Immunometabolism in the tumor microenvironment and neoantigenbased cancer vaccines: A new era in cancer therapy.

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Introduction

Cancer remains one of the most complex diseases, characterized by its ability to evade immune responses and adapt to various metabolic conditions. The tumor microenvironment (TME) plays a crucial role in modulating immune cell function and metabolic pathways, which in turn influence tumor progression. Immunometabolism, the intersection of immune cell function and metabolic processes, has emerged as a vital area of cancer research. Understanding how metabolic reprogramming within the TME affects immune responses can lead to more effective therapeutic strategies [1].

Simultaneously, advancements in cancer immunotherapy have led to the development of personalized treatment approaches such as neoantigen-based cancer vaccines. These vaccines harness the body's immune system to recognize and destroy tumor-specific antigens, offering a promising strategy for long-term cancer control. This article explores the relationship between immunometabolism in the TME and the potential of neoantigen-based vaccines to revolutionize cancer treatment [2].

The TME consists of tumor cells, immune cells, fibroblasts, and endothelial cells, all of which interact dynamically to influence tumor growth. Within this environment, metabolic alterations occur that support tumor survival while suppressing immune responses. For instance, cancer cells shift towards glycolysis, even in oxygen-rich conditions, a phenomenon known as the Warburg effect. This metabolic shift leads to the depletion of nutrients, such as glucose and amino acids, which are essential for immune cell activation [3].

Furthermore, the accumulation of immunosuppressive metabolites, such as lactic acid and kynurenine, impairs T-cell function and promotes the expansion of regulatory T cells (Tregs) and myeloid-derived suppressor cells (MDSCs). These immunosuppressive elements create a hostile environment that limits the effectiveness of anti-tumor immunity, highlighting the need for strategies that can counteract metabolic suppression [4].

Recent research has focused on modulating immunometabolism to enhance anti-cancer immune responses. Targeting metabolic pathways, such as glycolysis inhibitors and amino acid deprivation strategies, can help reprogram immune cells within the TME. Additionally, interventions like immune

checkpoint inhibitors (ICIs) have been shown to improve T-cell metabolism, restoring their ability to attack cancer cells [5].

Another promising approach involves using metabolic adjuvants in combination with immunotherapies. For example, drugs that inhibit lactate production or enhance oxidative phosphorylation in T cells can create a more favorable metabolic landscape, increasing the effectiveness of cancer treatments. These metabolic interventions pave the way for novel immunotherapeutic strategies, including cancer vaccines [6].

Neoantigen-based cancer vaccines represent a groundbreaking approach in precision medicine. Unlike traditional cancer vaccines, which target shared tumor-associated antigens, neoantigens are unique to individual tumors and arise from somatic mutations. This makes them highly immunogenic and less likely to trigger tolerance mechanisms that hinder immune responses [7].

Neoantigen vaccines work by presenting tumor-specific antigens to the immune system, stimulating cytotoxic T lymphocytes (CTLs) to recognize and eliminate cancer cells. Advances in next-generation sequencing (NGS) and bioinformatics have enabled the rapid identification of neoantigens, allowing for the development of personalized vaccines tailored to each patient's tumor profile [8].

Despite the promising potential of neoantigen-based vaccines, several challenges remain. One major hurdle is the tumor's ability to evolve and develop immune evasion mechanisms. Tumors can downregulate antigen presentation or modify their microenvironment to suppress immune responses, reducing vaccine efficacy. Overcoming these challenges requires innovative approaches, such as combination therapies that integrate neoantigen vaccines with ICIs, metabolic modulators, or adoptive T-cell therapies [9].

Additionally, optimizing vaccine delivery platforms, such as lipid nanoparticles or dendritic cell-based approaches, can enhance antigen presentation and improve immune responses. Ongoing clinical trials are investigating various strategies to maximize the effectiveness of neoantigen vaccines, offering hope for more durable and personalized cancer treatments in the future [10].

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Conclusion

The interplay between immunometabolism in the TME and the development of neoantigen-based cancer vaccines represents a promising frontier in oncology. By targeting metabolic pathways that suppress immune responses, researchers can enhance the efficacy of cancer immunotherapies. Neoantigen vaccines, with their personalized approach, offer a new level of precision in cancer treatment, potentially leading to long-lasting immune protection. As scientific advancements continue to unravel the complexities of the TME and neoantigen presentation, the integration of immunometabolic interventions with personalized cancer vaccines holds the potential to redefine cancer treatment. This innovative approach not only improves patient outcomes but also paves the way for a more effective and targeted battle against cancer.

References

- Khoiri S, Moussango VD. A Short Review on Harnessing Bioinformatics for Food Safety: Computational Approaches to Detecting Foodborne Pathogens. J Adv Healt Inform Res. 2024;2(3):109-14.
- 2. Taiwo OR, Onyeaka H, Oladipo EK, et al. Advancements in Predictive Microbiology: Integrating New Technologies for Efficient Food Safety Models. Inter J Microb. 2024;2024(1):6612162.
- 3. Taiwo OR, Onyeaka H, Oladipo EK, et al. Advancements in Predictive Microbiology: Integrating New Technologies for Efficient Food Safety Models. Intern J Microb. 2024;2024(1):6612162.

- 4. Pattabhiramaiah M, Mallikarjunaiah S. High-Throughput Sequencing for Detection of Foodborne Pathogens in Food Safety. Seq Tech Micro Food Saf Qual. CRC Press.
- Emamjomeh M, Hashim AM, Abdul-Mutalib NA, Mokhtar NF, et al. Profiling bacterial communities and foodborne pathogens on food-associated surface following contact with raw beef, chicken and pork using 16S amplicon metagenomics. Food Control. 2023;149:109698.
- 6. Mather AE, Gilmour MW, Reid SW, et al. Foodborne bacterial pathogens: genome-based approaches for enduring and emerging threats in a complex and changing world. Nat Rev Microb. 2024;22(9):543-55.
- 7. Li S, Tian Y, Jiang P, et al. Recent advances in the application of metabolomics for food safety control and food quality analyses. Crit Rev Food Sci Nutr. 2021;61(9):1448-69.
- 8. Beck KL, Haiminen N, Chambliss D, et al. Monitoring the microbiome for food safety and quality using deep shotgun sequencing. Sci Food. 2021;5(1):3.
- 9. Quek JJ, Wong JL, Tan JL, et al. Integrating Metagenomic and Culture-Based Techniques to Detect Foodborne Pathogens and Antimicrobial Resistance Genes in Malaysian Produce. Foods. 2025 Jan 22;14(3):352.
- 10. Vieira KC, Silva HR, Rocha IP, et al. Foodborne pathogens in the omics era. Crit Revi Food Sci ad Nutr. 2022;62(24):6726-41.