# Decoding cancer's identity: The power of immune profiling.

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# Introduction

The intricate relationship between the immune system and cancer is at the forefront of medical research, reshaping our understanding of disease and treatment. One of the most promising tools in this endeavor is immune profiling—an analytical approach that delves deep into the molecular and cellular intricacies of the immune response within tumors. Immune profiling not only provides insights into the tumor's interactions with the immune system but also holds the key to developing personalized and effective cancer therapies [1].

**Cancers don't exist in isolation**: they are surrounded by a dynamic ecosystem known as the Tumor Microenvironment (TME). This environment includes cancer cells, immune cells, stromal cells, blood vessels, and signaling molecules. Immune profiling offers a detailed snapshot of this complex landscape, shedding light on the interactions between different cell types and their impact on tumor progression.

**Flow cytometry**: This technique allows researchers to analyze individual cells within a tumor sample, identifying specific immune cell populations and their characteristics. Flow cytometry enables the quantification of immune cell types and the determination of their activation status [2].

**Immuno Histochemistry (IHC)**: IHC is a technique used to visualize specific proteins within tissue samples. By staining tumor sections with antibodies that target immune cell markers, researchers can visualize the presence, distribution, and localization of immune cells in the tumor.

**Single-cell RNA sequencing**: This cutting-edge technology enables the study of gene expression in individual cells within a tumor. It provides insights into the diversity of immune cell types, their functional states, and their communication within the TME.

**Mass Cytometry (CyTOF)**: Similar to flow cytometry, CyTOF analyzes multiple markers on individual cells simultaneously, allowing for high-dimensional profiling of immune cells. It offers the advantage of a larger number of markers compared to traditional flow cytometry [3].

Immunoscore: A Revolution in Cancer Prognosis

One breakthrough arising from immune profiling is the Immunoscore, a revolutionary method for predicting cancer outcomes. By quantifying the density of specific immune cell types within tumors, the Immunoscore provides a more accurate prediction of patient survival and treatment response than traditional methods. This information helps guide treatment decisions, enabling clinicians to tailor therapies based on a patient's individual immune profile.

Immune profiling holds immense potential for the development of personalized cancer immunotherapies. By understanding the immune landscape of a tumor, researchers can identify targets for intervention, such as immune checkpoint inhibitors or adoptive T-cell therapies. Additionally, immune profiling can help predict which patients are more likely to respond to specific immunotherapies, ensuring that treatments are both effective and tailored to the individual [4].

As technology advances and our understanding of immune responses deepen, immune profiling will continue to refine our approach to cancer treatment. Integrating immune profiling into clinical practice will empower oncologists to make informed decisions, leading to better patient outcomes and a new era of precision medicine [5].

### Conclusion

Immune profiling represents a quantum leap in our ability to comprehend the intricate dance between cancer and the immune system. By decoding the language of the tumor microenvironment, we unlock the potential to reshape cancer treatment strategies and offer patients the tailored therapies they deserve. As immune profiling continues to guide research and clinical decisions, we edge closer to a future where cancer treatments are as unique as the individuals they aim to heal.

#### References

- Gonzalez Castro LN, Tirosh I, Suvà ML. Decoding cancer biology one cell at a time. Cancer Discov. 2021;11(4):960-70.
- 2. Segal E, Sirlin CB, Ooi C, et al. Decoding global gene expression programs in liver cancer by noninvasive imaging. Nat Biotechnol. 2007;25(6):675-80.
- 3. Anzellotti S, Fairhall SL, Caramazza A. Decoding representations of face identity that are tolerant to rotation. Cerebral cortex. 2014;24(8):1988-95.
- 4. Volik S, Raphael BJ, Huang G, et al. Decoding the finescale structure of a breast cancer genome and transcriptome. Genome Res. 2006;16(3):394-404.
- 5. Snir T, Efroni S. T cell repertoire sequencing as a cancer's liquid biopsy—can we decode what the immune system is coding?. Curr Opin Syst Biol. 2020;24:135-41.

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