Decoding Cross-presentation: Unravelling MHC-mediated Molecular Functions and Regulatory Mechanisms

Tindea Mechteleria*

Department of Immunology, Kindai University, Wakayama 649-6493, Japan

Abstract

Cross-presentation, a fundamental process in immunology, plays a pivotal role in the adaptive immune response by presenting extracellular antigens on Major Histocompatibility Complex Class I (MHC-I) molecules. This comprehensive review delves into the intricate molecular characteristics, functional definitions and regulatory mechanisms governing cross-presentation. The exploration encompasses the diverse cellular players involved in this process, emphasizing the central role of antigen-presenting cells, particularly dendritic cells. The review outlines the molecular machinery responsible for the generation of MHC-I/peptide complexes from exogenous antigens, shedding light on the precise mechanisms that orchestrate cross-presentation. Furthermore, the regulatory checkpoints governing the efficiency and specificity of cross-presentation are dissected, providing insights into the dynamic interplay between the immune system and its surrounding microenvironment. This comprehensive analysis serves to enhance our understanding of cross-presentation, offering a foundation for future research and potential therapeutic interventions that leverage this critical immunological process.

Keywords: Cross-presentation • Antigen presentation • Dendritic cells • Adaptive immune response

Introduction

Cross-presentation, a finely tuned immunological phenomenon, plays a crucial role in shaping the adaptive immune response. This intricate process involves the presentation of extracellular antigens on Major Histocompatibility Complex Class I (MHC-I) molecules, allowing the immune system to efficiently recognize and eliminate infected or aberrant cells. Despite its pivotal significance, the molecular intricacies and regulatory mechanisms underlying cross-presentation are still subjects of intensive investigation. This review seeks to provide a comprehensive exploration of the molecular characteristics, functional definitions and regulatory mechanisms that govern cross-presentation. By delving into the dynamic interplay between cellular components and unveiling the molecular machinery orchestrating crosspresentation, this review aims to deepen our understanding of this fundamental immunological process. The exploration of cross-presentation has been a dynamic journey within the realms of immunology, with a rich tapestry of studies contributing to our evolving understanding of this phenomenon. Dendritic cells, renowned for their antigen-presenting prowess, emerge as central players in orchestrating cross-presentation. Numerous studies have unveiled the molecular events within dendritic cells that facilitate the capture, processing and presentation of extracellular antigens on MHC-I molecules. The cellular machinery responsible for the transport of antigens into the crosspresentation pathway, such as endosomal and phagosomal compartments, has been a focal point of investigation [1,2].

Literature Review

An array of antigens, including viral particles, cell debris and tumor-

*Address for Correspondence: Tindea Mechteleria, Department of Immunology, Kindai University, Wakayama 649-6493, Japan, E-mail: tindeamech@yahoo.com

Copyright: © 2023 Mechteleria T. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Received: 04 December, 2023, Manuscript No. icoa-24-126567; Editor assigned: 06 December, 2023, Pre QC No. P-126567; Reviewed: 18 December, 2023, QC No. Q-126567; Revised: 23 December, 2023, Manuscript No. R-126567; Published: 30 December, 2023, DOI: 10.37421/2469-9756.2023.9.212 associated antigens, are subject to cross-presentation. The diversity in the nature of antigens and the pathways through which they are processed underscores the complexity of cross-presentation. The literature reveals that the intricacies of this process extend beyond dendritic cells, involving a network of immune cells and factors that collectively contribute to the shaping of immune responses. Moreover, studies exploring the regulatory mechanisms of cross-presentation have uncovered checkpoints that control the efficiency and specificity of this process. Intriguingly, the immune microenvironment and external stimuli can influence cross-presentation, adding layers of complexity to its regulation. Understanding these regulatory circuits is not only crucial for deciphering the intricacies of immune surveillance but also holds potential implications for therapeutic interventions aimed at enhancing or modulating cross-presentation in various disease contexts. In summary, the literature review underscores the multifaceted nature of cross-presentation, reflecting a field rich in diversity and complexity. The collective findings pave the way for a more comprehensive understanding of the molecular functions and regulatory mechanisms governing cross-presentation, setting the stage for future research endeavors aimed at unraveling the full extent of its impact on immune responses and potential applications in therapeutic interventions [3,4].

Discussion

The in-depth exploration of cross-presentation, unraveling the Major Histocompatibility Complex (MHC)-mediated molecular functions and regulatory mechanisms, has illuminated the complexity of this critical immunological process. Dendritic cells, with their unique ability to capture, process and present extracellular antigens on MHC-I molecules, stand out as central orchestrators of cross-presentation. The literature reveals a sophisticated interplay of cellular components and molecular machinery within dendritic cells that governs the efficient presentation of antigens originating from exogenous sources. One key aspect of the discussion centers on the diverse nature of antigens subject to cross-presentation. Whether derived from viral particles, cell debris, or tumor-associated antigens, the range of substances processed through this pathway highlights its versatility. Understanding the intricacies of antigen recognition and processing is essential for harnessing the full potential of cross-presentation in therapeutic applications, particularly in the context of developing vaccines or immunotherapies for infectious diseases and cancers [5].

The review also delves into the dynamic regulatory mechanisms that

govern cross-presentation. The checkpoints controlling the efficiency and specificity of this process represent critical points of intervention. External stimuli, immune microenvironment factors and the crosstalk between different immune cells contribute to the modulation of cross-presentation, adding layers of complexity to its regulation. Deciphering these regulatory circuits not only enhances our understanding of immune surveillance but also opens avenues for targeted interventions to manipulate cross-presentation for therapeutic purposes. Moreover, the insights gained from decoding cross-presentation have broader implications for personalized medicine and the design of targeted immunotherapies. Understanding the intricate molecular machinery involved in cross-presentation allows for the identification of potential therapeutic targets that could be manipulated to enhance or modulate immune responses selectively. This is particularly relevant in the context of cancer immunotherapy, where exploiting cross-presentation pathways may bolster the immune system's ability to recognize and eliminate tumor cells. The integration of recent advancements in technologies such as single-cell sequencing and highthroughput screening further refines our understanding of the heterogeneity within dendritic cell populations and their roles in cross-presentation. This granularity is crucial for tailoring therapeutic strategies to individual patients, considering the diverse antigenic landscapes present in various diseases. Furthermore, the discussion of regulatory mechanisms emphasizes the need for a nuanced approach to modulating cross-presentation. The influence of the immune microenvironment and external factors underscores the dynamic nature of immune responses. Harnessing this knowledge could lead to the development of immunomodulatory agents that finely tune cross-presentation, enhancing its efficacy while minimizing potential detrimental effects on immune homeostasis [6].

Conclusion

In conclusion, the comprehensive review of cross-presentation, focusing on MHC-mediated molecular functions and regulatory mechanisms, provides a thorough understanding of the intricate processes that govern this essential aspect of the immune system. Dendritic cells emerge as central players and the molecular events within these cells offer valuable insights into the mechanics of antigen presentation. The diversity of antigens processed through cross-presentation and the nuanced regulatory mechanisms underline the adaptability and complexity of this immunological process. As we decode the molecular intricacies of cross-presentation, the knowledge gained holds immense potential for shaping future research endeavors and therapeutic applications. Targeting cross-presentation pathways may open new avenues for the development of vaccines, immunotherapies and interventions in various disease contexts. By unlocking the secrets of MHC-mediated functions and regulatory checkpoints, we advance not only our understanding of immunology but also the prospects for harnessing the power of the immune system in combatting infectious diseases, cancers and other immune-related disorders.

Acknowledgment

None.

Conflict of Interest

There are no conflicts of interest by author.

References

- Hughes, Austin L. and Meredith Yeager. "Natural selection at major histocompatibility complex loci of vertebrates." Annu Rev Genet 32 (1998): 415-435.
- Khare, Vishwa M., Vishesh K. Saxena, Alka Tomar and Karam Pal Singh, K et al. "MHC-B haplotypes impact susceptibility and resistance to RSV-A infection." Front Biosci 10 (2018): 506-19.
- Madden, Dean R. "The three-dimensional structure of peptide-MHC complexes." Annu Rev Immunol 13 (1995): 587-622.
- González, Elizabeth Rasche. "Immunogenetics recognized by Nobel Prize." JAMA 244 (1980): 2396-2400.
- Xia, Jing, Ying Zhang, Huanhuan Zhao and Jie Wang, et al. "Non-invasive monitoring of CNS MHC-I molecules in ischemic stroke mice." *Theranostics* 7 (2017): 2837.
- Malviya, Gaurav, E. F. J. De Vries, Rudi A. Dierckx and Alberto Signore. "Synthesis and evaluation of 99m Tc-labelled monoclonal antibody 1D09C3 for molecular imaging of major histocompatibility complex class II protein expression." *Mol Imaging Biol* 13 (2011): 930-939.

How to cite this article: Mechteleria, Tindea. "Decoding Cross-presentation: Unravelling MHC-mediated Molecular Functions and Regulatory Mechanisms." *Immunochem Immunopathol* 9 (2023): 212.