

TARGETING M2-TUMOR ASSOCIATED MACROPHAGES IN PROSTATE CANCER USING SURFACE ENRICHED GLYCOPROTEINS

Jelani C. Zarif¹, James R. Henandez¹, Weiming Yang², Hui Zhang², and Kenneth J. Pienta^{1, 3,4,5}

¹The Brady Urological Institute at the Johns Hopkins University School of Medicine Baltimore, MD 21287, ²Department of Pathology, ³Department Medical Oncology, Johns Hopkins School of Medicine and Sidney Kimmel Comprehensive Cancer Center, Baltimore, MD, ⁴Department of Pharmacology and Molecular Sciences, Johns Hopkins School of Medicine, Baltimore, MD, ⁵Department of Chemical and Biomolecular Engineering; Johns Hopkins University Baltimore, MD USA

Prostate cancer is a leading cause of cancer-related deaths of men in the U.S., and in the past year, over 30,000 men died from this disease. While localized prostate cancer is highly treatable by surgical resection and radiation, cancer that has metastasized remains incurable. Alternatively activated macrophages, also known as M2-macrophages, primarily scavenge debris and in the process, promote angiogenesis and wound repair. M2-macrophages are phenotypically similar to M2 tumor-associated macrophages (M2-TAMs) have been reported to associate with solid tumors such as prostate cancer to facilitate epithelial to mesenchymal transition (EMT), tumor invasiveness, metastasis, and resistance to therapy. As an invasive species within the tumor microenvironment, this makes M2-TAMs an ideal therapeutic target in prostate cancer. The purpose of this project is to develop novel therapeutics that will directly target M2-TAMs for destruction and subsequently attenuate prostate tumor growth, progression, and metastasis. Our hypothesis is to determine if targeting of M2-TAMs by using enriched surface antigens that are targeted by antibody-drug-conjugates (ADCs), be an effective therapy for lethal prostate cancer while simultaneously eliciting an immune response. To identify novel surface antigens expressed on M2-macrophages, we developed a novel method of creating homogenous populations of human macrophages from CD14⁺ monocytes *in vitro*. Our homogenous M1 macrophages secrete pro-inflammatory cytokines and our M2 macrophages secrete anti-inflammatory cytokines as well as VEGF. We then performed solid-phase extraction of *N*-linked glycopeptides (SPEG) followed by liquid chromatography-tandem Mass Spectrometry (LC-MS/MS) on CD14⁺ monocytes and homogenous macrophage populations. We discovered five novel peptides that are enriched exclusively on M2-macrophages relative to M1 macrophages and CD14⁺ monocytes. Lastly, we determined if these surface antigens, found enriched on M2 macrophages, were also expressed in human metastatic castrate-resistant prostate cancer (mCRPC). Using mCRPC tissues from rapid autopsies supplied by the Departments of Urology and Surgical Pathology, we were able to determine M2-macrophage infiltration by using immunohistochemistry and flow cytometry. The studies described here outline a method of altering the tumor immune microenvironment. To target M2 macrophages, we used small peptides that targeted an enriched surface glycoprotein CD206 that is expressed on M2-macrophages. We will then test their efficacies *in vivo* using a syngeneic prostate tumor mouse model to assess tumor shrinkage and effector T Cell infiltration. By identifying specific markers on M2-TAMs, we predict that this method of targeting will provide a better prognosis for patients who have been diagnosed with lethal prostate cancer.

Conflict of Interest: The authors declare no conflicts of interest.

Funding Acknowledgements: Research for this study was supported by National Cancer Institute grants CA163124, CA093900, CA143055, the UNCF/Merck Postdoctoral Science Research Fellowship award (J.C.Z.) and the Prostate Cancer Foundation's Young Investigator Award (J.C.Z.).