**. In the context of ICIs resistance, cancer's dysregulation of IL-34 showed how important M2-biased TAMs is in various cancer tissues. Pan-cancer analysis between IL-34 and M2 was conducted and uncovered that IL-34 has a positive correlation with infiltration of M2 among 30 cancer tissues with the highest positive association in prostatic adenocarcinoma (PRAD), testicular germ cell tumors (TCGT) and stomach adenocarcinoma (STAD) and negative correlation in low-grade gliomas (LGG). **Figure 2B**. Further analysis of the correlation difference in IL-34 expression between M1 and M2 with the same cancer types revealed a higher median value with M2 than M1 (0.33 versus 0.27) ($P = 0.002$). **Figure 2C**. **Supplementary Methods and Materials**. As a result, this may lead us to suggest that higher IL-34 expressions are closely related to M2-biased macrophage infiltration and decreased frequencies of the M1 type, and consequently, a higher possibility of resistance to ICI

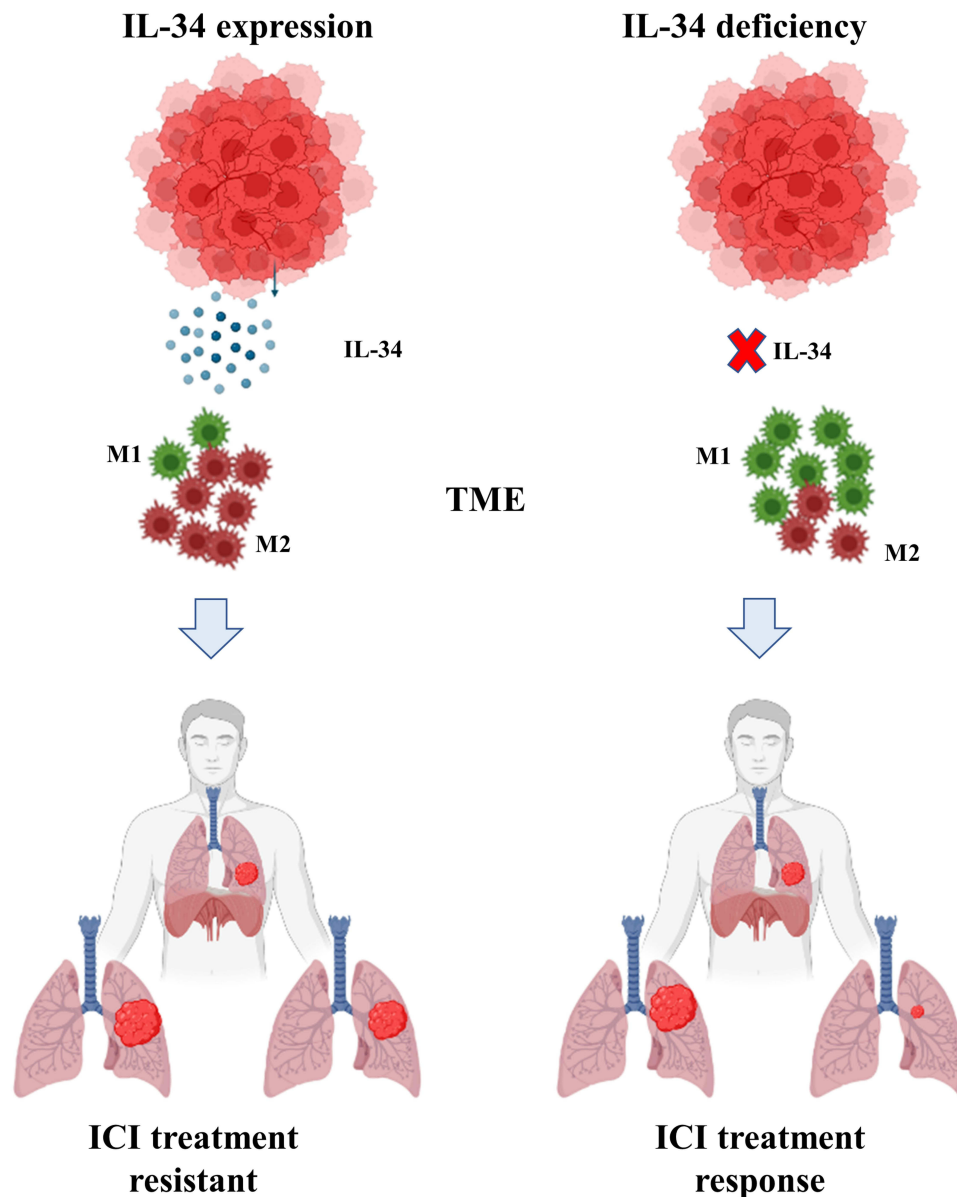


Figure 1 IL-34 expression status and response status in the context of immune checkpoint inhibitors. According to the literature, cases with high IL-34 expression are associated with macrophage M2 infiltration and lower frequencies of M1-type macrophages. Higher response status and decreased tumor tissue are associated with low or depleted IL-34 expression. Created with BioRender.com.

therapy. According to pan-cancer clinical correspondence data, cancer cells release IL-34 to create chemoprotective niches through the paracrine recruitment of M2-polarized TAMs with an elevated protumorigenic phenotype after treatment with ICIs. Therefore, new evidence for a novel targeted biomarker might be investigated by investigating the possibility that IL-34 plays a role in establishing therapeutic resistance to tumor ICIs.⁸ However, the molecular processes that underlie these discrepancies must be addressed in future research. In sum, it is conceivable that IL-34 might be exploited as a new therapeutic to improve the effectiveness of ICIs in treating a variety of cancers. However, more rigorous preclinical and clinical research are required to collaborate these findings.

Data Sharing Statement

We used <https://portal.gdc.cancer.gov/>, <http://ualcan.path.uab.edu/>, and <http://timer.cistrome.org/> for expression association. Also available in [Supplementary Materials and Methods](#) and [Supplementary Figure 1](#).

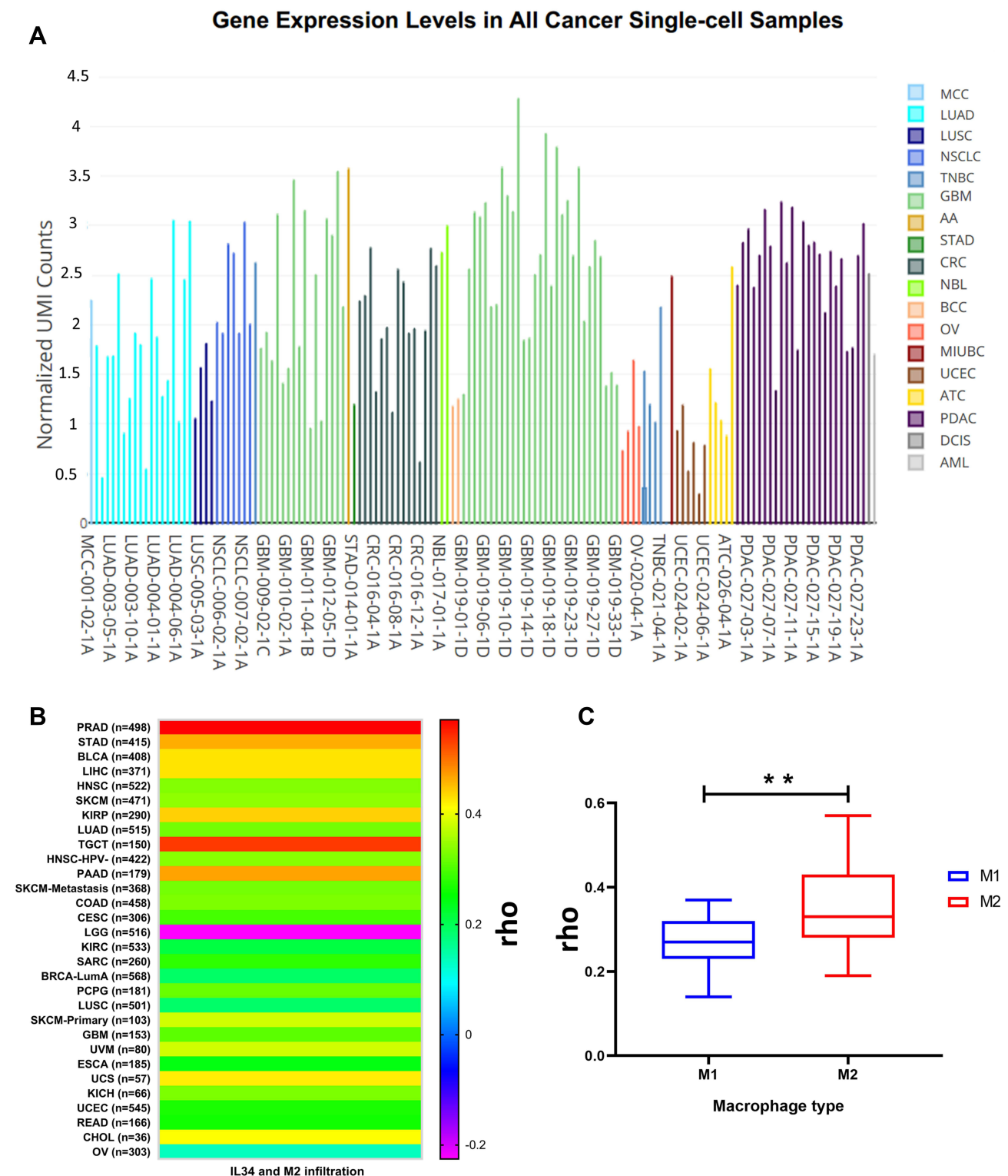


Figure 2 (A) Pan-cancer single-cell expression view among 638,341 cells. **(B)** Pan-cancer analysis for IL-34 expression and M2 associations by Spearman correlation test ($P < 0.05$). **(C)** An Independent test result of matched correlations values analysis (23 for each) between M1 and M2 macrophages. $** (P = 0.002)$.

Abbreviations: MCC, Merkel cell carcinoma; LUAD, Lung adenocarcinoma; LUSC, Lung squamous cancer; NSCLC, non-small cell lung cancer; TNBC, Triple-negative breast cancer; GBM, Glioblastoma; AA, Anaplastic astrocytoma; STAD, Stomach adenocarcinoma; CRC, Colorectal cancer; NBL, Neuroendocrine tumor; BCC, Basal cell carcinoma; OV, Ovarian carcinoma; MIUBC, Muscle-invasive urothelial bladder cancer; UCEC, Uterine cancer- endometrial carcinoma; ATC, Anaplastic thyroid cancer; PDAC, Pancreatic adenocarcinoma; DCIS, Breast ductal carcinoma in situ; AML, Acute myeloid leukemia.

Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis, and interpretation, or in all these areas; took part in drafting, revising, or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted, and agree to be accountable for all aspects of the work.

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Disclosure

The authors report no conflicts of interest in this work.

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