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MUC4 Based Immunotherapy for Pancreatic Cancer

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MUC4 BASED IMMUNOTHERAPY FOR PANCREATIC CANCER

By

KASTURI BANERJEE

A DISSERTATION

Presented to the Faculty of
The University of Nebraska Graduate College
In Partial Fulfillment of the Requirements
For the Degree of Doctor of Philosophy

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Graduate Program

Under the Supervision of Associate Professor Dr. Maneesh Jain

University of Nebraska Medical Center
Omaha, Nebraska

April 2018

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DEDICATION

I dedicate my PhD dissertation to my father Mr. Chandan Banerjee, my mother

Mrs. Sarbasree Banerjee, my younger sister Ms. Sanchari Banerjee

And my late grandmother Mrs. Kalyani Banerjee

For providing me constant love and support.

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ABSTRACT

MUC4 based immunotherapy for pancreatic cancer

Kasturi Banerjee, Ph.D.

University of Nebraska Medical Center, 2018

Supervisor: Maneesh Jain, PhD.

Pancreatic Cancer (PC) is a lethal disease claiming approximately 45000 lives in the US in 2018, and it establishes an elaborate immunosuppressive tumor microenvironment that aids in disease pathogenesis. Immunotherapy has emerged as a strategy to target tumor cells by reprogramming patient's immune system. Challenges present in PC immunotherapy are: i) identifying a tumor-associated antigen that could be targeted, ii) identifying adjuvants that could efficiently deliver antigens, iii) eliciting robust anti-tumor responses and iv) overcoming peripheral tolerance and immunosuppression elicited by the tumor.

Firstly, we detected circulating autoantibodies to MUC4 present in PC patients and observed that IgM autoantibodies to MUC4 peptides significantly correlate with overall PC patient survival, thus suggesting that MUC4 could potentially be targeted for PC immunotherapy. Our group is the first to successfully purify recombinant MUC4 β protein and characterize its immunogenic activity. We addressed the challenge of protein delivery by encapsulating MUC4 β in novel polyanhydride nanoparticles (MUC4 nanovaccine). In the second part of the dissertation, we characterized MUC4 nanovaccine in both *in vitro* and *in vivo* system. Our studies showed that MUC4 nanovaccine could robustly activate dendritic cells (DCs) and induce secretion of Th1 cytokines *in vitro*. High levels of Th1 IgG2b anti-MUC4 β antibodies were detected in MUC4 nanovaccine-immunized mice.

As described in the third part of the dissertation, we observed that *ex vivo* T-cells activated by MUC4 nanovaccine-pulsed DCs showed enhanced cytotoxic killing of miniMUC4 tumor cells, when compared to soluble MUC4 β mixed with empty nanoparticles (MUC4+NP). We validated our data in an *in vivo* subcutaneous PC tumor mouse model, and observed enhanced immune cells infiltration and corresponding levels of necrosis in miniMUC4 tumors corroborated with low tumor volume of miniMUC4 tumor (in comparison to contralateral vector control tumor) in MUC4-immunized mice. Furthermore, the presence of PD-L1 surface expression on miniMUC4 tumor cells indicated active immunosuppression lodged by tumor cells in response to IFN γ -secreting infiltrating cytotoxic T-cells.

Taken together, studies in this dissertation demonstrate that MUC4 nanovaccine could serve as a potential strategy for PC immunotherapy.

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LIST OF ABBREVIATIONS

ACT: Adoptive T-cell

ADCC: Antibody-Dependent Cellular Cytotoxicity

AMOP: Adhesion-associated Domain in MUC4 and Other Proteins

ANOVA: Analysis of Variance

APCs: Antigen presenting cells

BSA: Bovine Serum Albumin

CAR: Chimeric Antigen Receptors

CDNs: Cyclic Dinucleotides

CEA: Carcinoembryonic Antigen

ChARs: Chromatin-Accessible-Regions

CNP: Cationic Nanoparticles

CP: Chronic Pancreatitis

CPH: 1, 6-bis (*p*-carboxy phenoxy) hexane

CPTEG: 1, 8-bis (*p*-carboxy phenoxy)-3, 6-dioxaoctane

CR: Complete Response

CRS: Cytokine Release Syndrome

CTLA-4: Cytotoxic T-lymphocyte-associated protein 4

CTLs: Cytotoxic T-cells

DCs: Dendritic Cells

ECM: Extracellular Matrix

EGF: Epidermal Growth Factor

EGFR: Epidermal Growth Factor Receptor

ELISA: Enzyme-Linked Immunosorbent Assay

EMT: Epithelial to Mesenchymal

FAK: Focal Adhesion Kinase

FAP⁺: Fibroblast Activation Protein-positive

FBS: Fetal Bovine Serum

FGFR: Fibroblast Growth Factor Receptor

FNA: Fine Needle Aspirates

GAL-1: Galectin-1

Gal3: Galectin-3

GDPH: Glycine-Aspartate-Proline-Histidine

GM-CSF/neu: Granulocyte-Macrophage Colony-Stimulating Factor/neu

HER1: Human Epidermal Growth Factor Receptor 1

HER2: Human Epidermal Growth Factor Receptor 2

HLA: Human Leukocyte Antigen

HRP: Horseradish-peroxidase

IACUC: Institutional Animal Care and Use Committee

IDO: Indoleamine 2, 3-dioxygenase

IEDB: Immune Epitope Database and Analysis Resource

IF: Immunofluorescence

IFNs: Interferons

IFN β : Interferon-beta

IFN γ : Interferon-gamma

IHC: Immunohistochemistry

IL: Interleukin

IRB: Institutional Review Board

JAK: Janus Kinases

JNK: c-Jun N-terminal kinases

KIF20A: Kinesin-like Protein 20A

KPC: K-ras^{LSL.G12D/+}; p53^{R172H/+}; Pdx-1Cre mice

LAG3: Lymphocyte Activation Gene 3

LAK: Lymphokine-activated Killer

LM: *Listeria monocytogenes*

LPS: Lipopolysaccharides

mAbs: Monoclonal Antibodies

MALP-2: Macrophage-Activating Lipopeptide-2

MAPK: Mitogen-Activated Protein Kinases

MDSCs: Myeloid-Derived Suppressor Cells

MHC-I: Major Histocompatibility Complex I

MHCII- Major Histocompatibility Complex II

MMAE: Monomethyl Auristatin E

MMPs: Matrix-Metallopeptidase

MSD: Mean Survival Days

MSLN: Mesothelin

MST: Mean Survival Time

MUC: Mucin

NF- κ B: Nuclear Factor Kappa-light-chain-enhancer of Activated B Cells

NIDO: Nidogen-like Domain

NK cells: Natural Killer Cells

NR: Not Reported

NRI: Nebraska Research Initiative

OVA - Ovalbumin

PanIn: Pancreatic Intraepithelial Neoplasm

PBMC: Peripheral Blood Mononuclear Cells

PBS: Phosphate Buffered Saline

PC: Pancreatic Cancer

PD-1: Programmed Cell Death Protein 1

PD-L1: Programmed Death Ligand 1

PEG: Polyethylene Glycol

PI3K: Phosphatidylinositide 3-Kinases

PKA: Protein Kinase A

PMA: Phorbol 12-myristate 13-acetate

PR: Partial Response

PSCA: Prostate Stem Cell Antigen

PSCs: Pancreatic Stellate Cells

RA: Retinoic Acid

RIT: RetroNectinVR (CH296)-induced T-cell

RT: Room Temperature

RTK: Receptor Tyrosine Kinase

STAT-1: Signal transducer and activator of transcription 1

STING: Stimulator of interferon genes

TAA: Tumor Associated Antigens

TAM: Tumor Associated Macrophages

TBS: Tris-buffered Saline

TCGA: The Cancer Genome Atlas

TCR: T-cell Receptor

TGF β : Tumor Growth Factor-beta

TLRs: Toll-like Receptors

TMB: 3,3',5,5'-Tetramethylbenzidine

TME: Tumor Microenvironment

TNF: Tumor Necrosis Factor

TNFR-1: TNF Receptor 1

TNF- α : Tumor Necrosis Factor Alpha

UNMC: University of Nebraska Medical Center

UPMC: University of Pittsburgh Medical Center

VEGF: Vascular Endothelial Growth Factor

VEGFR1/2: VEGF Receptor1/2

VNTR: Variable Number Tandem Repeat

vWD: von Willebrand factor type D domain

WT1: Wilm's Tumor protein 1

CHAPTER 1A: INTRODUCTION

The material covered in this chapter is the subject of one review article

1. **Banerjee K.**, Kumar S., Ross K. A., Gautam S., Poelaert B., Nasser M. W., Aithal A., Bhatia R., Wannemuehler M.J., Narasimhan B., Solheim J. C., Batra S. K. and Jain M., *Emerging trends in the immunotherapy of pancreatic cancer*, **Cancer Letters**, **417** (2018) 35-46

1. Synopsis

Pancreatic cancer (PC) is the fourth leading cause of cancer-related deaths in the U.S., claiming approximately 45,000 lives every year. Much like other solid tumors, PC evades the host immune system and establishes itself by manipulating the immune system to establish an immunosuppressive tumor microenvironment (TME). Therefore, targeting and reinstating patient's immune system could serve as a powerful therapeutic tool. Indeed, immunotherapy has emerged in recent years as a potential adjunct treatment for solid tumors including PC. Immunotherapy modulates the host's immune response to tumor-associated antigens (TAAs), eradicates cancer cells by reducing host tolerance against TAAs and provides both short- and long-term protection against the disease. Passive immunotherapy like monoclonal antibodies or adoptive T-cell based therapy focuses on direct targeting of tumor cells, whereas, active immunotherapy like cancer vaccines specifically activates the patient's immune cells against cancer cells. Such strategies have been tested for antitumor responses alone and in combination with standard care in multiple preclinical and clinical studies. In this review, we discuss various immunotherapy strategies used currently and their efficacy in abrogating self-antigen tolerance and immunosuppression, as well as their ability to eradicate PC.

2. Introduction to Immunotherapy

It's forecasted that by 2030, pancreatic cancer (PC) will become the second leading cause of cancer-related deaths in the United States [1]. The current overall survival of PC patients by stage of disease progression is no more than 14% at Stage I, and the overall 5-year survival is approximately 8% [2]. The success rate of various treatment modalities for PC including surgery, chemotherapy, and radiation is limited and reoccurrence is typically inevitable [3]. In addition, late diagnosis of the disease further compounds the problem leading to high mortality rate. Recently immunotherapy has revolutionized cancer treatment especially in melanoma [4, 5]. It is increasingly being felt that immunotherapy if applied properly in combination with the standard of care can lead to better outcomes in solid tumors including PC. Studies support immunotherapy as a viable and metamorphic approach, which can boost and restore the immune system's ability to fight against cancer.

In this review, we describe the current understanding of different immunotherapeutic approaches including anti-cancer monoclonal antibodies (mAbs), T-cell-mediated immunotherapies and cancer vaccines, as powerful strategies for PC treatment. We will also discuss the clinical efficacy of immunotherapeutic strategies, challenges and assess their feasibility as next-generation treatment options, either alone or in combination with chemotherapy for PC treatment.

3. Immunoediting and Immunosuppressive Microenvironment

The interaction of the immune system with cancer cells is comprised of the three phases: Elimination, Equilibrium and Escape [6-8]. During the Elimination phase, the immune system can recognize and eliminate transformed cells. PC cells shed MICA that impairs cell surface expression of NKG2D receptor on T-cell and Natural Killer (NK) cells,

thus, helping tumor cells to escape immune surveillance by inhibiting the cytolytic activity of T-cell and NK cells [9, 10]. Transformed cells that escape the Elimination phase enter the Equilibrium phase, in which cancer cells undergo genomic editing and establish the tumor microenvironment (TME) that supports the growth of the early lesions. Finally, in the Escape phase, PC recruit immunosuppressive cells like myeloid-derived-suppressor cells (MDSCs), regulatory T-cells (T_{reg} cells) and tumor-associated macrophages (TAMs) [11, 12]. These immature MDSCs induce apoptosis in activated T-cells. Depletion of MDSCs in an autochthonous PC mouse model results in the unmasking of adaptive immune responses, tumor cell death and remodeling of tumor stroma [13, 14]. PC conditioned media-treated $CD4^+$ T-cells favored expansion of $CD69^+$ activated T_{reg} subset, which is known to suppress $CD4^+$ T-cell proliferation, thus promoting immunosuppression [15].

PC tumor and stromal cells secrete angiogenic chemokines like vascular endothelial growth factor (VEGF), signaling molecules like galectin-1 (GAL-1) [16], anti-inflammatory cytokines like Interleukin (IL)-10, IL-13 [17], and immune-inhibitory ligands like programmed death ligand 1 (PD-L1), cytotoxic T-lymphocyte-associated protein 4 (CTLA-4), which further facilitate escape from cytotoxic cell-mediated death by cluster of differentiation (CD) 8^+ T-cells and NK cells [13, 14, 18-22] (**Figure 1**). Additionally, exhausted $CD8^+$ T-cells have chromatin-accessible-regions (ChARs) that serve as an enhancer to maintain high levels of programmed cell death protein 1 (PD-1), which further keeps $CD8^+$ T-cells in an immunosuppressive state [23]. After escaping the immunosurveillance, immunologically and genetically modified tumor cells proliferate, which further prevent immunological [6, 8, 24]. Overall, cancer cells modify itself genetically and exploit the immunosuppressive behavior of the immune system to facilitate its escape from cell death mediated by $CD8^+$ T lymphocytes and NK cells.

4. PC Microenvironment and Immune Suppression

PC cells harbor limited genetic alterations and simultaneously modulate the TME to escape antitumor immune response. Several studies suggest that immunotherapeutic approaches are more effective in cancers with high mutation profile, displaying varied neoantigens, and having permeable tumor matrix architecture allowing better effector CD8⁺ T-cell infiltration [25, 26]. Furthermore, PC is an immunologically 'cold' tumor due to its low mutational load, dense desmoplasia and rigid extracellular matrix architecture, which restricts the access of effector immune cells to tumor islands, a phenomenon known as excluded infiltrate TME [27-31].

Activated PC stromal compartment excludes CD8⁺ T-cells, CD20⁺ B-cells, and CD56⁺ NK cells from the juxtatumoral region of the tumor. Administration of all-trans retinoic acid in a spontaneous K-ras^{LSL.G12D/+}; p53^{R172H/+}; Pdx-1Cre mice (KPC) mouse model induced quiescence in stromal cells, reduced fibronectin expression and correspondingly increased high influx of CD8⁺ T-cells into juxtatumoral compartment [32]. In PC tumor-bearing transgenic mice, ~55% of fibroblast activation protein-positive (FAP⁺) stromal cells were depleted by selective expression of the diphtheria toxin receptor, showed slow tumor growth only in the presence of CD4⁺ and CD8⁺ T-cells in the TME. Administration of anti PD1 and anti CTLA-4 antibodies along with FAP⁺ stromal cell deletion further reduced tumor growth in these mice [33]. Jiang et al. targeted focal adhesion kinase (FAK) using a selective FAK inhibitor VS-4718 and observed reduced tumor fibrosis and significantly increased survival of KPC mice, compared to untreated control mice. The authors demonstrated that FAK was hyperactive in neoplastic PC tumor and contributed to immunosuppressive TME by restricting cytotoxic CD8⁺ T-cells infiltration. Inhibition of FAK made TME penetrable, which increased adoptive cytotoxic anti-ovalbumin (OVA) CD8⁺ T-cells infiltration in the tumor and the efficacy of PD-1

antagonists in KPC mice when combined with low dose gemcitabine (25mg/kg) compared to combined treatment with vehicle, gemcitabine and anti PD-1 antibody [34]. These studies suggest that the presence of stroma in PC adds an obstacle to current immunotherapy strategies.

Modulation of stromal cells and their effects are influenced by galectins, which are soluble immune-modulating glycoproteins involved in T-cell homeostasis. For example, GAL-1 is expressed primarily by pancreatic stellate cells (PSCs), and GAL-1 overexpressing PSCs have been shown to induce apoptosis in co-cultured CD4⁺ and CD8⁺ T-cells compared to normal or quiescent PSCs [35, 36]. Another member of the galectin family, Galectin3 (Gal3), has also been demonstrated to play an important role in CD8⁺ T-cell biology by inducing the apoptosis in activated T-cells [37, 38]. Jaffee et al. found that neu-specific CD8⁺ T-cells have a high surface expression of lymphocyte activation gene 3 (LAG3), and Gal3 interaction with LAG3 induces immunosuppressive signaling in T-cells through the Gal3-LAG3 axis. This interaction reduces Interferon γ (IFN γ) production by CD8⁺ T-cells and abrogates their infiltration into the TME [39]. Immunization with GM-CSF/neu (granulocyte-macrophage colony-stimulating factor/neu) vaccine has higher efficacy in Gal3 knockout tumor-bearing mice when compared to wild-type tumor-bearing mice. After 60 days of treatment, approximately 80% of Gal-3-knockout mice were disease free, compared to 20% of wild-type mice. Neu-specific CD8⁺ T-cells produced high amounts of interferon-gamma (IFN γ) and granzyme B. In addition, Gal3 knockout mice had increased infiltration of plasmacytoid dendritic cells (DCs), which have a higher potential to activate CD8⁺ T-cells than conventional DCs. Thus, galectins play an important role not only in modulating T-cell function but also in the recruitment of immunosuppressive myeloid-derived suppressor cells (MDSCs) [39].

Apart from the establishment of stroma, tumor cells also modulate the cellular composition of their microenvironment. Tumor-associated macrophages (TAMs), which are M2 (pro-tumor) differentiated macrophages, express tolerance-inducing ligands such as PD-L1/L2, promote angiogenesis, and suppress adaptive immune responses through matrix remodeling carried out by matrix-metalloproteinases (MMPs) [40]. In addition, immunosuppressive CD4⁺FOXP3⁺ T_{reg} cells heavily infiltrate human PC tumors. In the KPC mouse model T_{reg} infiltration increases during the progression from pancreatic intraepithelial neoplasm (PanIN) to the advanced PC stage [13]. Similarly, increased numbers of CD68⁺ TAMs and MDSCs in circulation and TME are associated with disease invasiveness in PC patients [41, 42]. MDSCs also produce reactive oxygen species that further inhibit the antigen-specific response of CD8⁺ T-cells in TME [43].

PC tumor cells secrete anti-inflammatory cytokines and angiogenic chemokines that promote an immunosuppressive TME while facilitating metastasis [3, 32, 41, 44]. These cytokines activate T_{reg} and Th2 cells that prohibit anti-tumor responses elicited by other immune cells [24, 45-47]. Likewise, Indoleamine 2, 3-dioxygenase (IDO) enzyme is secreted by PC tumor cells that upregulate the growth of tumor cells by catabolizing tryptophan into kynurenine, and in turn inhibits T-cell and NK cells activation as well as induces T_{reg} differentiation leading to immunosuppressive TME [3].

Cytotoxic T-cells (CTLs), helper T-cells and DCs are functionally impaired in the immunosuppressive TME of PC, thus skewing to Th2 (tumor-tolerating) responses. Therefore, a proper understanding of these intricate cancer-immune system interactions is very essential to develop and monitor efficacious immunotherapies. The primary goal of PC immunotherapies is to target these interactions and the reprogramming of the immune system against PC tumor microenvironment

5. Immunotherapy Based Approaches

The goal of immunotherapy is to induce antitumor responses by reprogramming and augmenting immune surveillance and removing immune suppression. These anti-cancer immunotherapeutic approaches are divided into 'passive' and 'active' immunotherapies. Passive immunotherapeutic strategies involve mAbs, adoptive T-cell transfers and genetically engineered T-cells. Whereas the active immunotherapeutic approaches include vaccine-mediated immunity induced by the administration of tumor-associated antigens (TAAs) [48]. The TAAs could be delivered in the form of DNA or peptide vaccines, as well as modified tumor cells or antigen-pulsed DCs. Due to genetic alterations or post-translational modifications of proteins (such as glycosylation, phosphorylation, etc.), tumor cells can express proteins that differ from their counterpart in the normal cells or are aberrantly overexpressed in tumor tissues [49].

5.1 Passive Immunotherapeutic Strategies

Passive immunotherapy attacks cancer by directly targeting TAAs by the administration of diverse immune components that are engineered *ex vivo*. Following are the major passive immunotherapeutic strategies ongoing in preclinical studies or clinical trials and have been summarized in **Table II**.

5.1.1 Antibody-Mediated Passive Immunotherapy

Antibody-mediated immunotherapy involves targeting tumors using monoclonal antibodies (mAbs), antibody fragments, antibody-drug conjugates, or radio-immunotherapy conjugates to inhibit tumor signaling, immune suppression, or immune checkpoint blockade.

Anti CD40 mAbs

CD40 is a member of the tumor necrosis factor (TNF) receptor superfamily and is expressed primarily on APCs such as DCs, macrophages, monocytes, B-cells and some non-immune cells like cancer cells [50]. Anti-CD40 antibodies mimic the co-stimulatory signal of the CD40 ligand (CD40L). Tumor-bearing KPC mice with constitutive K-ras activation and gain-of-function p53 mutation when treated with anti-CD40 (clone FGK45, endotoxin-free), either alone or in combination with gemcitabine, showed detectable tumor regression. Treatment with anti-CD40 mAb bypassed the requirement for Toll-Like Receptors (TLRs), inflammasome, Type I interferons (IFNs), and stimulator of interferon genes (STING) to effectively prime adaptive T-cell responses against PC in these animals [51]. The mechanistic role of agonistic anti-CD40 mAb is to activate host antigen presenting cells (APCs) especially DCs, and also to induce clinically relevant antitumor T-cell responses, reverse tumor-induced immune suppression and induce T-cell-independent but macrophage-dependent tumor regression in PC patients [52]. In a clinical trial, 22 naïve patients with advanced PC were administered weekly doses of anti-CD40 mAb in combination with gemcitabine, which led to the increased B-cell surface expression of co-stimulatory molecules CD86, HLA-DR, and CD54 at 24-48 h post-treatment [52-55]. In another study on 21 chemotherapy-naïve and surgically incurable PC patients, treatment with gemcitabine and a human agonist anti-CD40 mAb (CP-870,893) for three weekly cycles showed the enhanced overall survival of 7.4 months compared to those who received gemcitabine alone with the median overall survival of 5.7 months. Upon biopsy, the tumors of anti-CD40 mAb-treated patients showed higher infiltration of macrophages, however, with accompanying absence of lymphocytes [55, 56].

Anti PD-L1mAbs

PD-1 (CD279) is a T-cell co-inhibitory receptor expressed on the surface of activated T-cells, T_{regs} and monocytes had extensively been exploited for immunotherapy.

PD-1 on T-cells interacts with two B7 family ligands, PD-L1 (CD274) and PD-L2 (CD273) expressed on tumor cells that leads to T-cell anergy or death and thus leading to tumor survival [57]. Presence of infiltrating PD-1⁺ T-cells in densely or loosely desmoplastic pancreatic tumors suggests tumor antigen-specific T-cell activation that correlates with increased overall survival, progression-free survival and distant-metastasis-free survival of PC patients [58]. Activated T lymphocytes infiltrating the TME express inflammatory cytokines like IFN γ that further stimulates PD-L1/L2 expression in the tumor cells [59, 60]. Blockade of PD-1 by an mAb abrogates the PD-1/PD-L1 axis and restore T-cell cytotoxic function [61]. In a preclinical study, combined treatment of Panc02 tumor-bearing mice with anti-PD-L1 mAb and gemcitabine significantly reduced average tumor volumes compared to gemcitabine and anti-PD-L1 mAb alone [62]. Due to the relative success of anti-PD-1 antibodies in both preclinical and clinical studies on selective solid tumors, the FDA recently approved two anti-PD-1 antibodies, pembrolizumab and nivolumab, for head and neck cancer, renal, melanoma, and non-small cell lung cancer treatment [63, 64].

Anti CTLA-4 mAbs

CTLA-4 is another co-inhibitory molecule expressed on the surface of activated T-cells and T_{reg} cells. CTLA-4 present on T-cells interacts with B7-1/B7-2 ligands on APCs, resulting in depletion and suppression of CD68-mediated T-cell-activation [65]. Ipilimumab, an antagonist mAb against CTLA-4, inhibits immunosuppression by T_{reg} cells and enhances the antitumor activity of effector T lymphocytes and innate immune cells. In a preclinical study, *in-vitro* treatment with ipilimumab significantly enhanced T-cell proliferation (preferentially promoting CD8⁺ T-cell expansion), Th1 cytokines release (IFN γ , IL-2, and IL-12), and increased cytotoxicity of CD8⁺ T-cells against Colo356/FG PC cells [66, 67]. In a Phase Ib clinical trial, patients with previously treated or histologically proven PC were given ipilimumab alone or in combination with GVAX. Post-treatment,

both the single and combination treatments enhanced mesothelin (MSLN) specific CD8⁺ T-cell populations that correlated with increased survival of >4.3 months, as well as a decline in CA-19.9 levels in 7 out of 15 patients compared to patients treated with ipilimumab alone (0 out of 15 patients) [68]. Combination therapy of anti-CD40, anti-CTLA-4 and anti-PD-1 antibodies with chemotherapy/nab-paclitaxel in KPC mice resulted in tumor regression in 39% of the animals (17 out of 44 mice), along with increased CD8⁺ T-cell infiltration and reduction in T_{reg} cells (7-fold CD8: T_{reg} ratio) in the PC TME. Furthermore, PC tumor cells implanted on the opposite flank were rejected with no additional treatment in 67-86% of mice, suggesting the development of immunological memory [69].

Other targeted therapy antibodies

Additionally, antibodies like cetuximab, pertuzumab, and trastuzumab that bind to the extracellular domain of the human epidermal growth factor receptor 1 (HER1)/epidermal growth factor receptor (EGFR) and inhibit their dimerization and internalization are under clinical investigation [70, 71]. In Phase II and III clinical studies, patients with metastatic PC were administered cetuximab in combination with gemcitabine leading to a stable disease in 63.4% of PC patients and a partial response in 12.2% of patients however there was no increase in the median overall survival of metastatic PC patients. [72]. Contrarily in IMPaCT clinical trial using next-generation sequencing technologies, personalized treatment of 5 PC patient's tumors with human epidermal growth factor receptor 2 (HER2) amplification with trastuzumab showed no successful outcome [73]. Similarly, another Phase II clinical trial showed no responses in 33 advanced PC patients treated with trastuzumab (3.0 or 4.0 mg/kg then 1.5 or 2mg/kg, weekly) and cetuximab (400mg/m² and then 250mg/m²) [74].

Scales et al., developed an anti-mesothelin (anti-MSLN) antibody to the unfolded, non-glycosylated MSLN extracellular domain and conjugated it to the microtubule-disrupting drug monomethyl auristatin E (MMAE). Humanized versions of anti-mesothelin (MSLN)-MMAE induced regression of MSLN-expressing HPAC xenografts in nude mice with a doubling delay to ≥ 74 days. Due to its robust and durable efficacy in mouse models, humanized anti-MSLN-MMAE is in Phase I clinical trials for PC [75]. Similarly, a Phase II clinical trial with the VEGF antibody bevacizumab in 19 out of 30 patients showed stable disease and partial response in 1 patient, although the antibody treatment alone did not improve the overall survival of patients [76].

MDSCs are known to secrete tumor-promoting factors, such as prokineticin 2 (PK2/Bv8). Anti-Bv8 antibody targeting the extracellular domain of Bv8 given in combination with gemcitabine reduced growth of orthotopic metastatic PC tumors, significantly reduced MDSCs infiltration, hypoxia and angiogenesis compared to mice treated with gemcitabine alone, indicating the significant potential of the anti-Bv8 antibody as a combinatorial or post-chemotherapy treatment in PC patients [77]. PC cells express antigens that are either unique to cancer or are being shared with other cancers with similar epithelial origin. The widely studied TAAs of PC that are currently utilized in vaccines in clinical trials are listed in **Table I**.

5.1.2 Passive T-cell-Mediated Immunotherapy

Monoclonal antibody (mAbs) based targeted therapy can elicit direct killing of tumor cells but has not provided long-term benefit to PC patients. Multiple studies are evaluating the strategies to develop passive T-cell-mediated immunotherapies including increasing the number of antigen-specific CD8⁺ T-cells, the responsiveness of the antigen-specific T-cells, or the affinity of the antigen-specific T-cell receptor (TCR) (e.g., with

transfected TCRs). Additional summary regarding the current clinical trials utilizing these strategies is provided in **Table III**.

a) *Adoptive T-cell transfer (ACT)*

The primary objective of ACT therapy is to isolate and expand T-cells *ex vivo* and transfer these autologous lymphocytes with antitumor activity in cancer patients. This method leads to the expansion of antitumor T-cell populations in the patient resulting in increased cytokine release and tumor cell targeting.

Kawaoka et al., developed CTLs by isolating T-cells from the blood of healthy volunteers expressing human leukocyte antigen (HLA)-A phenotype 24/26 and stimulating them with the MUC1-expressing human PC cell line YPK-1 (HLA-A phenotype 24/02) in combination with IL-2. The MUC1-specific CTLs killed five MUC1-expressing PC cell lines, irrespective of their HLA phenotype. 20 patients with either non-resectable or resectable PC were treated with MUC1-specific CTLs. Patients with non-resectable tumor did not show any improvement (median survival time (MST) of 5 months), however, 18 out of 20 patients with resectable PC responded with an MST of 17.8 months [78].

Murine PC cell lines have significant overexpression of telomerase activity. C57BL/6 mice were immunized with H2^b-restricted telomerase peptide emulsified with incomplete Freund's adjuvant, in complex with macrophage-activating lipopeptide-2 (MALP-2, a Toll-like receptor 2/6 agonist) to drive the generation of telomerase-specific CTLs. Orthotopically implanted syngeneic tumor-bearing mice were treated with IL-2-expanded anti-telomerase CTLs, which significantly reduced tumor volume compared to untreated mice. In addition, anti-telomerase CTL-treated mice developed higher numbers of both CD8⁺ central memory and effector antigen-specific T-cells [79]. Furthermore, in a clinical study, 46 PC patients with non-resectable and recurrent tumors received anti-CD3-stimulated lymphokine-activated killer (CD3-LAK) therapy (25 patients) or RetroNectinVR

(CH296)-induced T-cell (RIT) therapy (21 patients) at 2-week intervals. The ACT treated patients showed an increased circulating levels of IFN γ , IL-12, and IL-2, suggesting that the combined circulatory levels of these cytokines may serve as a predictive marker of the clinical response to ACT in patients [80].

b) *Chimeric antigen receptors (CAR) T-cells*

Highly antigen-specific autologous T-cells that are genetically engineered to express tumor antigen-specific TCRs or immunoglobulin-based fusion proteins are known as chimeric antigen receptors (CAR) T-cells. These engineered CAR T-cells are then cultured and expanded *ex vivo* for therapeutic purposes (**Figure 2**).

The differential glycosylation pattern of mucins provides a unique repertoire of antigenic epitopes that can be exploited for developing tumor-specific CAR T-cells. Posey et al. designed a scFv of a high-affinity antibody (5E5) to detect truncated O-glycopeptide MUC1 epitopes (GSTAP with one or two Tn O-glycans on the Ser/Thr residues) that are not expressed in normal tissues. MUC1 CAR T-cells (composed of 5E5 mAb scFV on a CAR backbone of CD8 α , transmembrane domain and costimulatory domains of 4-1BB and CD3 ζ) were generated that target the Tn/STn glycopeptide epitope on MUC1, and upon recognition of MUC1-9Tn, secreted high quantities of IL-2 and IFN γ , but not in response to the non-glycosylated MUC1-60-mer. Hs766T pancreatic tumor-bearing mice when treated with 5E5 CAR T-cells showed potent responses and improved survival to 113 days with 100% animals surviving compared to 40% and 33% of mice treated with non-transduced and CD19 CAR T-cells, respectively. In addition, many 5E5 CAR T-cells specifically accumulated in Hs766T tumors, in contrast to a small percentage of CD19 CAR T-cells [81, 82].

Carcinoembryonic antigen (CEA) is highly overexpressed on the surface of PC cells. Murine CEA binding domain (SCA431scFv)-containing CAR T-cells with intracellular

CD28-CD3 signaling domain were adoptively transferred into Panc02 CEA⁺ tumor-bearing CEA transgenic mice. Anti-CEA CAR T-cells significantly reduced the size of pancreatic tumors and produced long-term tumor elimination in 67% of the mice without inducing an autoimmune reaction. Upon re-challenge with CEA⁺ C15A3 cells, the animals rejected the cells and demonstrated increased serum levels of IL-1 β and IL-5 [83]. A similar study reported the eradication of CEA⁺ tumors in CEA-transgenic mice as a primary response to anti-CEA CAR T-cells with CD3 ζ endo-domain and rejection of CEA⁺ PC cells upon re-challenge. Based on the CAR T-cell model, there is evidence that antigen-specific CD8⁺ T-cells can be induced to overcome self-tolerance and eliminate cancer cells while sparing normal cells [84].

Prostate stem cell antigen (PSCA) is another highly expressed TAAs in PC patients as well as in tumor-derived cell lines. In a recent study, PSCA-specific CAR T-cells showed specific targeting and lysing of PSCA-expressing PC cells (ASPC1, Capan-1) while PSCA-negative 293T-cells showed no cytotoxicity [85]. In another study, anti-PSCA CAR T-cells were engineered using antigen-recognition domains derived from mouse or human antibodies with either one (CD28) or two (CD28 + 4-1BB) T-cell co-stimulatory molecules linked to the CD3 ζ endo-domain. These anti-PSCA CAR T-cells elicited antitumor responses in established human PC-derived xenograft tumors and 2 out of 5 mice showed complete tumor eradication [86].

MSLN is highly overexpressed on PC cells compared to its negligible expression in normal pancreas. Hingorani et al. developed MSLN peptide-specific high-affinity TCR₁₀₄₅ expressing CD8⁺ CAR T-cells that lysed KPC tumor cells *in vitro* and secreted IFN γ upon antigen recognition. A study in KPC mice showed that TCR₁₀₄₅ CAR T-cells infiltrated the pancreatic tumors four days post-injection and induced apoptosis of cancer cells after eight days of infusion. Upon the second infusion, TCR₁₀₄₅ CAR T-cells showed

10-fold increased retention in pancreatic tumor compared to non-specific (TCR_{gag}) T-cells. However, mice in both treatment groups developed progressive disease. TCR₁₀₄₅ cell recipient mice showed less metastasis (46%) and overall survival of 96 days compared to 64% metastatic lesions and survival of 54 days in TCR_{gag} treated mice. Overall, these data suggest that tumor antigen-specific engineered T-cell therapies are viable options for the treatment of invasive PC [87].

5.2 Active Immunotherapeutic Strategies

Active immunotherapy relies on stimulation of the immune system through immunological recognition of TAAs by T and B lymphocytes. TAAs have been widely explored as cancer vaccines for the treatment of PC in both *in vivo* mouse models and clinical trials. Cancer vaccines can be whole cancer cell-based vaccines, antigenic-peptide pulsed vaccines or DC-based vaccines. These vaccines are developed to exploit and activate both innate and active immune arms to eradicate tumor cells and evade future recurrence of the disease. Cancer vaccines currently being investigated in clinical trials in PC are summarized in **Table IV**.

5.2.1 Cancer Vaccines

Mucin (MUC) Vaccines

Mucins (MUCs) are glycoproteins that are differentially overexpressed in pancreatic tumor but is absent in normal pancreas. Some mucins (e.g., MUC1 and MUC4) have also been demonstrated to contribute to chemoresistance, and to enhance proliferation and survival of PC cells [88]. Therefore, mucins are being studied as potential candidates for vaccine development for PC. Studies conducted in human MUC1-transgenic (MUC1.Tg) mice treated with MUC1 cancer vaccines failed to show any detectable responses against MUC1⁺ tumor cells despite, MUC1-specific T-cells

generating IFN γ , IL-4 and IL-10 cytokines. The immune responses in these mice were not skewed to either type 1 or type 2 immune response thus rendering the vaccine ineffective against the B16.MUC1 tumor.[89]. The CD8⁺ T-cell killing of MUC1-expressing tumor cells was found to be mediated by perforin and FasL cytolytic pathways. Also, lymphotoxin- α , but not TNF receptor-1 (TNFR-1), played a critical and non-redundant role in the cell-mediated rejection of MUC1 expressing tumor cells [90]. In a Phase I clinical trial, MUC1-peptide (GVTSAPDTRPAPGSTAPPAH)-pulsed DC vaccines were administered to 7 patients with advanced PC. 2 out of 7 patients showed significantly increased mature DCs and peripheral blood mononuclear cell (PBMC)-mediated immune responses that were characterized by high IL-12p40 and IFN γ secretion, respectively. However, there was no tumor rejection in these patients [91]. A similar Phase I clinical trial in 20 advanced PC patients was performed with MUC1 peptide-pulsed DCs in combination with MUC1-specific CTLs. One patient with lung metastases showed complete remission, while five other patients demonstrated stable disease for at least six months post-therapy [92]. A study was conducted with 6 PC metastatic patient-derived DCs that were co-transfected with MUC4 and survivin mRNAs. These mRNA-loaded DCs activated CTLs against the MUC4 protein. Anti-MUC4 CTLs effectively targeted a human PC cell line (Capan-2) via major-histocompatibility-complex-I (MHC-I)-restricted recognition and released IFN γ . MUC4-mRNA-pulsed DCs stimulated more CTLs than survivin-mRNA-pulsed DCs, but comparatively elicited fewer CTLs activated by MUC4-survivin-mRNA-loaded DCs [93]. In another study, mature DCs were pulsed with a MUC4 epitope peptide (LLGVGTFVV) and co-cultured with CD8⁺ T-cells to generate MUC4-specific CTLs that could effectively kill HCT-116 colorectal cancer cells (MUC4⁺, HLA-A2⁺). However, the intensity of MUC4 surface expression on PC cell-line HPAC proportionally increased the apoptosis of MUC4-specific T-cells, thus rendering the therapy ineffective [94]. Based on these studies, other

mucins like MUC5AC, and MUC16 that are significantly overexpressed in PC may serve as potential vaccine candidates to develop novel immunotherapies.

Telomerase Vaccines

Due to overexpression of telomerase in PC patients, cancer vaccine containing telomerase-derived peptide (GV1001) vaccine is under clinical studies. However, so far PC patients treated with a combination of the GV1001 vaccine, GM-CSF, and gemcitabine showed transitory and weak Th1-type immune response, reduced infiltration of T_{reg} cells, and no significant increase in median overall survival. In a related clinical trial, the GV1001 vaccine failed to enhance the effects of chemotherapy (gemcitabine and capecitabine) [95, 96].

CEA Vaccines

CEA is overexpressed in >90% of PC patients making it a potential immunotherapeutic target. A Phase I clinical trial with CAP1-6D, an altered CEA peptide ligand/Montanide/GM-CSF vaccine, administered to 66 PC patients elicited robust CD8⁺ T-cell-mediated. In addition, 8 of the patients were found to have high IFN γ production, of which four patients showed progressive disease, three patients had stable disease, and one patient showed a complete response [97].

KIF20A Vaccines

Another cancer vaccine candidate is kinesin-like protein 20A (KIF20A), a member of the kinesin super family, which is significantly upregulated in PC. KIF20A-66, an HLA-A24-restricted epitope peptide derived from KIF20A peptide vaccine was injected into 29 PC patients in Phase II clinical trial. The tumor size (as confirmed by CT scan) was reduced in 8 patients and 16 patients showed increased CTL responses, which correlated with the increased overall survival of vaccinated patients [98]. In a similar Phase I trial,

nine advanced PC patients who previously received chemo/radiotherapy were treated with KIF20A-10-66 (KVYLRVRPLL) peptide vaccine along with gemcitabine. These patients showed increased activation of KIF20A-specific IFN γ -producing T-cells, had stable diseases and longer overall survival, indicating that KIF20A-based vaccines are possible immunotherapy candidates for PC [99].

K-ras Vaccines

K-ras is mutated in the majority of PC patients and is currently under investigation as an immunotherapy target. Inactivation of the oncogenic mutant K-ras enhances MHC I presentation [100]. In an *in vivo* study, mice were treated with lysates of human PANC-1 PC cells expressing mutant K-ras, with lysate in combination with K-ras mutant peptide (K-ras+peptide), or with lysate and peptide plus cationic nanoparticles (CNP) encapsulating K-ras mutant peptide (K-ras+peptide-CNP). The K-ras+peptide-CNP activated CTLs induced specific killing of K-ras-positive tumors efficiently and improved the survival time of K-ras mutant tumor-bearing nude mice compared to the K-ras mutant lysate and K-ras peptide treatment group (132). In addition, a clinical trial testing a mutant Ras peptide vaccine (administered in combination with GM-CSF) in 11 patients with advanced PC showed higher antitumor responses (92% exhibiting an immune response) with two patients having a complete response and overall survival of 20.8 months [101, 102].

WT1 Vaccines

The Wilm's Tumor protein 1 (WT1) is another suitable vaccine target for PC due to its differential overexpression in tumor cells but not in the normal pancreas [103]. In a recent study, 32 HLA-A*24:02⁺ advanced PC patients were treated with an HLA-A*24:02-restricted, modified 9-mer WT1 peptide (CYTWNQMNL) emulsified with Montanide ISA51 adjuvant (WT1 vaccine). The MST of patients who responded to the WT1 vaccine was

10.9 months. Further, these patients developed strong effector T-cell responses along with generation of WT1-specific CD8⁺ memory T-cells, whereas unresponsive patients showed MST of only 3.9 months [104]. In a separate clinical trial, nine patients with advanced PC were vaccinated with WT1 peptide vaccine and 8 out of 9 patients demonstrated stable disease. However, no WT1-specific T-cells were observed in the circulation of these patients [105].

VEGF Vaccines

VEGF is another protein that is overexpressed in PC. In a trial, nine advanced PC patients were vaccinated with four peptides comprised of KIF20A, cell division cycle-associated 1 (CDCA1), VEGFR1, and VEGFR2. Patients demonstrated increased anti-CDCA1 and anti-VEGFR2 CD8⁺ T-cells in circulation. Additionally, 4 out of 9 patients presented with stable disease post-vaccination [106]. In a subsequent study, vaccination with WT1 and VEGFR2 peptides generated HLA-A24-restricted CTLs, which demonstrated strong cytotoxicity towards PC cells that were HLA-A24-positive and expressed corresponding TAAs [107].

Prophylactic Cancer Vaccines

Prophylactic vaccines have recently been investigated as immunotherapy tools to target endogenous neoantigens by utilizing attenuated bacteria/virus to stimulate antitumor adaptive immune responses [108]. *Listeria monocytogenes* (LM) is a gram-positive bacterium that induces robust CD4⁺ and CD8⁺ T-cell responses by its selective infection of APCs (via *actA* virulence gene) over non-phagocytic cells (*inIB* gene). An LM $\Delta actA/\Delta inIB$ strain engineered to express human MSLN (CRS-207 vaccine) was administered to 10 PC patients, which resulted in induction of expression of Th1 cytokines (IL-12, TNF- α). In addition, 6 out of the 10 patients developed MSLN-specific CD8⁺ T-cells [109, 110]. Jaffee et al. conducted a clinical trial on 93 metastatic PC patients, in which 69

patients received two doses of cyclophosphamide with GVAX (Cy/GVAX) followed by four doses of CRS207 (Arm A) and 21 patients received six doses of Cy/GVAX (Arm B). Patients in Arm A showed increased overall survival of 9.7 months compared to 4.6 months in Arm B, suggesting that the success of this immunotherapy might depend on the proper patient selection [111]. The Kras^{G12D} oncogene prophylactic vaccine (LM-K-ras vaccine) has been studied in KPC mice, either alone or in combination with T_{reg} depletion (anti-CD25 antibody, PC61, and low-dose Cy). KPC mice that received the vaccine at early PanIN 1 stages in combination with T_{reg} depletion showed prolonged survival compared to mice that received the vaccine alone at either early or late PanINs stage, suggesting the potential of T_{reg} depletion therapy as the prophylactic approach for PC [112].

STING activation as an adjuvant to vaccines and other immunotherapies

STING is a transmembrane protein that resides on the endoplasmic reticulum, which upon activation through cyclic dinucleotides (CDNs), synthetic CDNs, or bacterial infection in the host induces interferon beta (IFN β) and nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) pro-inflammatory responses via IRF3 and Stat6 pathways [113]. It has been shown that endogenous STING activation via CDNs in the tumor microenvironment enhanced inflammatory responses, thereby inhibiting tumor progression and distant metastasis [114]. Recently, intraperitoneal injection of DMXAA, an activator of the mouse STING pathway, activated CD8⁺ T-cells that led to tumor rejection [114, 115]. Similarly, synthetic STING activators known as RR-CDGs have shown efficacy in the regression of primary pancreatic tumors, and distant metastatic lesions through T-cells recruitment in a TNF- α -dependent manner [116]. Furthermore, these novel synthetic activators of STING have demonstrated enhanced adjuvant activity to accelerate adoptive immune responses in the presence of radiation therapy [116]. The

cGAMP-induced activation of endothelial cell-specific STING enhanced the antitumor responses of CD8⁺ T-cells and improved the responses of anti-CTLA-4 and anti-PD-1 immunotherapies [117].

6. Challenges in Pancreatic Cancer (PC) Immunotherapy

The immune system has the potential to selectively target tumor cells upon strategic activation in cancer patients leading to better therapeutic outcomes. However, tumors employ extensive measures to escape immune surveillance, suggesting the necessity to develop novel counteracting strategies for the improved efficacy. Therefore, recent immunotherapeutic approaches alone or in combination with conventional treatment modalities need to be re-evaluated for successful therapeutic outcome in terms of improved patient survival.

6.1 Identification of Tumor Associated Antigens for PC Immunotherapy

Tumor-associated antigens are autologous cellular antigens that are specifically expressed by cancer cells and are negligibly expressed by or absent in normal cells. Mutation-derived tumor antigens are generated by somatic mutations inherited by tumor cells during malignant transformation that may be identified by immune surveillance [118]. The Cancer Genome Atlas (TCGA) database analysis showed that notably PC afflicted patients carry limited genetic mutations between 4 to 4000 neoantigens [119, 120]. When compared to melanoma patients who could likely express 14000 neoantigens that is a significantly low number of neoantigen repertoire in PC patients. Further expression neoantigens mutation load correlated negatively with T-cell tumor infiltration and corresponding overall survival of PC patients [119]. Differentially and spatiotemporally overexpressed or post-translationally antigens could potentially serve as tumor antigens for immunotherapy purposes [121]. A preferred way utilized to validate the likelihood of

these tumor antigens to be immunogenic in patients, is to detect autoantibodies against them in cancer patient serums and this strategy has met with moderate success [122, 123]. Multiple proteins like mucins (MUC1), WT1, VEGF, etc. overexpressed in PC patients, have been extensively exploited for PC immunotherapy but have achieved limited success as discussed previously in the review. It has been challenging to identify eligible TAAs and thus, necessitates coming up with a strategy to discover novel tumor antigen that could be targeted efficaciously by immunotherapy in PC patients.

6.2 Generation of Anti-Tumor Responses Against PC Microenvironment

Strong immunosuppressive TME, rigid tumor matrix architecture, T_{regs} infiltration, constant antigen exposure mediated T-cell exhaustion, and upregulation of inhibitory receptors like PD-1 provide physical obstacles to effector T-cells function and generates tolerance towards tumors [124, 125]. Thus, checkpoint inhibitors and depletion of T_{regs} could potentially reverse T-cell exhaustion of effector T-cells. Improper homing and inefficient infiltration of CAR T-cells to the tumor bed occur due to tumor blood vessels not responding to inflammatory stimuli. Anti-angiogenic therapy matures tumor blood vessels, facilitating pericyte recruitment and increasing tumor perfusion, which consequently increases the efficacy of CAR T-cell immunotherapy [126]. Apart from improper homing, CARs recognize TAAs that are also found at a lower level in normal tissues (which can cause toxicities in PC patients), thus if CAR T-cells survive for long periods of time in patients they increase the risk of developing autoimmunity in the future [127]. Furthermore, activated CAR T-cells containing co-stimulatory domains like CD27, CD28 or 4-1BB release a variety of inflammatory cytokines like IL-2, IL-6, and IFN γ after encountering tumor cells which induce macrophages to release more inflammatory cytokines thus establishing a positive cytokine-based feedback loop to enhance T-cell activity causing cytokine release syndrome (CRS), which could be fatal for patients [128, 129].

Development of short-lived CAR T-cells or combination treatment with an IL-6 receptor inhibitor like tocilizumab could be effective in reversing the effects of CRS without affecting the activity of CAR T-cells [128, 130].

Human PC malignancy is complex because of its TME architecture and cytokine milieu that is inefficiently recapitulated in KPC or xenograft mouse models. Secondly, both tumor cells as well as normal cells express common antigens (though usually at a lower level than normal cells), and therefore cancer vaccines can potentially cause toxicities in the patients. Thirdly, peptide-based cancer vaccines do not capture all unique immunogenic epitopes present on original tumor antigens. Pancreas-specific transgene expressing spontaneous PC mouse models, and either protein fragments or intact proteins as immunogens could address the limitations faced in this field, thereby increasing the cancer vaccine efficacy. Additionally, selection of PC patient based on both tumor stage and tumor infiltrating lymphocytes (TILs) like CD8⁺ and PD-1⁺ T-cells could further increase the response to cancer vaccines [131].

6.3 Overcoming Immunosuppression in PC Microenvironment

Nanoparticles are capable of encapsulating multiple proteins, ligands, nucleic acids and other materials, thus increasing the epitope repertoire. Nanoparticles can also incorporate immune-stimulatory adjuvants (such as TLR agonists) or chemotherapeutic drugs to enhance the overall immunogenicity, stability, delivery and/or direct cytotoxicity of the vaccine, therefore overcoming the limitations of current cancer immunotherapies [132]. For example, mice immunized with Doxorubicin-CpG-PLGA microparticles showed a reduced tumor burden at lower drug concentrations compared to mice that received doses of the soluble drug. When combined with an anti-CTLA-4 antibody, the treatment successfully reduced aggressive tumor burden at both the injected and distant tumor sites in tumor-bearing mice [133]. This co-encapsulation of multiple therapeutics and immune

stimulatory molecules may provide dose-sparing capabilities, reducing the cost and toxicity of cancer therapeutics [134].

Encapsulation into biodegradable nanoparticles protects the payload until release [135, 136]. In addition, tuning the polymer chemistry enables sustained and controlled release of encapsulated payloads [137] and immunomodulatory capabilities [138]. Particularly, it has been demonstrated that varying chemistries of polyanhydride nanoparticles were efficiently internalized by APCs, leading to the upregulation of MHC I, major histocompatibility complex II (MHC II) and costimulatory molecules, as well as inducing the secretion of cytokines [139-141]. In addition, amphiphilic nanoparticles promoted the production of long-lived, high avidity antibody [142] with suboptimal doses of antigen [143], suggesting the development of long-lived plasma cells. Polyanhydride nanoparticles loaded with OVA-induced memory CD8⁺ T-cells that were recruited and responded to subsequent challenges with OVA-secreting tumor cells [144]. Finally, many nanoparticles can be functionalized with ligands or antibodies that may increase selectivity and reduce the side effects of chemotherapeutics on healthy tissues [145]. Targeting moieties are often attached to the nanoparticle surface via a polyethylene glycol (PEG) linker [145]. This method of PEGylation allows for flexibility of the targeting moiety and may enhance interactions with cancer cell receptors [145, 146]. For example, PLGA nanoparticles covalently modified with folate via PEG demonstrated an increased association and uptake with cancer cells *in vivo* [147].

The limited success of immunotherapeutic studies performed in PC provides a generous room for improvement. Tailoring immunotherapy to PC patients by identifying unique tumor-specific antigens through genetic screening and expression studies [49, 148] and combining it with continuous collection and screening of tumor samples in clinical

trials to understand immunotherapy resistance, will further improve the response rates and survival benefits of PC immunotherapy.

Table I: Tumor associated antigens targeted for PC immunotherapy.

Tumor Associated Antigen (Cancer Antigen)	Expression		Description
	Normal Pancreas	Pancreatic Cancer	
Mucins MUC1 MUC4 MUC5AC	MUC4 and MUC5AC are undetectable, while MUC1 is expressed at low levels.	Aberrantly overexpressed and glycosylated in PC patients	<p>Mucins are glycoproteins that are differentially overexpressed in PC but are not expressed in the normal pancreatic epithelium (except for MUC1 which is expressed at low level). These mucins (e.g., MUC1, MUC4, and MUC5AC) are involved in PC pathogenesis, provide chemoresistance and enhance proliferation and survival of PC cells. Their overexpression has been correlated with poor prognosis in patients.</p> <p>[Ref. No. 54,64,69-70]</p>
Telomerase	Absent	Expressed in 80-90% of PC patients	<p>Telomerase is a ribonucleoprotein enzyme that catalyzes the synthesis of telomeric DNA. It is involved in the formation and protection of the telomere, which prevents cells from undergoing senescence. Telomerase activity has been detected in pancreatic juice samples of PC patients. hTERT expression and telomerase activity are predictors of poor outcome in pancreatic cancer.</p> <p>[Ref. no. 55, 71-72]</p>
Carcinoembryonic antigen (CEA)	Absent	Expressed in 77% of PC patients and detected in patient serum.	<p>Carcinoembryonic antigen (CEA), a glycosylated protein of MW 180 kDa, is related to tumor burden of PC due to its close association with cancer</p>

			<p>cell adhesion, metabolism, and proliferation. In clinical practice, CEA is often used to predict the outcomes of patients with resectable PC.</p> <p>[Ref. no. 60,73]</p>
Mutated K-ras (G12D)	Absent	Expressed in 89.8-94.9% of PC patients	<p>K-ras is mutated in PC cells and inactivation of the oncogenic mutant K-ras enhances MHC I presentation. K-RAS belongs to the superfamily of small G proteins and plays crucial roles in signal transduction in cells. K-RAS mutations in PC transform and alter the biological behavior in PC cells including metabolism reprogramming, thus playing a crucial role in PC pathogenesis.</p> <p>[Ref. no. 77-80]</p>
Vascular endothelial growth factor (VEGF)	Absent	Expressed in 77-93% of PC patients	<p>VEGF, primarily VEGF-A and its receptors (VEGFR1 & VEGFR2), are primarily involved in the angiogenesis process in PC cancer. Increased vascularization of pancreatic tumors promotes their growth and metastasis by providing nutritional flow. Neovascularization also facilitates infiltration of pro-tumor immune cells (e.g., MDSCs).</p> <p>[Ref. no. 82-83]</p>
Mesothelin	Absent	Expressed in ~86% of PC patients	<p>Mesothelin (MSLN) is a glycoprotein overexpressed in various epithelial cancers like mesothelioma and pancreatic, ovarian, and lung cancers. MSLN is synthesized as a 71 kDa precursor protein, which is processed to a 30 kDa megakaryocyte-potentiating factor and a 40 kDa MSLN</p>

			<p>protein. It is attached to the plasma membrane by a glycosylphosphatidylinositol anchor and is involved in cell adhesion. MSLN serves as a marker of neoplastic transformation of pancreatic epithelial cells.</p> <p>[Ref. no. 86-87]</p>
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Table II: Clinical trials testing antibody-based immunotherapies for pancreatic cancer.

Antigen & Drug	Clinical Trials .gov Identifier	Phase	No. of Patients	Status	Median overall survival (months)	Median progression-free survival (months)	Objective response rate (%)
CD40 (CP-870,893) 0.2 mg/kg	NCT00711191 [Ref. No. 35]	I	22	Completed	7.4 (5.5 to 12.8)	5.6 (4.0 to 7.4)	7.7 (0.2 to 36.0)
CD40 (CP-870,893) + Gemcitabine	NCT01456585*	I	10	Completed	NR ^b	NR	NR
PD-1 (CT-011) alone or in combination with Gemcitabine	NCT01313416*	II	29	Suspended	NR	NR	NR
PD-L1 (pembrolizumab)	NCT02362048*	II	73	Active	NR	NR	NR
PD-L1 (pembrolizumab)	NCT02009449*	I	350	Active	NR	NR	NR
CTLA-4 (ipilimumab)	NCT00112580	II	27	Completed	NR	NR	1 patient had PR ^c
CTLA-4 (ipilimumab) + Pancreatic Cancer Vaccine	NCT00836407*	I	30	Completed	5.7 (4.3 to 14.7)	NR	NR

* Data obtained from <https://clinicaltrials.gov/>

HER1 (Cetuximab) + Irinotecan + Docetaxel	NCT00 042939 *	II	87	Compl eted	5.3 (4.5 to 9. 4)	4.5 (2.7 to 5. 6)	0.07 (0.024 to 0.198)
HER1 (Cetuximab) + Gemcitabine + Radiotherapy	NCT00 225784 *8	II	37	Compl eted	17.3 (2 to N/A)	9.1 (2 to N/A)	10 out of 37 had PR
HER1 (Cetuximab) + Ixabepilon e	NCT00 38314*	II	54	Compl eted	7.6 (5.5 to 1 2.2)	3.9 (2.6 to 4. 4)	4 patients had PR
HER1 (Cetuximab) + Irinotecan + Oxaliplatin	NCT00 871169 *	II	58	Compl eted	NR	NR	6.9 (1.91 to 1 6.7)
HER1 (Cetuximab) + Gemcitabine + VEGF (Bevacizuma b)	NCT00 326911 *	II	30	Termi nated	5.41 (3.84 to 6.74)	3.55 (2.00 to 5 .59)	4 patients had either PR or CR ^d
HER1 (Cetuximab) + Gemcitabine + VEGF (Bevacizuma b)	NCT00 091026 *	II	71	Compl eted	7.9 (5.5 to 9. 5)	5.0 (3.7 to 5. 5)	21 (12 to 32)
HER1 (Cetuximab) + Gemcitabine + Oxaliplatin	NCT00 338039 *	II	69	Compl eted	19.2 (14.2 to 24.2)	NR	NR

HER1 (Cetuximab) + Gemcitabine + Capecitabine + Radiation	NCT00 305877 *	II	65	Compl eted	0.38 (0.26 to 0.50)	0.17 (0.08 to 0 .26)	0.30 (0.19 to 0. 42)
HER2 (Trastuzumab) + Interleukin 12	NCT00 004074 *	I	15	Compl eted	NR	NR	NR
HER2 (Trastuzumab) + HER1 (Cetuximab)	NCT00 923299 [Ref. No. 51]	I & II	44	Compl eted	4.6 (2.7-6.6)	1.8 (1.7-2.0)	NR
Mesothelin (SS1(dsFv)- PE38 immunotoxin)	NCT00 006981 *	I	NR	Compl eted	NR	NR	NR
VEGF (Bevacizuma b) + Gemcitabine + accelerated Radiation Therapy	NCT00 557492 *	II	43	Ongoi ng	19.7 (16.5 to 28.2)	12.9 (7.0 to 18 .7)	2.3 (0.1 to 12)
VEGF (Bevacizuma b) + Octreotide Acetate + Everolimus	NCT01 229943 *	II	75	Ongoi ng	36.7 (31.8 to N/A)	16.7 (12.6 to 1 9.7)	31

* Data obtained from <https://clinicaltrials.gov/>

^b NR (not reported).

^c PR (partial response).

^d CR (complete response).

Table III: Clinical trials testing T cell-mediated immunotherapies for pancreatic cancer.

Antigen & Drug	ClinicalTrials.gov Identifier	Phase	No. of Patients	Status
Anti-CEA CAR-T cells	NCT02416466*	I	8	Ongoing
Autologous T cells transfected with chimeric anti-mesothelin immunoreceptor SS1	NCT01897415*	I	16	Ongoing
CART-meso-19 T cells + Cyclophosphamide	NCT02465983*	I	12	Ongoing
GI-4000 Vaccine + Activated T Cells	NCT00837135*	I	NR ^b	Withdrawn
MFE23 scFv-expressing autologous anti-CEA MFEz T lymphocytes	NCT01212887*	I	14	Terminated due to safety concerns and lack of efficacy
Autologous Natural Killer / Natural Killer T Cell Immunotherapy	NCT00909558*	I	24	Suspended
Prostate Stem Cell Antigen (PSCA) Specific CAR T Cells (BPX-601) + Rimiducid	NCT02744287*	I	30	Recruiting

* Data obtained from <https://clinicaltrials.gov/> ^bNR (not reported).

Table IV: Clinical trials testing cancer vaccines for pancreatic cancer.

Antigen & Drug	ClinicalTrials.gov Identifier	Phase	No. of Patients	Status
MUC1 Vaccine (Cvac vaccine)	NCT02310971*	II	0	Withdrawn
Falimarev (MUC1 PANVAC-F vaccine) + Inalimarev (MUC1 PANVAC-V vaccine) + Sargramostim (GM-CSF vaccine)	NCT00669734*	I	18	Ongoing
Telomerase vaccine (GV1001) + gemcitabine + Sargramostim + tadalafil + Radiation Therapy	NCT01342224*	I	11	Active
Telomerase vaccine (GV1001) + Sargramostim + capecitabine + gemcitabine	NCT00425360*	III	1110	Completed
CEA vaccine (ALVAC + vaccinia) + aldesleukin (IL-2) + Sargramostim	NCT00003125*	II	24	Completed
CEA vaccine (AVX701)	NCT00529984*	I & II	28	Completed
Recombinant fowlpox-CEA(6D)/TRICOM vaccine + GM-CSF vaccine + Sargramostim	NCT00028496*	I	48	Completed
Recombinant fowlpox-CEA(6D)/TRICOM vacci	NCT00128622*	I	24	Completed

ne + denileukin diftiox + therapeutic autologous dendritic cells				
CEA vaccine (TRICOM-CEA(6D))	NCT00027534*	I	14	Completed
CEA RNA-pulsed DC cancer vaccine	NCT00004604*	I	24	Completed
CEA vaccine (carcinoembryonic antigen peptide 1-6D) + incomplete Freund's adjuvant + sargramostim	NCT00012246*	II	7	Terminated
K-ras vaccine (TG01) + GM-CSF + Gemcitabine	NCT02261714*	I & II	32	Active
Aldesleukin + ras peptide cancer vaccine + sargramostim + DetoxPC	NCT00019331*	II	11	Completed
HLA-A*02:01-restricted VEGFR1-derived peptide vaccination	NCT00683085*	I & II	2	Terminated
VEGFR-2 DNA vaccine VXM01	NCT01486329*	I	72	Completed
Mesothelin vaccine (CRS-207) + GVAX vaccine + gemcitabine + capecitabine + 5-FU + irinotecan or erlotinib + cyclophosphamide	NCT02004262 [Ref no. 87, 88]	II	303	Completed
GVAX Pancreas + Mesothelin vaccine	NCT01417000	II	93	Ongoing

(CRS-207) + Cyclophosphamide	[Ref. no. 88]			
Mesothelin vaccine (CRS-207)	NCT00585845*	I	17	Terminated
Cancer stem cell vaccine	NCT02074046*	I & II	40	Completed

* Data obtained from <https://clinicaltrials.gov/>

Figure 1: Pancreatic cancer cells establish an immunosuppressive TME.

Cancer cells secrete various anti-inflammatory cytokines like IL-10, TGF- β , IL-23, along with angiogenic chemokines (e.g., CXCL1-3, CXCL5, CXCL12, CCL2, and VEGF-A), which generate an immunosuppressive tumor microenvironment (TME) and facilitate cancer initiation, progression and metastasis. Upregulation of the expression of these cytokines shifts the balance in TME, which facilitates the evasion from immune surveillance during PC progression [6, 8, 20, 28]. The PC immunosuppressive microenvironment also includes crosstalk between cancer cells and various myeloid and lymphoid subsets. Tumor-associated macrophages (TAMs) and myeloid-derived suppressor cells (MDSCs) express immuno-inhibitory ligands and reactive oxygen species that inhibit infiltration and activation of T and NK cells [3, 11-13]. MDSCs and cancer cells also secrete VEGFs that promote angiogenesis, which aids in the metastasis of the cancer cells [17]. PC tumor cells and pancreatic stellate cells (desmoplasia) secrete inhibitory cytokines and chemokines, and express inhibitory surface ligands such as programmed death ligand-1 (PD-L1) and galectin-1 (Gal-1) that lead to inactivation and apoptosis of cytotoxic (CD8⁺) and helper (CD4⁺) T-cells by programmed death receptor-1 (PD-1) or Gal-1 binding receptor respectively [16, 18, 21, 22]. T_{reg} cells suppress the functions of activated T-cells and NK cells in the TME [13, 125]. In addition, the rigid architecture of pancreatic tumor bed provides a physical barrier to T-cells infiltration thereby excluding them to the edge/boundary of the tumor and thus rendering the pancreatic tumor as an immunologically 'cold' tumor [28-31]. All these cells are involved in the maintenance of the immunosuppressive TME, and cancer progression.

Figure 1

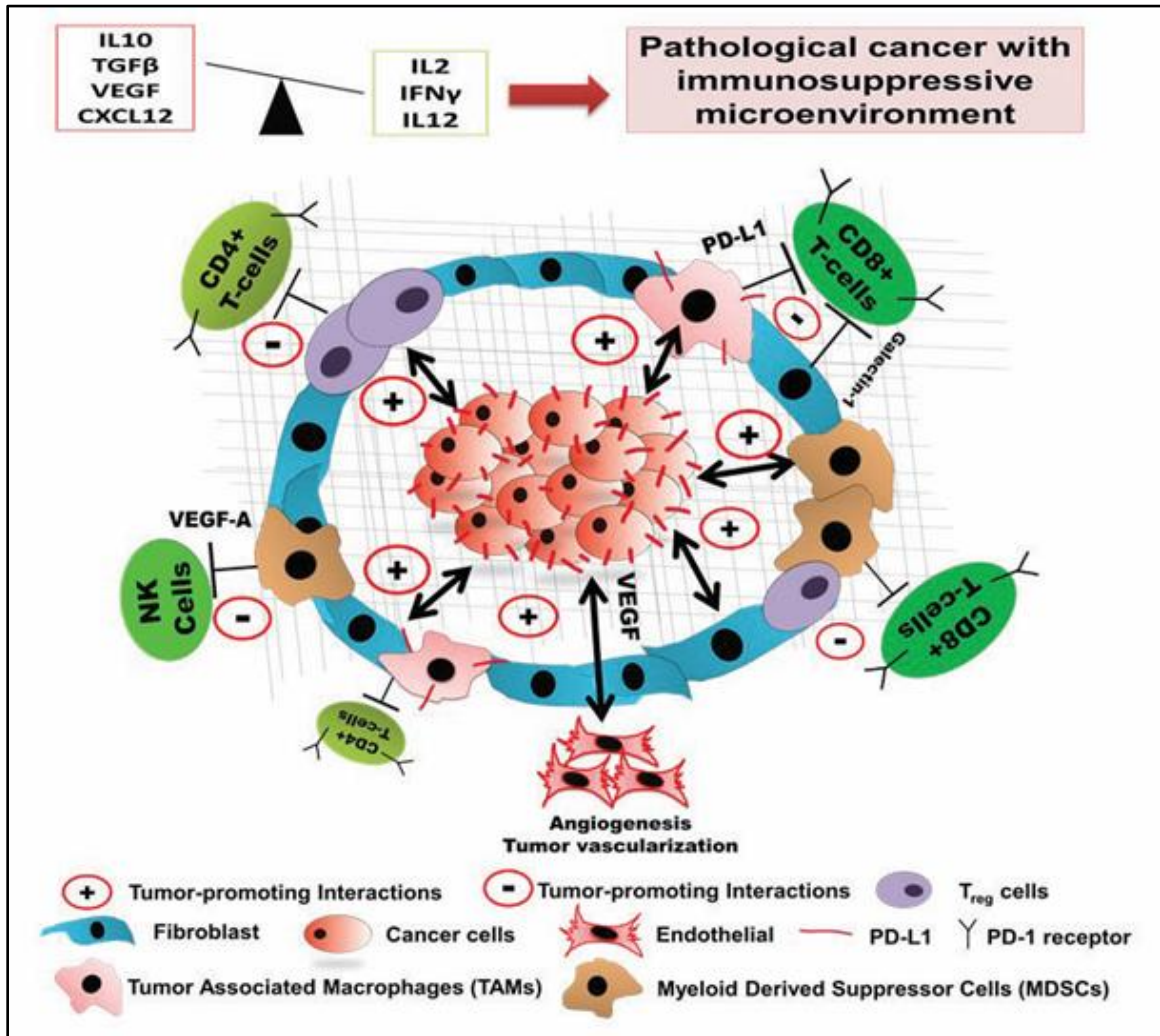
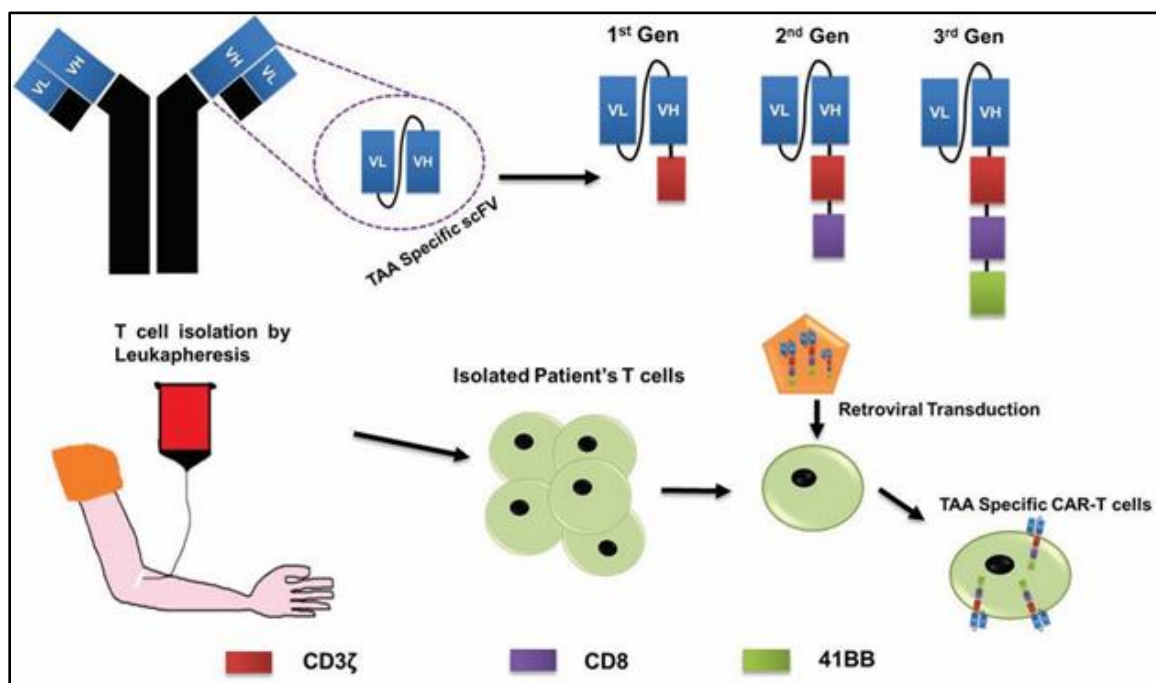


Figure 2: Schematic representation of engineering of CAR-T cells.

CAR T-cells are genetically engineered T-cells expressing tumor antigen-specific chimeric TCR [149, 150]. The modified receptor is a chimera of a signaling domain of the TCR complex and an antigen-recognizing domain, such as a single chain fragment (scFv) of an antibody [151, 152]. CAR T-cells are not dependent on antigen presentation by MHC molecules expressed on APCs for antigen-specific activation. Adoptive cell transfer of CAR T-cells involves the isolation, stimulation, expansion, transduction, and ultimately re-infusion of human T lymphocytes [153, 154]. First-generation TCRs included only the intracellular domain of the CD3 ζ chain but did not show any significant *in vivo* efficacy in transgenic mouse model studies [155]. Second-generation CARs introduced additional co-stimulatory domains such as CD28, which significantly augmented CAR signaling, and improved cytokine production and T-cell proliferation, as well as differentiation, and survival [156, 157]. Third-generation CARs contain multiple co-stimulatory domains such as 4-1BB (CD137), and whether they have a clinical benefit over second-generation CAR T-cells is still under investigation [152, 158, 159].

Figure 2



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**CHAPTER 1B: MUCINS-BASED
VACCINES AND MUC4 IN
PANCREATIC CANCER**

1. Synopsis

Mucins are high-molecular-weight, heavily O-glycosylated glycoproteins that are differentially overexpressed in pancreatic cancer. Literature surveys has shown the instrumental role mucins play in the pathobiology of PC disease. Overexpression, aberrant glycosylation and pathological role proposes mucins as lucrative targets for developing targeted therapies. Immunotherapy is one of the strategies that specifically target tumor-associated antigens and simultaneously reprograms the patient's immune system towards the targeted killing of tumor cells. MUC1 has been one of the well-explored mucins for developing PC immunotherapies but achieved limited success in PC patients. MUC4 is one of the differentially overexpressed mucins in PC and unlike MUC1, MUC4 has unique tumor specificity since it is undetectable in normal pancreas. MUC4 plays a crucial role in PC disease progression and metastasis. Further MUC4's expression has been demonstrated to be regulated by cytokines such as IFN γ and tumor growth factor beta (TGF β), suggesting a possible interrelationship between immune tumor microenvironment and PC disease aggressiveness. Previous studies have provided evidence that makes a strong case to evaluate MUC4 as an immunotherapeutic candidate, but to date only limited studies have been made. Therefore, in this section we have reviewed existing literature and summarized their findings to make a case for investigating MUC4 as a potential candidate for PC immunotherapy.

2. Mucins: A Target for Pancreatic Cancer Immunotherapy

PC cells undergo a cellular and genetic transformation that contributes to the expression of proteins that are mostly absent in *de novo* pancreas cells. One of the families of proteins that is aberrantly overexpressed in PC tumor cells and has been reported to play a crucial role in the pathogenesis of the disease is mucins. MUCs are high molecular weight glycoproteins with O-glycosylated variable number tandem repeat (VNTR) region and consists of 21 family members. Fourteen MUC proteins out of 21 mucin family members fall under the category of classical mucins, which is characterized by high molecular weight O-glycoprotein, secreted in the mucus layer, presence of VNTR sequence, predicted peptide domain containing high percentage of serine and threonine residues, and lastly a complex mRNA expression of these mucins [1]. MUC glycoproteins are classified according to their structure and function into two categories: transmembrane/membrane-bound mucins comprising of MUC1, MUC3A, MUC3B, MUC4, MUC12 and MUC17; and secretory/gel-forming mucins consisting of MUC6, MUC2, MUC5AC and MUC5B [2].

Transmembrane mucins like MUC1, MUC4 and MUC16 are differentially overexpressed by PC cancer cells and their expression gradually increases with the progression of the disease [3-9]. MUC1 and MUC4 interact with various receptors such as receptor tyrosine kinases (RTKs) including epidermal growth factor receptor (EGFR), ERBB2 & ERBB3, and fibroblast growth factor receptor (FGFR), that promotes proliferation, invasion, metastasis and resistance to therapeutic anti-RTK (Receptor Tyrosine Kinase) antibodies [10-14]. MUCs expressed on the PC cancer cell surface due to their large structure interact with the TME that induce immune evasion and oncogenic signaling-mediated angiogenesis and metastasis [2]. Studies have demonstrated that MUCs like MUC1 on PC cells interact with TAMs through sialoadhesin [15]. Interaction

between M2 macrophages with PC cells via CA125 (carbohydrate epitope located on MUC16) differentiates these macrophages to immunosuppressive phenotype and induces secretion of IL10 cytokine [16]. In addition to MUC's crucial role in pathogenesis and immunosuppression of PC, they undergo aberrant glycosylation due to the altered expression and localization of glycosyltransferases [17]. Thus, these aspects make MUCs an important therapeutic target for PC immunotherapy strategies.

2.1 Mucin-based Cancer Vaccines for PC

Due to their contributions in PC pathogenesis, MUCs have been targeted by immunotherapy strategies like cancer vaccines to efficaciously treat PC patients. We have previously discussed MUC vaccines studied and tested in both pre-clinical and clinical trials for PC immunotherapy in our previous chapter [18]. In addition, a recent pre-clinical study with MUC1-based DNA vaccine (pVAX1-MUC1-VNTR_n DNA vaccine) showed that MUC1-VNTR₆- and MUC1-VNTR₉- transfected DCs were able to activate IFN- γ producing CTLs, increase the cytotoxicity of CTLs and suppress Panc02-MUC1 PC tumor cell's growth in tumor-bearing mice [19]. Mice immunized with MUC1-tandem repeat B-cell peptide conjugated with Ttox (MUC1-Ttox vaccine) produced high titers of anti MUC1 IgG antibodies. These IgG MUC1 antibodies could specifically differentiate between human normal and PC tumor cells [20]. A clinical trial (NCT03114631) with DCs pulsed with MUC1/WT-1 peptides is currently under investigation for treating both resectable and unresectable PC patients.

2.2 Limitations of Mucin Vaccines for PC Immunotherapy

Efficacious MUC cancer vaccines have been a challenge to develop because tumor antigens like MUC1 are also expressed by normal cells (self-antigens) causing the immune system to develop self-tolerance against them, which leads to

hyporesponsiveness of APCs upon exposure to MUC1 cancer vaccines. Efficient activation of DCs is instrumental to activate cellular and humoral responses in an antigen-specific phenotype. In a very detailed study done in MUC1.Tg mice, it was observed that MUC1p cancer vaccine couldn't successfully activate APCs post vaccination in MUC1 expressing MUC1.Tg mice and was unable to restrict the growth of both transplanted and spontaneous PC tumors [21]. Whereas in WT mice, MUC1p cancer vaccination activated DCs and induced significant expression of surface markers like MHC-II, CD80, CD86 and CD40. Upon further investigation they discovered that MUC1.Tg immunized mice had a higher percentage of Fox3⁺ T_{regs} cells and these T_{regs} inhibited the expression of pancreatic enzymes such as trypsin and CBP1 transcript in DCs recovered from spleens of MUC1.Tg mice. Specific depletion of T_{regs} by anti-CD25 treatment or blocking of IL-10 by anti-IL10R antibody prior to MUC1p cancer vaccination rescued the expression of these enzymes in DCs to the similar levels of DCs isolated from WT mice [21]. This study suggests the necessity of discovering TAAs that are distinct from their counterparts being expressed on normal cells to prevent the formation of self-tolerance against them. A strategy to recognize MUCs against which self-tolerance is likely compromised in naïve PC patients who haven't received any treatments or surgery, could enable identifying the immunogenic MUC candidate that will probably overcome the hyporesponsiveness of APCs and generate corresponding anti-tumor responses.

Another limitation of MUC vaccines is the number of available epitopes to activate APCs. Peptides from VNTR region of MUC proteins have been utilized to develop cancer vaccines like peptide-toxin vaccines [20] or DC vaccines [19]. But these vaccines have achieved only limited success in clinical trials. Peptide-based vaccines have certain limitations such as peptides are mostly designed to have an MHC-I restriction that leads to the presentation of individual peptides on only certain HLA types. In addition, if selected

peptides have low affinity for MHC, they may be poorly immunogenic and induce weak or transient immune responses by APCs upon exposure [22]. As previously discussed in our review [18], protein provides an array of epitopes having multiple immunogenicities and thus could be presented by different MHC-I HLA types, potentially leading to the generation of strong immune responses. Expression, isolation and purification of transmembrane MUC glycoproteins come with its challenges. Due to the high molecular weight of MUCs, their purification and maintenance of their native antigenicity have not been feasible to date.

3. MUC4 as Tumor-Associated Antigen for PC

3.1 MUC4 in Pancreatic Cancer

In contrast to MUC1, MUC4 expression in normal organs is mostly restricted and is undetectable in normal pancreas and inflammatory diseases of the pancreas [23, 24] thus providing a better tumor specificity for targeting purposes. MUC4 is one of the most differentially overexpressed multi-domain transmembrane glycoprotein in PC [2, 7, 23, 25]. MUC4 protein carries a high percentage of allelic polymorphism in the VNTR domain that is rich in serine, threonine, and proline residues and it extensively undergoes mucin-type O-linked glycosylation [2, 23, 26] thus contributing to the high molecular weight of the apoprotein. MUC4 has a putative cleavage site comprising of Glycine-Aspartate-Proline-Histidine (GDPH) that can undergo autocatalysis and generate two subunits: MUC4 α upstream of GDPH cleavage site and MUC4 β downstream of the site [26-28]. MUC4 α subunit is the large N-terminal domain that is composed of VNTR domain, nidogen-like domain (NIDO), and adhesion-associated domain in MUC4 and other proteins (AMOP). MUC4 β on the other hand is a smaller subunit composed of von Willebrand factor type D

domain (vWD), three epidermal growth factor (EGF)-like domains, a transmembrane domain and a short cytoplasmic tail [23].

MUC4 has been demonstrated to play a fundamental role in the pathobiology of PC. Over the years studies have shown that MUC4's interaction with EGFR family members, HER2 and HER3 is mediated by the 3 EGF-like domains present in the MUC4 β subunit [29-31]. These interactions subsequently activate an intracellular cascade of signaling events including mitogen-activated protein kinases (MAPK), c-Jun N-terminal kinases (JNK), and signal transducer and activator of transcription 1 (STAT-1) that promote cell proliferation, migration and metastasis [12, 29-32]. The NIDO domain on MUC4 interacts with fibulin-2 that competitively disrupts its interaction with extracellular matrix (ECM) proteins present in the basement membrane, thus abrogating normal ECM protein-protein interactions [33]. An overall survey of the literature shows that MUC4 can modulate diverse pathways such as drug resistance [34], epithelial to mesenchymal transition (EMT) and metastasis [35], tumor cell proliferation and invasion [12], and thus PC patients with a MUC4 expression on tumors have a worse prognosis [36]. Given the tumor-promoting nature and PC-specific expression of MUC4, various studies have focused on utilizing MUC4 as a diagnostic/prognostic marker. MUC4 has emerged as a useful diagnostic tool for PC in fine needle aspirates (FNAs) [37, 38]. Also studies with patient serum samples for MUC4 serving as biomarkers have been investigated [39, 40]. In summary, MUC4 has been undisputedly established as a PC tumor-specific molecule that plays an instrumental role in progression and metastasis of PC disease, as well as serves as a useful diagnostic marker.

3.2 Immune Regulation of MUC4 in Pancreatic Cancer

Multiple signaling pathways regulate MUC4 expression in PC through the binding of different transcription factors at the MUC4 promoter site [41, 42]. Previous studies have

shown cytokine-mediated expression of MUC4 in PC cells. IFN γ and retinoic acid (RA) synergistically upregulates the expression of MUC4 in PC cells through the dual activation of STAT-1 and transforming growth factor beta-2 (TGF β) pathways [43, 44]. TGF β can upregulate MUC4 expression either by cooperative activation of Smad3 and Smad4 signaling pathways, or by activating MAPK, phosphatidylinositide 3-kinases (PI3K) and protein kinase A (PKA) signaling pathways [45]. Further, in other cancers such as gastric and colon cancer cell lines, it has been observed that interleukin (IL)-4, IL9, IL6, & IL24 could upregulate MUC4 expression by activating JAK/STAT pathway [46, 47] or in a STAT-3 dependent manner [48, 49]. PC has an elaborate TME composed of multiple immune cells such as TAMs, MDSCs, M2 macrophages etc. that secrete Th2 cytokines such as IL4, IL9 and IL24 (an IL-10 cytokine family member) to establish and maintain immunosuppression [18]. In addition, aberrant glycosylation of MUC4 appears to induce MUC4-specific immune responses such as generation of both Th1 and Th2 cytokines in MUC4-expressing pancreatic tumors [44]. Cumulatively these studies demonstrate that immune cells in TME can regulate the expression of MUC4 on PC cells by secretion of cytokines, which reveals an unexplored relationship between MUC4 and PC immunosuppression.

4. Summary and Conclusion

PC immunotherapy faces some major challenges in the selection of a suitable antigen. It has to meet three criteria: tumor-specific expression, antigen availability due to sufficient expression, and high immunogenicity to elicit strong immune responses. Due to its deregulated overexpression in PC and functional role in pathogenesis, MUC4 has emerged as a promising diagnostic and therapeutic candidate. MUC4's central role in eliciting tumor proliferation, invasiveness, drug-resistance and metastasis, has led to considerable interest in targeting MUC4 to avail therapeutic benefits. Further, due to

aberrant glycosylation [50] and cleavage [26-28], MUC4 expressed on PC tumor cells will be distinct from the expression on normal tissues. Circulating autoantibodies to MUC4 has been detected in colorectal patients [51] and preliminary study with MUC4 peptide-induced DNA vaccine showed that MUC4 DC vaccine could efficiently present the antigen and elicit potent MUC4-specific CTL response [52].

Further, the structural complexity and various glycoforms of MUC4 adds a challenge in designing strategies for immunotherapy. Previous studies have been focused on peptides located in the VNTR region. Since MUC4 could putatively get cleaved at the GDPH site into alpha and beta subunits, thus there is a possibility of MUC4 α (containing VNTR domain) to be not attached to the cell surface of tumor cells because MUC4 fragments have been discovered in secretions [53, 54].

In conclusion, MUC4 is a cancer-specific tumor-promoting glycoprotein that could serve as an effective antigen to develop vaccines against PC. However, investigations exploring MUC4 as a candidate for immunotherapy are in infancy. It requires multiple optimizations like identifying optimal biodegradable adjuvant delivery systems, identification of helper epitopes and developing platforms with good adjuvant properties that conserve the antigenicity of the epitopes. Thus, further studies are necessary to establish MUC4 as a suitable target for PC immunotherapy.

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**CHAPTER 1C: DISSERTATION GENERAL
HYPOTHESIS AND OBJECTIVES**

1. Background and Rationale

PC is the third leading cause of cancer-related death in the U.S. claiming 45,000 lives every year because of its poor prognosis and resistance to conventional therapies [1]. Much like other solid tumors, PC evades immune surveillance by manipulating immune cells to establish an immunosuppressive TME. Immunotherapy has emerged as an alternative approach for PC immunotherapy. It reprograms the patient's immune system to selectively target and kill cancer cells thus reducing the non-specific side-effects. PC immunotherapy faces multiple challenges such as: i) identification of tumor-associated antigen that could be targeted; ii) preservation of TAA antigenicity and its sustained delivery; iii) generation of robust anti-tumor responses; and iv) overcoming immunosuppression on effector immune cells in PC tumor microenvironment.

PC harbor limited genetic alterations which render it as an immunologically 'cold' tumor. Overexpression and altered glycosylation of mucins in tumor cells can trigger humoral and cellular immune responses. Among various mucins, MUC4, heterodimeric transmembrane mucin is aberrantly overexpressed in PC but is undetectable in normal pancreas and is associated with poor prognosis in PC patients [2]. In PC, MUC4 is aberrantly glycosylated, and thus could serve as a potential source of neoantigenic epitopes, which could be exploited for cancer immunotherapy. Additionally, due to the transmembrane nature of MUC4 only a very limited amount is released into the blood, and that makes MUC4 a very strong candidate for vaccine [3-6]. A survey of literature has shown that mucins like MUC1 have been targeted by immunotherapy approaches for PC treatment but have not fared well in clinical trials. Further detection of circulating serum autoantibodies to MUC4 has shown its potential to be immunogenic in colorectal cancer patients [7]. However, MUC4 immunogenicity in PC patients has yet not been studied.

These summarized findings suggest that MUC4 has potential to be a candidate for PC immunotherapy.

Polyanhydride nanoparticles represent an ideal vaccine adjuvant/delivery platform. Polyanhydride nanoparticles have been demonstrated to provide sustained release and enhanced stability of encapsulated antigens (peptides, proteins etc.) [8]. The ratio of polyanhydride nanoparticle's formulation provides an immune modulation property that could be utilized to tune the immune responses to either Th1 or Th2 phenotype [9, 10]. In addition, their adjuvant properties enhance antigen internalization by antigen-presenting cells and stimulate both antibody- and cell-mediated immunity [11-13]. Due to their pathogen-mimicking, immune-modulation and biodegradable nature, the polyanhydride nanoparticles platform could be utilized to develop vaccines for PC immunotherapy.

2. Hypothesis

Based on previous studies demonstrating the tumor-specific overexpression of MUC4 in PC and suitable qualities of polyanhydride nanoparticles to serve as a vaccine delivery platform, we *hypothesized* that i) MUC4 is immunogenic in PC patients, and ii) delivery of encapsulated MUC4 in unique amphiphilic polyanhydride nanovaccine will elicit robust immune responses in an antigen-specific manner.

3. Objectives

Aim 1: To determine whether tolerance will be broken against MUC4 in pancreatic cancer patients and elucidating it as a suitable target it for immunotherapy studies.

Aim 2: To evaluate whether MUC4 nanovaccine can elicit robust immune responses.

Aim 3: To evaluate whether MUC4 nanovaccine can elicit antigen-specific anti-tumor responses in PC-subcutaneous tumor-bearing mouse model.

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CHAPTER 2:

MATERIALS AND METHODS

1. Human Serum Sample Description

Human sera were obtained after written informed consent was acquired from individuals {the protocol was approved by the UPMC Review Board (IRB number PRO07030072), and UNMC Review Board (IRB number 209–00)} from the University of Pittsburgh Medical Center (UPMC) and shipped to UNMC. Patients with benign pathologies or abnormal imaging of pancreas were categorized as Healthy; PC staging was determined surgically by pathologists; chronic pancreatitis (CP) was defined based on standard clinical practices at UPMC. Sera from PC patients were collected close to the time of the first diagnosis of cancer, prior to surgery and before chemotherapy treatment. All samples collected at UPMC were shipped by overnight mail to the UNMC. After receiving the samples, they were stored following clotting of sera for 30 min, aliquoted and immediately frozen at -70°C within 60 min of collection. The sample set was comprised of 21 CP patients, 27 Healthy, and 31 PC patients, which could be divided among 16 early-stage patients (Stage I–II) and 15 late-stage patients (Stage III–IV) who were recruited at UPMC. To evaluate the presence of autoantibodies against MUC4 peptides, randomized patient sera was selected to make a cohort of 10 Healthy, 10 CP patients and 22 PC patients. Details on the groups of patients are provided in the supplementary section, and patient demographic data are described in **Supplementary Table 1 & 2**.

2. Prediction of MUC4 Immunogenic Peptides by Bioinformatics

MUC4 immunogenic peptides were predicted by multiple software: T-cell epitopes for human MHC-I (most common: HLA-A*-02:01) were predicted and scored by Immune Epitope Database and Analysis Resource (IEDB) [24-31] and TepiTool [32], and combined human HLA isoforms scores were predicted by NetCTL-1.0 [33] online prediction tools. Human MUC4 peptides that could be loaded on C57BL/6 mouse MHC-I & MHC-II complexes were predicted and scored by IEDB online prediction tool; and linear B cell

epitopes were predicted and scored by IEDB and BCPred [34] online prediction tools. Lowest percentile rank scoring peptides were selected to be synthesized for our study. For our study, we selected the amino acid sequence derived from the MUC4 sequence isolated and characterized at www.uniprot.org (UniProtKB/Swiss-Prot: Q99102.4) [20, 22, 35, 36].

3. MUC4 Peptide and MUC4 β Purification

Predicted MUC4 9-mer-amino-acid peptides, MUC4 VNTR peptide (TR) and pancreatic differentiation 2 (PD2) peptide used in this study were locally synthesized and purified at UNMC. Seven peptides from the randomized regions of MUC4 α (excluding the TR region) and four peptides from the MUC4 β region (upstream of the transmembrane domain) were used to detect autoantibodies in PC patients, CP patients and Healthy sera.

Recombinant human MUC4 β protein was expressed in *E. coli* R-2 (DE3) strain transformed with MUC4 β -6-His-Tag expression plasmid and purified by AKTA Ni-NTA affinity chromatography. Eluted protein fractions were assessed by Coomassie Blue stained 10% SDS-PAGE gels and by immunoblotting using anti-His tag antibody (Thermo Fisher, Pierce # 31430), confirming the presence of MUC4 β fractions. MUC4 β fractions were concentrated by using Amicon Ultra centrifuge filters and further purified by dialysis in ultra-purified endotoxin-free water. Purified MUC4 β fractions were passed through an endotoxin removal spin column (Thermo Fisher, Pierce # 88277) and the final endotoxin level measured by Pierce LAL Chromogenic Endotoxin assay kit (Pierce, # 88282) was less than <1. The purified recombinant MUC4 β protein fraction was quantified by BCA protein assay kit (Thermo Fisher, Pierce # 23225) for the study.

4. Encapsulation of MUC4 β in Polyanhydride Nanoparticles

The 20:80 CPTEG:CPH polymer was synthesized via melt polycondensation [23]. Next, nanoparticles encapsulating MUC4 β were synthesized using a solid-oil-oil double emulsion technique previously described in [28]. Briefly, purified MUC4 β was dialyzed to nanopure water and lyophilized. Next, 20:80 CPTEG: CPH polymer containing three wt.% MUC4 β was dissolved 20 mg/mL in methylene chloride. The solution was sonicated for 30 s to ensure even distribution of the protein. The nanoparticles were precipitated into chilled pentane (-10°C; 1:250 methylene chloride: pentane) and collected via vacuum filtration. Nanoparticle morphology was verified by scanning electron microscopy (FEI Quanta 250, FEI, Hillsboro, OR) and their size subsequently analyzed with ImageJ (ImageJ 1.48v, NIH). The release kinetics of MUC4 β from 20:80 CPTEG: CPH nanoparticles were monitored as previously described [22]. Briefly, nanoparticles were incubated in PBS at 37°C. Periodically, the samples were centrifuged, supernatant collected, and particles were resuspended in fresh buffer. The amount of protein in the collected supernatant was quantified using a microBCA assay. At the end of approximately one month, the buffer was exchanged with 40 mM sodium hydroxide to quickly degrade the nanoparticles and release any remaining protein. The encapsulation efficiency was determined by comparing the total amount of protein released to the amount theoretically encapsulated.

5. Primary Dendritic cells (DCs) Isolation

C57BL/6 mice were kept under SPF conditions at UNMC animal facilities in accordance with UNMC Institutional Animal Care and Use Committee (IACUC) standards. Femurs and tibiae of female, 6-8 weeks old female C57BL/6 mice were removed and purified from the surrounding muscle tissue by rubbing with kleenex tissues. Thereafter intact bones were disinfected in phosphate buffered saline pH 7.2 (PBS) containing

penicillin (100 U/mL, Sigma) and streptomycin (100 mg/mL, Sigma) solution for 2-5 min. Then both ends were cut with scissors and the marrow was flushed with PBS using a 1 mL syringe attached to 25 G needle. Homogenous pipetting disintegrated clusters within the marrow suspension. Cells were centrifuged at 2000 rpm for 2.5 min and the cell pellet was resuspended in 10 mL of 1X RBC lysis buffer in the dark for 5 min. Ten mL of RPMI was added to stop the lysis. Cells were centrifuged at 2000 rpm and washed 3 times in 10 mL of RPMI. After the last wash, cells were resuspended homogeneously in a 1 bone:1 mL of media ratio. Resuspended cells were passed through a single cell strainer to remove clumps. Single cell suspensions were poured in 10 mL RPMI containing 10% FBS, penicillin (100 U/mL, Sigma) and streptomycin (100 mg/mL, Sigma) media in a 10 cm tissue culture plate that was kept in an incubator for 3 h. After incubation, we collected all floating cells and washed the plate with media twice to collect rest of the suspended cells in a 50 mL falcon tube. The cells were pelleted by centrifugation and the media aspirated. The cells were resuspended in 10 mL RPMI media and transferred into T75 flasks for our *in vitro* studies.

6. DC Maturation and Pulsing

To enhance the DC population, we added 100 ng/ μ l of rmGM-CSF and 50 ng/ μ l rmIL-4 reconstituted and diluted in serum-free RPMI to freshly isolated DC cultures on Day 0. On Day 3, 5, and 7, we collected the nonadherent cells in 50 ml centrifuge tubes, pelleted them and resuspended them in a total volume of 10 ml having 75% of fresh media containing 10% FBS + antibiotics and 100 ng/ μ l of GM-CSF and 50 ng/ μ l IL-4 in a fresh T75 flask. At day 9, immature DCs were counted and seeded in 24 well plate for activation studies. Polyanhydride nanoparticles were suspended in complete culture medium, sonicated briefly (30s on ice), and added to the DC cultures at day 9 at a concentration of 100 μ g/mL. DCs were pulsed in the following groups: 3 μ g/ml free MUC4 β protein

(MUC4), free MUC4 β protein (3 μ g/ml) mixed with blank nanoparticles (100 μ g/mL) (MUC4+NP) and MUC4 nanovaccine (100 μ g/mL). Unstimulated DCs (US) and DCs stimulated with LPS (200 ng/mL) were used as negative and positive controls, respectively. Cultures were incubated for 48 h (37 °C, 5% CO₂). Activated dendritic cells were harvested from 24-well plate and centrifuged at 1000 x g for 5 minutes to collect the culture medium and the pellet was processed further for flow cytometry studies. Supernatant was collected for ELISA studies from each treatment groups.

7. Flow Cytometry of Activated DCs

Dendritic cells were resuspended and washed 3 times in FACS buffer {1X PBS (pH 7.2) + 1% Fetal bovine serum (FBS)} to remove any residual culture medium. DCs were fixed in 4% paraformaldehyde for 15 min at RT and washed by centrifugation with excess FACS Buffer. After that, the 1X10⁵ DCs were resuspended in 100 μ l volume of conjugated antibody cocktail for detection of DC surface markers consisting of CD11c, MHC-I, MHC-II, CD40, CD80, CD86, and CD205 was prepared at 1:300 dilution in FACS Buffer. Corresponding isotype controls were also prepared at 1:300 dilution in FACS buffer. The washed DCs were suspended in either DC surface markers antibody cocktail or in isotype-antibody cocktail and incubated for 60 min on ice (4⁰C) in the dark. The labelled cells were again washed and analyzed using a BD LSR-II Green Flow Cytometer (BD Biosciences). FlowJo® and BD FACSDIVA software were used to analyze the data.

8. Cytokine Analysis by ELISA

Supernatants were preserved at -80⁰C and thawed on ice for cytokine analysis. IL-6, IFN- γ and IL-12/IL-23p40 cytokines in supernatants were measured by enzyme-linked immunosorbent assay (ELISA) kits from BioLegend and the manufacturer's protocol was followed for the assay. 96-well ELISA strips were coated with capture antibodies (1:200 dilution) diluted in carbonate-bicarbonate buffer (0.5M, pH 9.6) and incubated overnight

at 4°C. The next day coated strips were washed 4 times with 1X PBST (0.05% Tween 20) and blocked with 3% bovine serum albumin (BSA) in PBS for 2 hours at 37°C. Strips were then washed 4 times and 100 µl of supernatants were added to coated strips and incubated for 2 hours at 37°C. Plates were washed with PBST for 4 times, followed by incubation with detection antibody (1:200) for 1 hour at 37°C. Secondary antibody was washed away with 4 PBST washes. Avidin (1:1000) was added to ELISA strips and incubated for 30 min at RT in dark. Excess Avidin was washed away with 5 PBST washes, followed by addition of TMB (3,3',5,5'-Tetramethylbenzidine) substrate in the dark and incubated at RT for color to develop. Absorbance was measured after the reaction was stopped (~15 mins) with 1N H₂SO₄ at 450 nm using a SpectraMax® Plus384 microplate reader (Molecular Devices LLC, Sunnyvale, California). For serum cytokines analysis, serum samples were collected post-animal sacrifice and stored at -80°C. Serum samples were added at 1:10 dilution to capture-antibody coated plates for cytokine analysis. IL-6, IFN-γ and IL-12/IL-23p40 were measured by ELISA kits from BioLegend using a similar manufacturer's protocol.

9. Mice Immunization for Anti-MUC4 Antibodies Generation

Eight weeks C57BL/6 mice were immunized with various formulations of recombinant MUC4β protein, including protein alone (20 µg/mouse/dose), encapsulated MUC4β protein (300 µg/mouse/dose), protein plus nanoparticle, and saline control, constituted in Freund's adjuvant. First booster dose was given after 2 weeks of primary immunization. The blood was withdrawn from submandibular vein after 1 week of a single booster dose. The serum was isolated from the blood after overnight storage at 4°C and stored at -20 °C for further use.

10. MUC4 Autoantibody ELISA

To detect circulating MUC4 autoantibodies in sera of the sample set, a novel modified sandwich ELISA-assay was developed. Briefly each 96-well plate was coated with five $\mu\text{g/mL}$ of recombinant MUC4 β protein or with one $\mu\text{g/mL}$ of MUC4 peptides or control peptides (PD2 & TR) in carbonate-bicarbonate coating buffer (0.5 M, pH 9.6) and incubated overnight at 4 $^{\circ}\text{C}$. The plate was washed two times with 1X PBST and blocked with 3% BSA in PBS for 3 hours at 37 $^{\circ}\text{C}$. Further, the plate was washed four times and then a serial dilution (1:16000) of the serum samples were added to MUC4 β -coated plate and a serial dilution of 1:2000 was added to MUC4 peptide-coated plates. ELISA plates with primary serum dilutions were incubated overnight at 4 $^{\circ}\text{C}$. Plates were washed with PBST for four times, followed by incubation with horseradish-peroxidase (HRP) conjugated goat anti-human IgM+ IgG+ IgA (H+L) (Jackson ImmunoResearch), or for isotyping HRP conjugated IgM, IgG (H+L) and IgA (Jackson ImmunoResearch) respectively for 1 hour at 37 $^{\circ}\text{C}$. The excess secondary antibody was washed away with 5 PBST washes and followed by addition of TMB substrate in the dark. Absorbance was measured after the reaction was stopped (~15mins) by adding 1N H_2SO_4 at 450 nm using a SpectraMax[®] Plus384 microplate reader (Molecular Devices LLC, Sunnyvale, California).

11. Anti-MUC4 β Antibodies Detection in Immunized Mice Serum Using ELISA

Serum samples were collected post-animal sacrifice and stored at -80 $^{\circ}\text{C}$ for detection of anti-MUC4 β antibodies using ELISA techniques. To detect anti-MUC4 β antibodies, we used a modified ELISA protocol. A 96-well plate was coated with 5 $\mu\text{g/mL}$ of recombinant MUC4 β protein in carbonate-bicarbonate coating buffer (0.5 M, pH 9.6) and incubated

overnight at 4°C. Plate was washed 2 times with 1X PBST and blocked with 3% BSA in PBS for 3 hours at 37°C. Further, the plate was washed 4 times and then serial dilutions of the serum samples were incubated in MUC4 β coated plate for 2 hours at 37°C. Plates were washed with PBST for 4 times, followed by incubation with horseradish-peroxidase (HRP) conjugated goat anti-mouse IgG (total H+L) (Thermo Fisher), IgG1 (Abcam) and IgG2b (Abcam) for 1 hour at 37°C. Excess secondary antibody was washed away with 5 PBST washes and followed by addition of TMB substrate in the dark. Absorbance was measured after the reaction was stopped (~15 min) with 1N H₂SO₄ at 450 nm using a SpectraMax[®] Plus384 microplate reader (Molecular Devices LLC, Sunnyvale, California).

12. Generation of MUC4-Expressing Murine PC Cell Lines

A MUC4-expressing mouse pancreatic cancer cell line derived from KPC mice (KPC960) was developed by transfection of KPC960 cell lines with miniMUC4-pSecTagC plasmid using the Lipofectamine method (Invitrogen) and single colonies were obtained by zeocin selection as published previously in Moniaux *et.al* 2007 and confirmed by immunoblotting with a MUC4 peptide mouse monoclonal antibody (8G7). The miniMUC4 expressing KPC960 cells were kindly developed, validated and gifted by Dr. Shailendra Gautam for our *in vivo* and *ex vivo* studies.

13. Tumor Implantation and Mice Immunization

For *in vivo* characterization of MUC4 nanovaccine in mice, 8 weeks C57BL/6-FBP mixed background mice were immunized with different formulations of recombinant MUC4 β protein, including protein alone (MUC4- 35 μ g/mouse/dose), free MUC4 protein mixed with nanoparticles (MUC4+ NP) (35 μ g/mouse/dose of MUC4 + 500 μ g/mouse/dose empty nanoparticles), MUC4 nanovaccine (500 μ g/mouse/dose), and unimmunized mice (negative control). First booster dose was given after two weeks of primary immunization.

Two weeks after first booster, mice were injected with 1×10^6 cells of mini-Muc4 or Vector expressing KPC derived mouse pancreatic cell lines(KPC960) into right and left flanks of mixed background mice (n=5), respectively. Tumor growth was monitored till the tumor volume reached to 100mm^3 volume. Mice were then immunized with second booster dose but the MUC4 protein dose was reduced to $20 \mu\text{g}/\text{mouse}/\text{dose}$ in all treatment groups except for saline control. Tumors were harvested at 23rd-day post second booster immunization and preserved in formalin for further use.

14. T-cell Isolation and *In Vitro* Activation

Naïve T-cells were isolated from 8-10weeks old C57BL/6-FBP mixed background mice. Mice were sacrificed, and spleens were isolated in RPMI+10%FBS+Pen-Strp (RPMI) medium in 50 ml Falcon tube on ice. Media was removed by suction inside the laminar hood and one spleen was added to one of the wells of 6 well plates. Spleens were teased with a 25G syringe needle and 1 ml pipette tip. Teased-out splenocytes were transferred to 15 ml falcon tube in RPMI media and was centrifuged for 2-3mins at 2000 rpm. The supernatant was removed and the pellet was resuspended in 10-12ml of 1X RBC lysis buffer, then the tube was kept for 5 min in the dark. The tube was centrifuged again for 2-3 min at 2000 rpm and the supernatant was aspirated. The pellet was resuspended and splenocytes were washed for 3 times with RPMI media. After the last wash, the pellet was resuspended in 2 ml of RPMI media and counted total cells/ml. We centrifuged the cell suspension at 2000 rpm for 10 min and removed the supernatant completely. We resuspended the cell pellet in $40 \mu\text{L}$ of MACs Buffer (1X PBS pH 7.2, 0.5% BSA, 2mM EDTA) per 10^7 cells. We added $10 \mu\text{L}$ of Pan T cell Biotin-Antibody Cocktail (MACS Miltenyi Biotech, Inc.) per 10^7 cells and mixed it well. We incubated the mixture for 5 min in the refrigerator ($2-8^\circ \text{C}$) and then added $30 \mu\text{L}$ of MACs buffer per 10^7 cells, and followed by addition of $40 \mu\text{L}$ of Anti-Biotin Microbeads per 10^7 cells. We mixed all the

components well & incubated for additional 10 min in the refrigerator (2-8° C). We took the mixture and processed the sample in an autoMACS® Pro Separator (MACS Miltenyi Biotech, Inc.). Tubes were placed in the following Chill Rack positions: position A = sample, position B = negative fraction position C = positive fraction. We prepared and primed the instrument and followed the instructions that were given in the user manual. T cells were isolated by magnetic separation and we used the 'Depletes' program to do the separation into positive (other splenocytes) and negative (T cells) population.

Naïve T-cells were then cultured and expanded in RPMI on day 0. On day 1, purified T cells were incubated with phorbol 12-myristate 13-acetate (PMA) (25 ng/mL) Ionomycin (500 ng/mL, Abcam) for 48 hours at 37°C. On day 3, these T-cells were co-cultured in 12-well plate with 48hour-pulsed DCs in the following groups: 3 µg/ml free MUC4β protein (MUC4), free MUC4β protein (3 µg/ml) mixed with blank nanoparticles (100 µg/mL) (MUC4+NP) and MUC4 nanovaccine (100 µg/mL). Unstimulated DCs (US) and DCs stimulated with lipopolysaccharides {LPS} (200 ng/mL) were used as negative and positive controls, respectively. T-cells were co-cultured for 48 hours with pulsed DCs and then collected for further analysis. The experiment was done in triplicates.

Since the Pan T cell Isolation kit can purify both naïve and activated T cells, a similar T-cell isolation protocol was utilized to isolate T-cells from immunized mice and expanded *in vitro* by treatment with PMA/Ionomycin (25 ng/mL/500ng/mL, Abcam) for 48 hours at 37°C. On day 3, the expanded T-cells were analyzed by flow cytometry for phenotyping T-cells and cytokine secretion studies.

15. Cytotoxicity Assay of T-cells

The miniMUC4 KPC960 cells and KPC960 vector control cells (500 cells/well) were seeded in 96-U bottom plates and cultured overnight in a 37°C incubator. The plate

was centrifuged and the media was removed. Activated T-cells were added from each of the treatment groups to target cancer cells at 10:1 dilution into their respectively labeled wells. T-cell mediated cytotoxicity was measured by the CytoTox 96® Non-Radioactive Cytotoxicity Assay Kit (Promega) and the manufacturer's protocol was followed for the assay. The plate was centrifuged at 250 × g for 4 minutes and the plate was incubated at 37°C for 4 hours. At 45 minutes prior to supernatant harvest, Lysis Solution was added to wells of both cancer cell lines that didn't have T-cells added to them, which served as our Target Cell Maximum LDH Release Control. The plate was centrifuged at 250 × g for 4 minutes and 50 µl of the supernatant from each well of the assay plate was transferred to the corresponding well of a flat-bottom 96-well enzymatic assay plate. Substrate Mix was reconstituted using Assay Buffer and 50 µl of the reconstituted Substrate Mix was added to each well of the plate. The plate was incubated at room temperature in the dark for 30 min. Absorbance was measured after the reaction was stopped with 50 µl of the Stop Solution at 450 nm using a SpectraMax® Plus384 microplate reader (Molecular Devices LLC, Sunnyvale, California).

16. Flow cytometry of Activated T-cells

T-cells isolated from immunized mice were expanded by PMA + Ionomycin treatment. Brefeldin A (2 µg/mL) was added to the treated flask and untreated flask and cells were incubated for 4 hours in a 37°C incubator. T-cells were resuspended and washed three times in FACS buffer {1X PBS (pH 7.2) + 1% Fetal bovine serum (FBS)} to remove any residual culture medium. After that, the 1×10^5 T-cells were resuspended in 100 µl volume of conjugated antibody cocktail for detection of T-cell surface markers (consisting of Th1 phenotype CD4, CD8, and Tbet) which was prepared at 1:300 dilution in FACS buffer. Corresponding isotype controls were also prepared at 1:300 dilution in FACS buffer. The washed T-cells were suspended in either DC surface marker's antibody

cocktail or in isotype-antibody cocktail and incubated for 60 min on ice (4°C) in the dark. The labelled cells were again washed with FACs buffer. T-cells labelled with surface antibodies were fixed in 4% paraformaldehyde for 15 min at RT and washed by centrifugation with excess FACs buffer. For intracellular labeling of cytokines, antibodies and their corresponding isotype controls were prepared in pre-chilled Phosflow Buffer™ (BD Biosciences) at 1:300 dilution and keep it on ice. Fixed T-cells were centrifuged, and 4% paraformaldehyde was discarded by gentle flicking and washed one time with FACs buffer. Fixed T-cells were permeabilized by adding 100 ul of PhosFlow Buffer™ in the dark and on ice for 30 min. T-cells were centrifuged and the permeabilization buffer was decanted. For intracellular cytokine staining, 100 ul of IL2, IL12, TNF α and IFN γ (eBioscience) antibody cocktail or isotype control antibody cocktail added to T-cells and incubated for 60 min on ice (4°C) in the dark. The labelled cells were again washed with FACs buffer and analyzed using a BD LSR-II Green Flow Cytometer (BD Biosciences). FlowJo® and BD FACSDIVA software were used to analyze the data.

17. Immunohistochemistry and Immunofluorescence staining of tumor tissue sections

The miniMUC4-expressing and vector control murine subcutaneous tumor sections were evaluated for PD-L1 expression by immunohistochemistry (IHC) and for CD4 and CD8 T-cell's infiltration by immunofluorescence (IF). Harvested tumors were fixed in 10% neutral buffered formalin, embedded in paraffin, and 4 μ m thick sections were placed on slides. After deparaffinizing with xylene and rehydrating with decreasing alcohol gradients with final rehydration in MilliQ water, for IHC staining, tumor sections were treated with 5% H₂O₂ in 100% methanol for 1 hour at room temperature in the dark for quenching of endogenous peroxidases. For IF, tumor sections were treated with 100% methanol for 1 hour at room temperature in the dark for fixing the tissue. Antigen retrieval

was done using Tris-EDTA buffer, pH 9.0 in the microwave (1100W) at high power for 15 min. For simple hematoxylin and eosin staining of tumor tissue sections, tissue slides were counterstained with hematoxylin and eosin, followed by dehydration with increasing alcohol gradients, xylene washes, and mounting with a cover slip. All hematoxylin and eosin-stained slides were shown to the pathologist for analysis and quantification.

Tissues were blocked with 2.5% normal horse serum for 3 hours, and then incubated with a primary anti-mouse PD-L1 antibody (cc-50298, Santa Cruz Biotechnology Inc.) at 1:200 dilution in Tris-buffered saline (TBS) at 4°C overnight. For T-cell's IF staining, two separate tissue sections were incubated with anti-mouse CD4 or CD8 antibodies (Thermo Fisher Scientific) respectively at 1:400 in PBS at 4°C overnight. PD-L1 stained slides were rinsed with TBST and T-cells stained slides were washed with PBST. After wash, PD-L1 labeled slides were incubated with an HRP-conjugated goat anti-rabbit secondary antibody for 1 hour. T-cell's labeled sections were incubated with Alexa Fluor 488 conjugated goat-anti-mouse secondary antibody for 1 hour in the dark. IHC slides were rinsed with 2 washes of TBST and 3 washes of TBS, and subsequently, using DAB substrate kit (Vector Laboratories) the color was developed. Sections were counterstained with hematoxylin, followed by dehydration with increasing alcohol gradients, then xylene washes and mounting with a cover slip. IF slides were again washed with 4 washes with PBST and two washes with PBS, followed by covering the section with a glass slide and the anti-fade VECTASHIELD mounting media (Vector Laboratories Inc., Burlingame, CA, USA). IF staining was observed under the Zeiss 510 LASER SCAN confocal microscope and quantified by ImageJ software (<https://imagej.nih.gov/>).

18. Statistical Analysis

For our autoantibodies analysis, the differences among group means were tested by one-way analysis of variance (ANOVA) F-test using JMP® Data Analysis Software (SAS Institute Inc., NC). If the F-test was significant, paired Wilcoxon sign rank t-tests were performed for pairwise comparisons of group means. Significance was defined as $p < 0.001$.

For our nanovaccine *in vitro*, *ex vivo* and *in vivo* studies differences among group means were tested by one-way analysis of variance (ANOVA) F-test using GraphPad Prism v. 7.0 (GraphPad, La Jolla, CA). If the F-test was significant, Tukey's t-tests and Student's t-tests were performed for pairwise comparisons of group means. Significance was defined as $p < 0.05$.

19. Table I: List of antibodies

S.No.	Antibodies	Company	Catalogue No.
1	Alexa Fluor® 700 anti-mouse CD11c (clone N418)	eBioscience (San Diego, CA)	56-0114-80
2	FITC conjugated anti-mouse/rat MHC Class II (I-Ek, clone 14-4-4S)	eBioscience (San Diego, CA)	11-5980-81
3	(APC) anti-mouse CD40 (clone 1C10)	eBioscience (San Diego, CA)	17-0401-81
4	(PE)-Cy7 conjugated anti-mouse F4/80 (clone BM8)	eBioscience (San Diego, CA)	25-4801-82
5	Anti-Mouse MHC Class I (H-2Kd) eFluor® 450 (Clone: SF1-1.1.1)	eBioscience (San Diego, CA)	48-5957-80
6	Alexa Fluor® 700 conjugated Armenian hamster IgG (clone eBio299Arm)	eBioscience (San Diego, CA)	56-4888-80
7	FITC IgG2a κ (clone eBR2a)	eBioscience (San Diego, CA)	11-4321-80
8	Rat IgG2a K Isotype Control APC (clone eBR2a)	eBioscience (San Diego, CA)	17-4321-41 17-4321-81
9	PE-conjugated rat IgG2a (clone eBR2a)	eBioscience (San Diego, CA)	12-4321-80
10	Rat IgG2a K Isotype Control PE-Cyanine7 (clone eBR2a)	eBioscience (San Diego, CA)	25-4321-81
11	PE/Cy7 anti-mouse CD86 (Clone GL1)	BD Biosciences	560582
12	PE conjugated anti-mouse CD80 (clone 16-10A1)	eBioscience (San Diego, CA)	12-0801-81

13	PE/Cy 5.5 antimouse CD205 (MMR, clone NLDC-145)	BioLegend	138207
14	PerCP/Cy5.5 Rat IgG2a, κ Isotype Ctrl Antibody	BioLegend	400531
15	Rat IgG2a K Isotype Control eFluor® 450 (clone eBR2a)	eBioscience (San Diego, CA)	48-4321-80
16	T-bet (Anti-Human/Mouse T-bet PerCP-Cyanine5.5)	eBioscience (San Diego, CA)	45-5825-80
17	GATA-3 (Anti-Human/Mouse Gata-3 eFluor® 660)	eBioscience (San Diego, CA)	50-9966-41
18	CD4 (Anti-Mouse CD4 eFluor® 450)	eBioscience (San Diego, CA)	48-0042-80
19	CD8 (Anti-Mouse CD8a PE)	eBioscience (San Diego, CA)	12-0081-81
20	IL-2 (Anti-Mouse IL-2 PE-Cyanine7)	eBioscience (San Diego, CA)	25-7021-80
21	IL-10 (Anti-Mouse IL-10 FITC)	eBioscience (San Diego, CA)	11-7101-41
22	IL-12 (Anti-Mouse IL-12/IL-23 p40 Alexa Fluor® 488)	eBioscience (San Diego, CA)	53-7123-80
23	IL-13 (Anti-Mouse IL-13 PE-Cyanine7)	eBioscience (San Diego, CA)	25-7133-80
24	IFN γ (Anti-Mouse IFN gamma APC)	eBioscience (San Diego, CA)	17-7311-81
25	Rat IgG2b Isotype Control eFluor® 660	eBioscience (San Diego, CA)	50-4031-80
26	Rat IgG2a K Isotype Control Alexa Fluor® 488	eBioscience (San Diego, CA)	53-4321-80

27	Rat IgG2b K Isotype Control FITC	eBioscience (San Diego, CA)	11-4031-81
28	Rat IgG1 K Isotype Control eFluor® 450	eBioscience (San Diego, CA)	48-4301-80
29	CD4 Monoclonal Antibody (RIV6)	ThermoFisher Scientific	MA1-7631
30	CD8 Monoclonal Antibody (RIV11)	ThermoFisher Scientific	MA1-7632
31	Pdcd-1L1 (H-130) {Discontinued Antibody}	SantaCruz Biotechnology INC.	Sc-50298
32	Goat Anti-mouse IgG2b-HRP	ThermoFisher Scientific	M32407
33	Goat Anti-mouse IgG1-HRP	ThermoFisher Scientific	A10551
34	Peroxidase AffiniPure Goat Anti-Human IgM, Fc5µfragment specific	Jackson ImmunoResearch Laboratories, INC.	109-035-129
35	Peroxidase AffiniPure Goat Anti-Human IgG (H+L)	Jackson ImmunoResearch Laboratories, INC.	109-035-003
36	Peroxidase AffiniPure Goat Anti-Human IgA + IgG + IgM (H+L)	Jackson ImmunoResearch Laboratories, INC.	109-035-064
37	Peroxidase AffiniPure Goat Anti-Human Serum IgA, α Chain Specific	Jackson ImmunoResearch Laboratories, INC.	109-035-011
38	Goat anti-Mouse IgG (H+L) Cross-Adsorbed Secondary Antibody, Alexa Fluor 488	ThermoFisher Scientific	A-11001
39	Goat anti-Mouse IgG (H+L) Secondary Antibody, HRP	ThermoFisher Scientific	31430
40	Goat anti-Rabbit IgG (H+L) Secondary Antibody, HRP	ThermoFisher Scientific	31460

Supplemental Table 1. Patient characteristics, all subjects

		PC (n=22)	Healthy (n=27)	CP (n=21)	p-value
Age	Median (range)	71 (56-85)	64 (28-90)	54 (30-81)	0.0003
Gender	Female	9 (41%)	18 (67%)	8 (38%)	0.12
	Male	13 (59%)	9 (33%)	13 (62%)	
Race	African American	1 (5%)	1 (4%)	5 (24%)	0.090
	Caucasian	21 (95%)	26 (96%)	16 (76%)	
Stage	I/II	12 (55%)	-	-	
	III/IV	10 (45%)			

Supplemental Table 2. Patient characteristics, limited subjects

		PC (n=22)	Healthy (n=10)	CP (n=10)	p-value
Age	Median (range)	71 (56-85)	62 (33-86)	48.5 (30-81)	0.016
Gender	Female	9 (41%)	6 (60%)	5 (50%)	0.61
	Male	13 (59%)	4 (40%)	5 (50%)	
Race	African American	1 (5%)	0	2 (20%)	0.27
	Caucasian	21 (95%)	10 (100%)	8 (80%)	
Stage	I/II	12 (55%)	-	-	
	III/IV	10 (45%)			

CHAPTER 3: MUC4 AS A VACCINE CANDIDATE FOR PC IMMUNOTHERAPY

The material covered in this chapter is the subject of a research article

2. Banerjee K., Kshirsagar P., Kaur S., Brand RE., Smith L., Batra SK., and Jain M. *Presence of MUC4 autoantibodies in the serum of pancreatic cancer patients.* (Manuscript under Preparation)

1. Synopsis

A major challenge in pancreatic cancer (PC) immunotherapy is identifying tumor-associated-antigens (TAAs) that can be targeted. In PC, Mucin-4 (MUC4) is differentially overexpressed and aberrantly glycosylated, thus potentially serving as a source of neo-antigenic epitopes, which could be exploited for cancer immunotherapy. Literature review has shown that an immune response developed in the form of autoantibodies to various tumor antigens could serve as a surrogate for immunogenicity (and compromised self-tolerance) in cancer patients. The presence of MUC4 autoantibodies was analyzed in serum samples taken from patients at the time of pancreatic cancer diagnosis prior to any treatments, from patients with chronic pancreatitis, and healthy controls. By using indirect-ELISA techniques we were able to detect autoantibodies against recombinant human MUC4-beta (MUC4 β) protein in 77.41% of PC patient's sera, whereas only 33.33% of age-matched healthy controls and 23.81% of chronic pancreatitis (CP) patients had positive sera. In addition, we used a panel of 7 MUC4-alpha region peptides and 4 MUC4 β peptides to identify highly immunogenic MUC4 epitopes. Furthermore, isotyping these autoantibodies in PC patient sera indicates that IgM antibodies to peptides A2, D4 and B1 correlate with better prognosis. Our novel study suggests that MUC4 expressed by pancreatic tumors is immunogenic in PC patients and could be used as a target for cancer immunotherapy. Further these studies suggest that segregation of PC patients based on the presence of MUC4 autoantibodies against multiple MUC4 peptide, could be helpful in identifying patients for personalized MUC4-based immunotherapy treatment of pancreatic cancer.

2. Background and Rationale

Pancreatic cancer (PC) is a fatal disease with a poor 5-year overall survival of merely 8% [1]. Conventional therapies for PC such as chemotherapy, radiation therapy and surgery have fared with limited success, thus necessitating the need to develop alternative treatment strategies [2, 3]. Immunotherapy has emerged as a viable option but had stunted success in the efficacious treatment of pancreatic cancer patients [4, 5]. PC has limited genetic mutations, notably at the *KRAS* codon position with 12 mutations, and bear at the maximum 4000 potential immunogenic neoantigens, whereas melanoma has a predicted neoantigen count of 14000 that corroborates with success seen in melanoma immunotherapy [6-8]. Thus, one of the major challenges in PC immunotherapy is the identification of tumor-associated antigens (TAAs) that could be specifically targeted to cancer with negligible side-effects on healthy cells/organs.

TAAs expressed by cancer cells are mostly recognized as self-antigen due to the overriding of the immune system by immunoediting, therefore establishing a potent cancer-mediated tolerance against its TAAs [3, 9, 10]. Abnormal expression of antigens by cancer cells being recognized by immune surveillance that leads to the generation of autoantibodies is reflective of the immune response generated in cancer patients [10]. Expression of the antigen at the aberrant location where it is neither expressed in normal organ nor during other malignancy affecting that organ makes it a lucrative target for immune cells. In addition, post-translational modifications of such antigens affect their processing and loading on major histocompatibility complex (MHC molecules). The modified antigen-loaded MHC interacts with T cell receptors (TCRs), hence activating CD4⁺ T helper cells to mount a humoral autoantibody response, overriding immunological tolerance against the TAA [10, 11].

Mucin 4 (MUC4), a high molecular weight glycoprotein, is aberrantly overexpressed by and glycosylated in PC cancer cells, whereas it is negligibly present in normal pancreas and chronic pancreatitis (CP) patients [12-14]. Presence of autoantibodies against the VNTR region of another aberrantly overexpressed mucin in PC, Mucin 1 (MUC1), has been detected in platinum based drug resistant lung and ovarian cancers [15, 16], but there were no significant increase in MUC1 autoantibodies detected in PC patients when compared to age-matched controls [17]. Unlike MUC1, the MUC4 expression is restricted in most of the normal organs except for low levels being expressed in urogenital tracts and trachea in lungs [18]. In addition, autoantibodies against MUC4-TR region glycopeptides have been detected in colorectal cancer patients [19]. Thus, we hypothesized that MUC4 could be potentially immunogenic because of its spatial expression and post-translational modifications leading to tolerance against MUC4 being compromised in PC patients.

MUC4 has a potential autocatalytic Gly-AspPro-His (GDPH) cleavage site that generates the N-terminal MUC4 α subunit (containing the TR region) and a membrane-tethered MUC4 β subunit [20-22]. In contrast to the published studies on MUC4 autoantibodies [19], our study encompassed the entire length of MUC4 to demonstrate that MUC4 is immunogenic in PC patients. This design provided us a unique opportunity to assess the development of the humoral response to the entire protein backbone of MUC4 in PC patient serum. Due to the large molecular weight of MUC4 α we were unable to express and purify the recombinant protein in a bacterial system. We successfully purified the MUC4 β recombinant protein, which was used in our autoantibody study. Our lab is interested in utilizing MUC4 to develop an immunotherapy strategy to treat PC. Therefore, we also predicted potential major histocompatibility complex-I (MHC-I) HLA-binding immunogenic peptides using a reverse immunology approach [23] across the

entire protein sequence of MUC4. Most of the available studies on cancer vaccine immunotherapies primarily focus on HLA-binding peptides to induce T-cell mediated responses. Humoral response analysis against such peptides have not been investigated and thus excludes an equivalently crucial arm from factoring in the overall clinical outcome status of vaccine-immunized patients. We predicted the B-cell probability score for the same peptides and were keen to observe if an autoantibody signature was present against these T-cell peptides. Our study reported the presence of circulating autoantibodies against both MUC4 β protein and MUC4 peptides present in PC patient serum when compared to CP patient's and healthy individual's serum. This interesting observation suggests that MUC4 protein could potentially stimulate both T-cells and B-cells that may affect the overall clinical outcome. Further, we elucidated the various isotypes of autoantibodies present produced against MUC4 in PC patients. In addition, since the humoral response against MUC4 serves as a surrogate of compromised tolerance, we stratified and predicted patients who could have a higher probability of response to personalized MUC4-based immunotherapy.

3. Results

3.1 Analysis of Circulating MUC4 Autoantibody in PC patients

The primary aim of this study was to develop a novel platform for selective detection of circulating autoantibodies against the human MUC4 β recombinant protein. To our knowledge we are the first group to purify a recombinant human MUC4 β protein, which we have been utilized to develop the assay. Circulating autoantibodies against recombinant human MUC4 β protein was observed to be upregulated in PC patients as compared to healthy individuals and chronic pancreatitis patients. The low level of antibody positive sera and high level of antibody positive sera was defined at its cut-off values of <1.283 and ≥ 1.283 respectively (**Figure 1A**). Based on the cut-off values we

were able to determine that 76.7% of PC patients had high levels of MUC4 autoantibodies, whereas only 37% of healthy and 28.6% of CP patients showed high levels of MUC4 β autoantibodies (**Figure 1A**). We didn't observe a significant advantage in survival between PC patients with high levels of MUC4 autoantibodies and low levels of MUC4 autoantibodies (**Figure 1B**). Further we performed stage-wise analysis and observed that 86.7% of early-stage PC patients and 66.7% of late-stage patients had circulating MUC4 autoantibodies in their sera (**Figure 1C**). PC patient survival days analysis demonstrated that patients with higher serum reactivity had mean survival days (MSD) of 179 days for Stage 1-II and 309 days for Stage II-IV, compared to the patients with lower serum reactivity had MSD of 131 days for Stage 1-II and 244 days for Stage II-IV (**Figure 1C**).

3.2 Bioinformatics Analysis Predicts Potential MUC4 Peptides

Due to the difficulty in purification of the MUC4 α domain, we predicted potential adequate HLA-binding immunogenic MHC-I peptides inside both MUC4 α and MUC4 β domains to capture the entire MUC4 protein. A prerequisite for an immune response to arm, immunogenic peptides of MUC4 must be loaded on major histocompatibility complexes that would be cross-represented to effector immune cells (i.e. T-cells and B-cells). HLA-A2 is one of the widely used HLA-subtype for peptide predictions that have been used in designing vaccinations for melanoma and lung cancer [29, 37-39]. We used 3 prediction tools NetCTL1.0 and IEDB (Tepi Tool and Peptide processing) for determining HLA A2-binding T cell epitopes. NetCTL1.0 (based on a combined prediction of peptide MHC binding, proteasomal C terminal cleavage and TAP transport efficiency) software analyzed MUC4 sequence and predicted potential HLA-binding 9-mer amino acid sequences scored across 10 major human HLA-A2 isoforms. Peptides with combined high scores have higher specificity, sensitivity and higher affinity binding to HLA-A2 MHC-I complex. We further analyzed these peptides using IEDB (<http://tools.iedb.org/tepitool/>)

prediction TepiTool which is based on consensus method employing artificial neural network, combinatorial library networks and SMM-align. TepiTool scored the NetCTL1.0 predicted peptides based on their likelihood to bind to human HLA-A2*02.1 (mostly used in vaccine studies [39]), as well as, C57BL/6 mouse HLA-D^b and HLA-K^b MHC-I isotypes. In addition IEDB Processing (<http://tools.iedb.org/processing/>) analysis, which scores based on proteasomal C terminal cleavage, TAP transport efficiency and MHC-I complex loading, ranked these predicted peptides for both mouse and human HLA MHC-I loading. Lower predicted rank score suggests a higher probability of these peptides to be loaded on MHC-I complex and cross-presented to T-cells for activating them in an antigen-specific manner (**Figure 2**).

Our autoantibody analysis demonstrated MUC4 β specific antibodies being generated specifically in PC patients. Bioinformatics analysis predicted MHC-I peptides in MUC4 β region that made us curious to analyze the potential of these peptides to serve as B-cell immunogenic epitopes. For the same, we ran and scored these peptides on the IEDB B-cell prediction tool (<http://tools.iedb.org/bcell/>) and performed BCPred analysis (<http://ailab.ist.psu.edu/bcpred/predict.html>). Our B-cell epitope analysis showed that there was a difference in predicted scores between each peptide. Only 13 out of 24 predicted peptides were picked up by both prediction software and scored for the potential of them getting recognized by B-cells. We ranked these peptides based on their BCPred scores (**Figure 3**), and randomly selected peptides that represent all differentially scored B-cell epitope regions of both MUC4 α and MUC4 β to elucidate the plausibility of autoantibodies generated against these predicted peptides.

3.3 Screening of Autoantibody Signature against MUC4 Peptides

Our initial experiment elucidated the presence of autoantibodies against a MUC4 β recombinant protein which provides a large repertoire of epitopes that could be recognized

by B cells. In addition, our bioinformatics analysis predicted peptides that have the potential to serve as both T-cell and B-cell epitopes. To validate our observations, we developed a modified indirect ELISA to detect autoantibodies if any, generated against these epitopes. Our ELISA studies revealed differential serum reactivity to each of the 7 MUC4 α peptides whereas significantly low/ reactivity was seen against TR and PD2 peptides and 5 μ g/ml of BSA protein (**Figure 4A**). Further, we were curious to understand whether any peptide specifically fared better when compared to the survival status of PC patients. Our study showed that patients with high serum reactivity to peptides D3 and F3 survived longer than their low serum reactivity counterparts, but the rest of the 5 MUC4 α peptides didn't show similar relationship (**Figure 4B**).

Similarly, PC patient serum significantly reacted with all 4 peptides from MUC4 β region and low reactivity was observed with control peptides and protein (**Figure 5A**). In addition, high serum reactivity to MUC4 β peptides didn't provide a survival advantage (**Figure 5B**). Thus, our study shows for the first time that MUC4 MHC-I HLA-binding T-cell peptides could potentially be recognized by B-cells and might have a dual (activate both cellular and humoral immune pathway) epitope property. Further, we stratified PC patients with high autoantibody levels detected in their sera and observed that we could potentially design a selection strategy to elucidate patients who likely might have compromised peripheral tolerance (**Table I**). To ensure the specificity of the detected antibodies, we compared the serum reactivity of randomly selected 10 healthy and 10 chronic pancreatitis individuals and we observed that PC patient specific reactivity was seen to all 11 MUC4 peptides and negative in healthy or CP individuals (**Supplemental Figure 1 & 2**).

3.4 Isotype Analysis of MUC4 Autoantibodies in PC Patients

Autoantibody isotypes such as IgM autoantibodies in cancer patient serum have been shown to correlate with poor prognosis and overall survival of those patients [15].

Activation of B-cells to its mature form after receiving secondary stimulation from CD4⁺ T-cells induces isotype switching to generate various classes of antibodies with the same variable antigen binding region as the original antibody generated to the antigen by the VD (J) recombination pathway [40, 41]. Our observations demonstrated that autoantibodies are present against both recombinant MUC4 protein and its peptides, but higher levels of these autoantibodies didn't significantly correlate with patient survival data. Isotyping of mucin autoantibodies could provide some insight into understanding whether activation of B-cells confers any survival advantage to PC patients. We measured different isotypes (IgA, IgM and IgG) of MUC4 autoantibodies present in PC patient sera and correlated with overall patient survival and prognosis. All three MUC4 β protein autoantibody isotypes were detected, however presence of IgG isotype autoantibodies in serum segregated PC patients in two groups. Despite the segregation, we didn't observe any significant correlation between autoantibody isotypes and patient survival **(Supplemental Figure 3)**.

Isotype analysis of autoantibodies against MUC4 α peptides demonstrated differential levels of isotypes present in PC patient serum **(Figure 6A)**. Our data further elucidated that high levels of IgM autoantibody against A2 and D4 MUC4 α peptides correlate strongly with overall patient survival ($P < 0.001$) **(Figure 6B)**. Analysis of MUC4 β peptides also demonstrated that all isotypes were detected **(Figure 7A)**. Similarly, Kaplan-Meier graph shows that high levels of IgM autoantibody to B1 peptide correlated strongly with PC survival **(Figure 7B)**. Our data showed that MUC4 is not only immunogenic in PC patients but also induces maturation of B-cells. Interestingly, for all MUC4 peptides, patients with IgM isotype autoantibodies had better median survival than patients with either IgG or IgA autoantibodies **(Supplemental Figure 4)**.

4. Discussion

The present study investigated the possibility of MUC4 being immunogenic in PC patients. We developed a modified ELISA strategy and tested both MUC4 recombinant protein and peptides to determine the presence of autoantibodies in patient sera. MUC4 is aberrantly overexpressed in PC patients and its expression increases with gradual progression of the malignancy [42]. MUC4 overexpression and aberrant glycosylation in PC and its absence in the normal pancreas [14] makes it a potential antigen that might get recognized by the immune system (specifically by B-cells). The modified indirect ELISA discovery platform for detecting human serum autoantibodies was developed for efficient antigen-antibody binding and provide a remarkably low background. Cancer-associated autoantibodies to recombinant MUC4 β protein were identified in PC patient sera, whereas healthy individuals and CP patients were negligible for MUC4 autoantibodies, although survival analysis didn't reveal any robust correlation with levels of autoantibodies present in PC patient sera. Presence of circulating MUC4 autoantibodies in PC patients therefore supports the hypothesis that immunological tolerance is compromised against MUC4 and gets recognized by B-cells.

In our study, we utilized both recombinant protein and peptides to investigate whether different domains/regions of MUC4 is immunogenic other than just the TR region sequence that is well studied. Bioinformatics analysis predicted HLA-binding peptides that have potential to serve as a dual epitopes for both cellular and humoral immune pathways in human as well as in mouse. To capture MUC4 α region we randomly selected peptides corresponding to that region due to unavailability of purified recombinant MUC4 α protein. The rationale behind using peptides of MUC4 β region was to recognize immunodominant region present in MUC4 β protein and it was investigated on a peptide-based ELISA platform. Our study revealed that only PC patient serum contained autoantibodies against

MUC4 peptides, whereas there was negligible reactivity detected in control individuals. Our patient stratification strategy based on their serum reactivity with respective MUC4 peptides elucidate that the tolerance against each peptide is compromised in patient-to-patient basis. Such case basis-compromised tolerance might correlate with the differential response of PC patients to MUC4-based immunotherapy that needs to be investigated in the future.

Detection of specific autoantibody isotypes suggests activation and maturation of B-cells leading to the systemic availability of neutralizing antibodies protecting against cancer, which could be translated to the observed negative correlation of immature IgM antibodies with the survival of patients [15, 19]. Our data in contrast showed that high levels of IgM autoantibodies against A2, D4 and B1 have a statistically strong correlation with longer survival of PC patients. Further, PC patients with high levels of IgM autoantibodies have better survival than those who have IgG/IgA circulating autoantibodies, thus making a case for a better understanding of the role of humoral responses in the survival of patients.

Overall our studies provide enough evidence to suggest that entire MUC4 is immunogenic and peripheral tolerance to MUC4 is compromised in PC patients. In a recent study, it has been shown that CD8⁺ T cell-specific MUC16 neoantigen epitopes provide a survival advantage to PC patients [43]. Interestingly our data for the first time reveals that T-cell epitopes located on various regions of the MUC4 protein sequence (excluding VNTR region), could simultaneously be recognized by B-cells and activate the humoral arm of the immune system. Further, circulating IgM MUC4 autoantibodies against particular peptides, showed to provide protection to the PC patients and correlates with increased overall survival status. Cancer vaccines have emerged as an alternative treatment modality for cancer patients who respond poorly to traditional therapies. An

efficient vaccine-mediated immunity requires strong activation of both humoral (antibody and memory B-cell generation) and cellular responses (activation of both CD4⁺ helper T-cells, CD8⁺ cytotoxic T cells and memory T-cells), which provides optimal protection from the disease [44]. We propose that the dual antigenic (activate both B & T cells) epitope quality and aberrant expression of MUC4 in all the stages of PC tumor development qualifies it to be a strong candidate for immunotherapy strategies like cancer vaccines.

Table I: Autoantibodies reactivity against MUC4 peptides can predict patients with compromised tolerance

Patient No.	Stage	Alpha peptide recognized	Beta peptide recognized
12	IIB	A2, E4, D3, C4, F1, F3	B1, B3, B4, E3
6	III	A2, E4, D3, C4, F1	B1, B3, B4, E3
18	IIB	A2, C4, D4, F3	B1, E3
5	IIB	A2, E4, F3	B1, B4
7	IV	E4, D3, C4, F1	B4
9	IIA	A2, D4, F1, F3	B1
30	III	A2, E4	B4, E3
16	IV	E4	B3, B4
25	IIB	D4, C4	B3
22	IIB	F3	E3
15	IIA	D4, F1, F3	
3	III	D4, F1	
24	IV	C4	
10	IV		B1
20	IIB		E3

Figure 1: Autoantibodies against MUC4 β were detected in PC patient serum.

Indirect ELISA study revealed that PC patients had high expression of circulating autoantibodies to recombinant human MUC4 β protein (A). The high levels of MUC4 autoantibodies didn't correlate with PC survival as seen in Kaplan-Meier graph (B). Further analysis showed that patients with high levels of MUC4 autoantibodies, both at early (Stage I-II) and late (Stage IV) stages of PC, had longer survival than patients with low levels of autoantibodies (C).

Figure 1

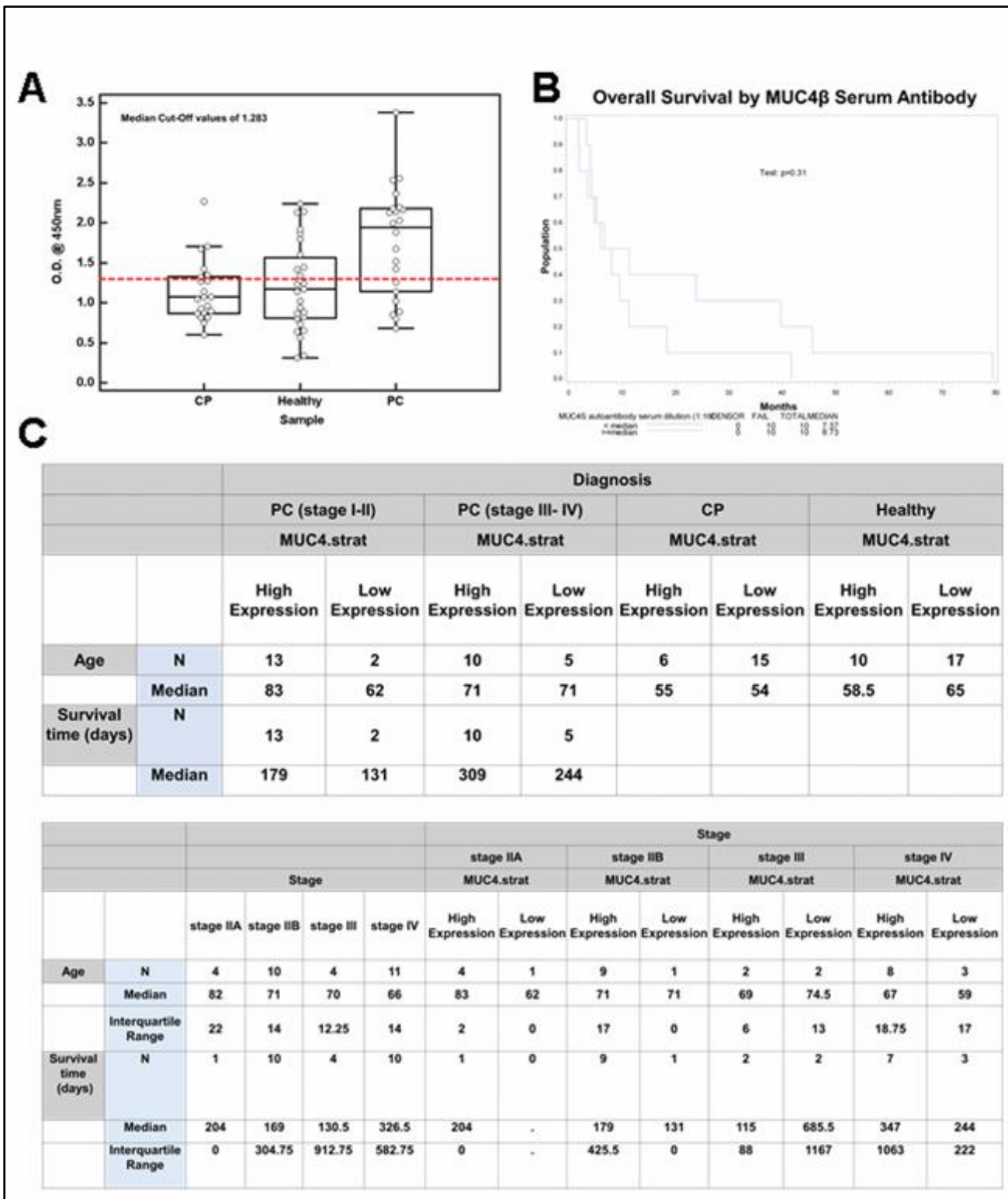


Figure 2: Bioinformatics analysis predicted T-cell immunodominant epitopes of MUC4.

NetCTL1.0 and IEDB software analysis based on combined prediction of peptide MHC binding, proteasomal C terminal cleavage and TAP transport efficiency, predicted and ranked T-cell immunodominant epitopes. Lower predicted rank score suggests a higher probability of these peptides to be loaded on MHC-I complex and cross-represented to T-cells for activating them in an antigen-specific manner.

Figure 2

Name	Peptide Seq	Human T cell Score (NetCTL 2.0)	Human MHC-I (HLA-A*02:01) TepiTool Percentile rank	Human MHC-I (HLA-A*02:01) Peptide Processing rank	Murine MHC-I (H-2-Kb & H-2- Db) TepiTool Percentile rank	Murine MHC-I (H-2-Kb & H-2-Db) Peptide Processing rank
A2	MMTSEKITV	0.5474	1.8	-0.52	H-2-Kb : 8.1 H-2-Db : 16.0	H-2-Kb : -2.62 H-2-Db : -2.74
E1	QATDTFSTV	0.4943	14	-2.64	H-2-Kb : 3.45 H-2-Db : 17.0	H-2-Kb : -2.56 H-2-Db : -3.05
F4	TLANSVVST	0.2844	6.4	-2.15	H-2-Kb : 52 H-2-Db : 63	H-2-Kb : -3.60 H-2-Db : -3.83
C3	AAMTHTHQA	0.4335	8.4	-2.80	H-2-Kb : 10.1 H-2- Db : 2.1	H-2-Kb : -2.78 H-2-Db : -2.43
E2	SVHNVGTGV	0.5395	5.5	-2.35	H-2-Kb : 11.1 H-2- Db : 16.0	H-2-Kb : -2.52 H-2-Db : -3.19
D4	VMETVTQET	0.4484	15	-3.02	H-2-Kb : 33.5 H-2- Db : 37.0	H-2-Kb : -3.58 H-2-Db : -3.77
C4	NVMETVTQE	0.2021	30	-4.12	H-2-Kb : 43 H-2- Db : 48	H-2-Kb : -4.23 H-2-Db : -4.31
A1	MMTSKITIM	0.5697	3.6	-0.86	H-2-Kb : 1.35 H-2- Db : 3.2	H-2-Kb : -1.98 H-2-Db : -2.25
F1	SVHNVMETV	0.5137	6.7	-1.36	H-2-Kb : 12.6 H-2- Db : 5.7	H-2-Kb : -2.36 H-2-Db : -2.65
D2	SVIQMIKTA	0.3433	14.0	-2.58	H-2-Kb : 18.0 H-2- Db : 5.3	H-2-Kb : -2.93 H-2-Db : -3.03
F2	TLTQMMTST	0.2905	8.3	-3.34	H-2-Kb : 44.5 H-2- Db : 32.0	H-2-Kb : -3.97 H-2-Db : -4.22
C2	HTSQQITTA	0.6948	23	-3.49	H-2-Kb : 77.5 H-2- Db : 29.0	H-2-Kb : -3.66 H-2-Db : -3.67
D3	HTTQSTTEL	0.8685	22	-2.71	H-2-Kb : 26.5 H-2- Db : 8.7	H-2-Kb : -2.56 H-2-Db : -2.49
C1	MTSKITITMT	0.7004	21	-3.66	H-2-Kb : 41 H-2- Db : 61	H-2-Kb : -3.78 H-2-Db : -4.05
A3	V TSAALVTV	0.6861	4.4	-1.03	H-2-Kb : 9.1 H-2- Db : 16.0	H-2-Kb : -2.42 H-2-Db : -2.98
B3	VVEAFlyHV	0.5953	2.7	-0.74	H-2-Kb : 1.15 H-2- Db : 26.0	H-2-Kb : -1.85 H-2-Db : -3.32
B4	QLLA AVVEA	0.3776	1.8	-0.71	H-2-Kb : 59.5 H-2- Db : 26.0	H-2-Kb : -3.32 H-2-Db : -3.34
E3	CLYNQTSRV	0.4851	0.2	-0.80	H-2-Kb : 3.3 H-2- Db : 35	H-2-Kb : -2.05 H-2-Db : -3.34
A4	GVSQEIFTL	0.4879	5.4	-1.07	H-2-Kb : 30.0 H-2- Db : 27.0	H-2-Kb : -2.30 H-2-Db : -2.52
D1	LVTSKVFRM	0.4660	12.0	-2.61	H-2-Kb : 2.8 H-2- Db : 25.0	H-2-Kb : -2.31 H-2-Db : -3.32
B2	TLIQYTSNA	0.3345	3.1	-1.31	H-2-Kb : 14.6 H-2- Db : 40.0	H-2-Kb : -2.41 H-2-Db : -3.50
B1	SMAEVNASV	0.6073	0.3	0.57	H-2-Kb : 6.6 H-2- Db : 15.0	H-2-Kb : -2.27 H-2-Db : -2.67
F3	TISAITSKV	0.5102	3.2	-1.54	H-2-Kb : 26.3 H-2- Db : 62.0	H-2-Kb : -2.81 H-2-Db : -3.29
E4	AITSKVSTI	0.4367	12.0	-2.31	H-2-Kb : 7.1 H-2- Db : 11.0	H-2-Kb : -2.42 H-2-Db : -2.86

Figure 3: Bioinformatics analysis predicted B-cell immunodominant epitopes of MUC4.

IEDB B-cell prediction software analysis based on sequence characteristics of the antigen using amino acid scales and HMMs, predicted and ranked B-cell immunodominant epitopes. High predicted rank score suggests higher probability of these peptides to be recognized by B-cells.

Figure 3

Name	Peptide Seq	BCPRED	IEDB (Epitope)
A2	MMTSEKITV	MMTSEKITVTTSTGSLGNP (0.999)	-
E1	QATDTFSTV	SQATDTFSTVPPTPSITSS (0.998)	MAQTQRTGTSRGSDTISLASQATDTFSTVPPTPP SITSSGLTSPQTQHTLSPSGSGKTF
F4	TLANSVVST	TLANSVVSTPGGPEGQWTS (0.998)	SVVSTPGGPEGQWTSASASTSPDTAAAMTHTHQ
C3	AAMTHTHQA	SASASTSPDTAAAMTHTHQA (0.997)	SVVSTPGGPEGQWTSASASTSPDTAAAMTHTHQ AESTEASGQT
E2	SVHNVGTGV	SPSVHNVGTVSQKTSPSGE (0.985)	QTSTLHRTTSTPSFSPSVHNVGTVSQKTSPSG ETATSS
D4	VMETVTQET	VHNVMETVTQETAPPDEMTT (0.975)	ETVTQETAPPDEMTTSFPSSVT
C4	NVMETVTQE	VHNVMETVTQETAPPDEMTT (0.975)	ETVTQETAPPDEMTTSFPSSVT
A1	MMTSKITIM	MMTSKITIMTTSTDSTLGNP (0.969)	TSMMS
F1	SVHNVMETV	SPSVHNVMETVTQETAPPDE (0.966)	-
D2	SVIQMIKTA	YSVIQMIKTATSPSSPMLD (0.963)	-
F2	TLTQMMTST	TETTSKAQDTLTQMMTSTL (0.959)	-
C2	HTSQQITTA	MLDRHTSQQITTAPSTNHST (0.958)	TATSPSSPMLDRHTSQQITTAPSTNHSTIHSTST P
D3	HTTQSTTEL	FSSNPSRDSHTTQSTTELLS (0.946)	EGISTSGETTRFSSNPSRDSHTTQSTTELLSASAS HGAIPVSTGMASIVPG
C1	MTSKITIMT	TNTLMMTSKITIMTTSTDST (0.905)	-
A3	VTSAAALVTV	PGETSSVPVTGSLMPVTSAA (0.822)	KITVTTSTGSLGNPGETSSVPVTGSLMPV
B3	VVEAFLYHV	VVEAFLYHVPRRSEEPNDV (0.901)	-
B4	QLLAAVVEA	EFQYRPRGPVIDFLNNQLLAAVVEAFLYHV PRRSEEPNDV (0.901)	-
E3	CLYNQTSRV	HCNAESQCLYNQTSRVGNSS (0.848)	CLYN
A4	GVSQEIFTL	GRPHERSSFSFGVSQEIFTL (0.763)	TLSTALSPSSLPPKISTAFHTQQSEGAETTGRPHE RSSFSFGVSQ
D1	LVTSKVFRM	TISSPSSVSNTFLVTSKVFR (0.758)	-
B2	TLIQYTSNA	GRVIEAYKGQTTLIQYTSNA (0.745)	REVSKNYEQANATLNQYPPSINGGRVIEAYKGQT TLIQYTSNAEDANFT
B1	SMAEVNASV	-	EENASMAEVN
F3	TISAITSKV	-	TISAI
E4	AITSKVSTI	-	-

Figure 4: Reactivity against predicted MUC4 α epitopes was found in PC patients.

Indirect ELISA study demonstrated that PC patient serum contains circulating autoantibodies against MUC4 α immunodominant predicted epitopes, whereas low reactivity to PD2 internal protein control was observed. The serum reactivity to MUC4 α was observed to be significantly higher than PD2 and TR control peptides (A). Presence of autoantibodies against these peptides didn't confer any survival advantage (B). ANOVA analysis of group means was significant ($P < 0.001$), ** and ** denotes paired T-test between PD2 vs. MUC4 peptides and TR vs peptides respectively ($P < 0.001$)

Figure 4

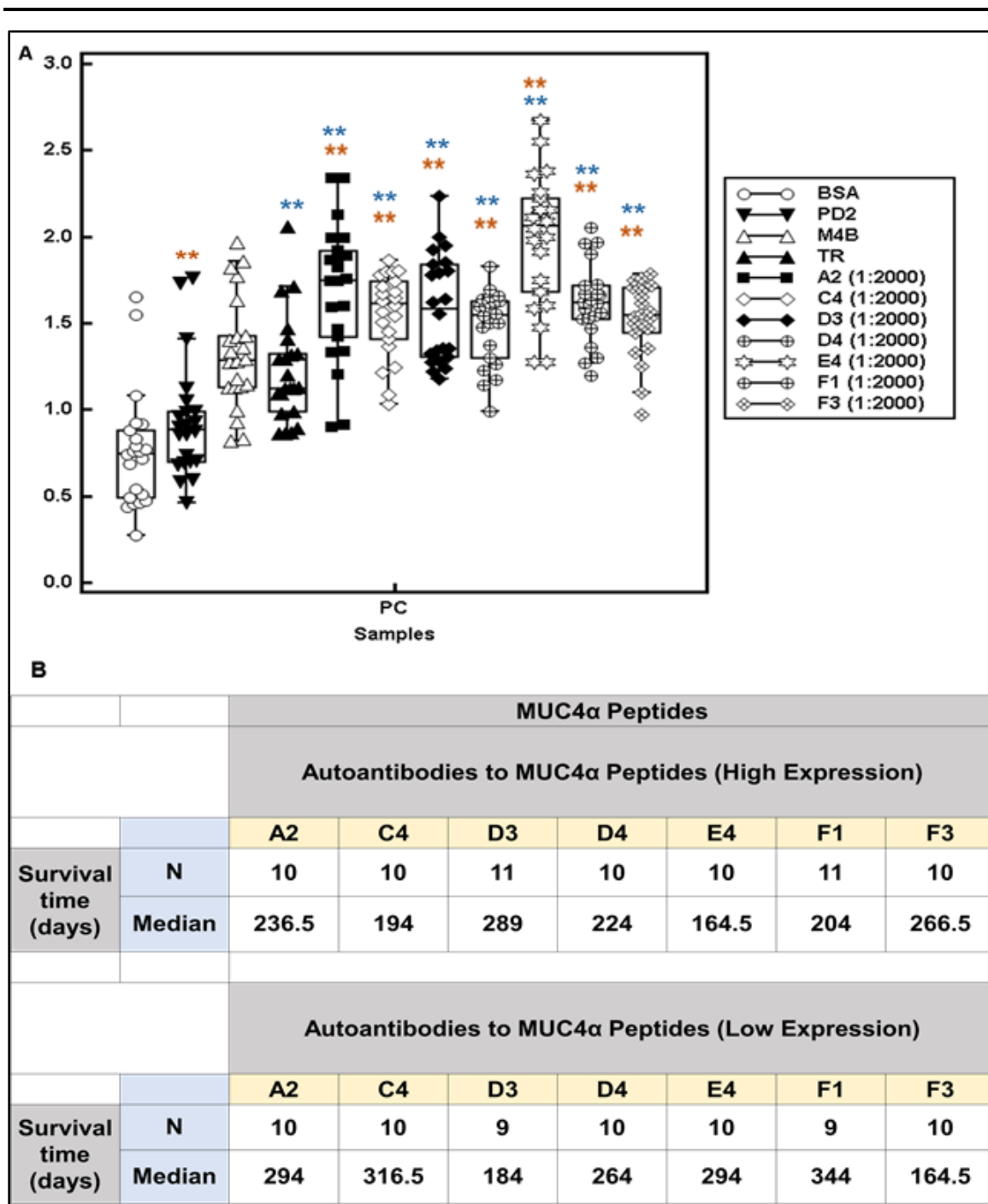


Figure 5: Reactivity against predicted MUC4 β epitopes was found in PC patients.

Indirect ELISA study demonstrated that PC patient serum contains circulating autoantibodies against MUC4 β immunodominant predicted epitopes, whereas low reactivity to PD2 internal protein control was observed. The serum reactivity to MUC4 α was observed to be significantly higher than PD2 and TR control peptides (A). Presence of autoantibodies against these peptides didn't confer any survival advantage (B). ANOVA analysis of group means was significant ($P < 0.001$), ** and ** denotes paired T-test between PD2 vs. MUC4 peptides and TR vs peptides respectively ($P < 0.001$)

Figure 5

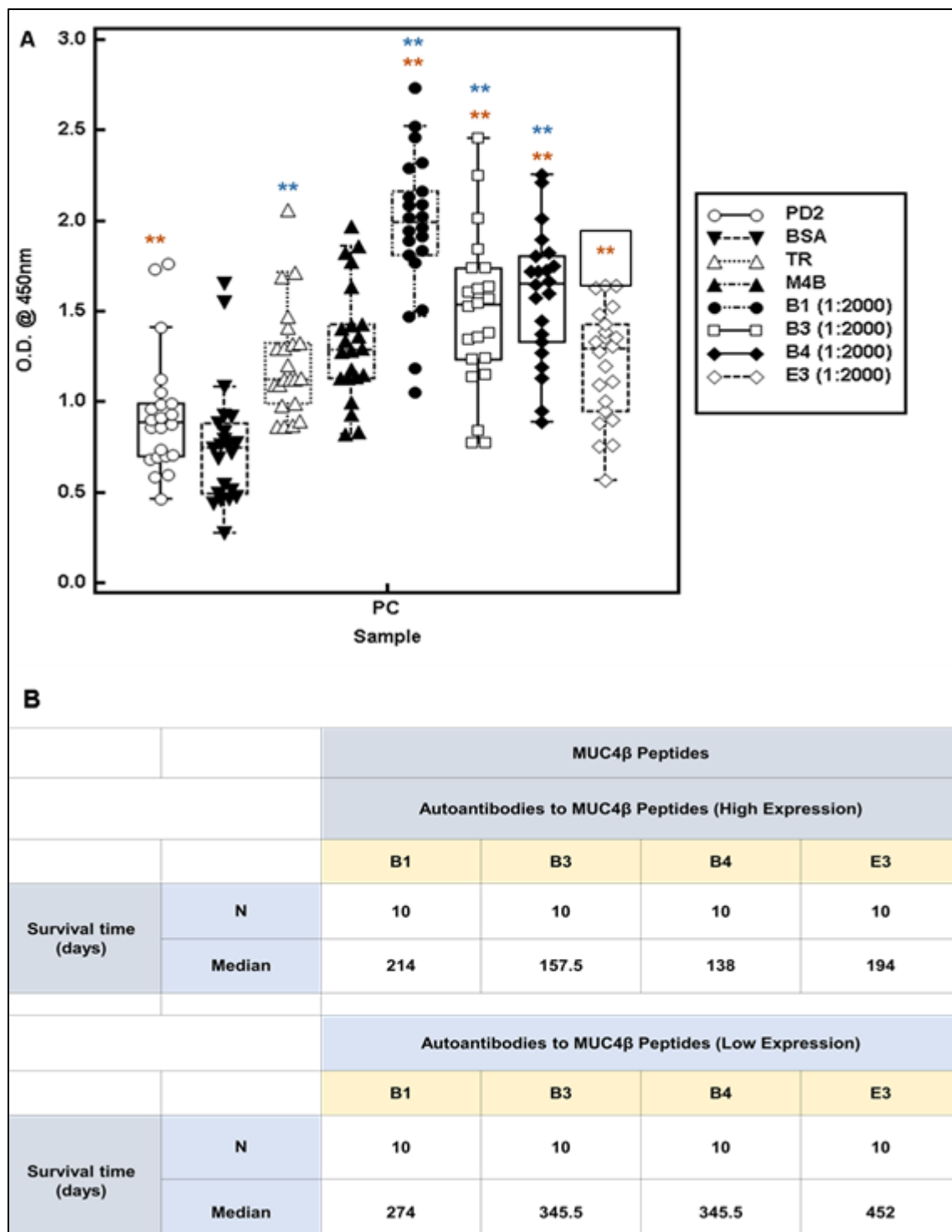


Figure 6: IgM autoantibodies to MUC4 α peptides show a better prognosis in PC patients.

Isotyping analysis revealed that differential levels of all three IgM, IgG and IgA isotypes have been detected against all MUC4 α peptides (A). Kaplan-Meier graph shows that high levels of IgM autoantibodies to A2 and D4 (blue arrows) significantly correlated with PC patient survival (MSD: 344 days & 535 days respectively).

Figure 6

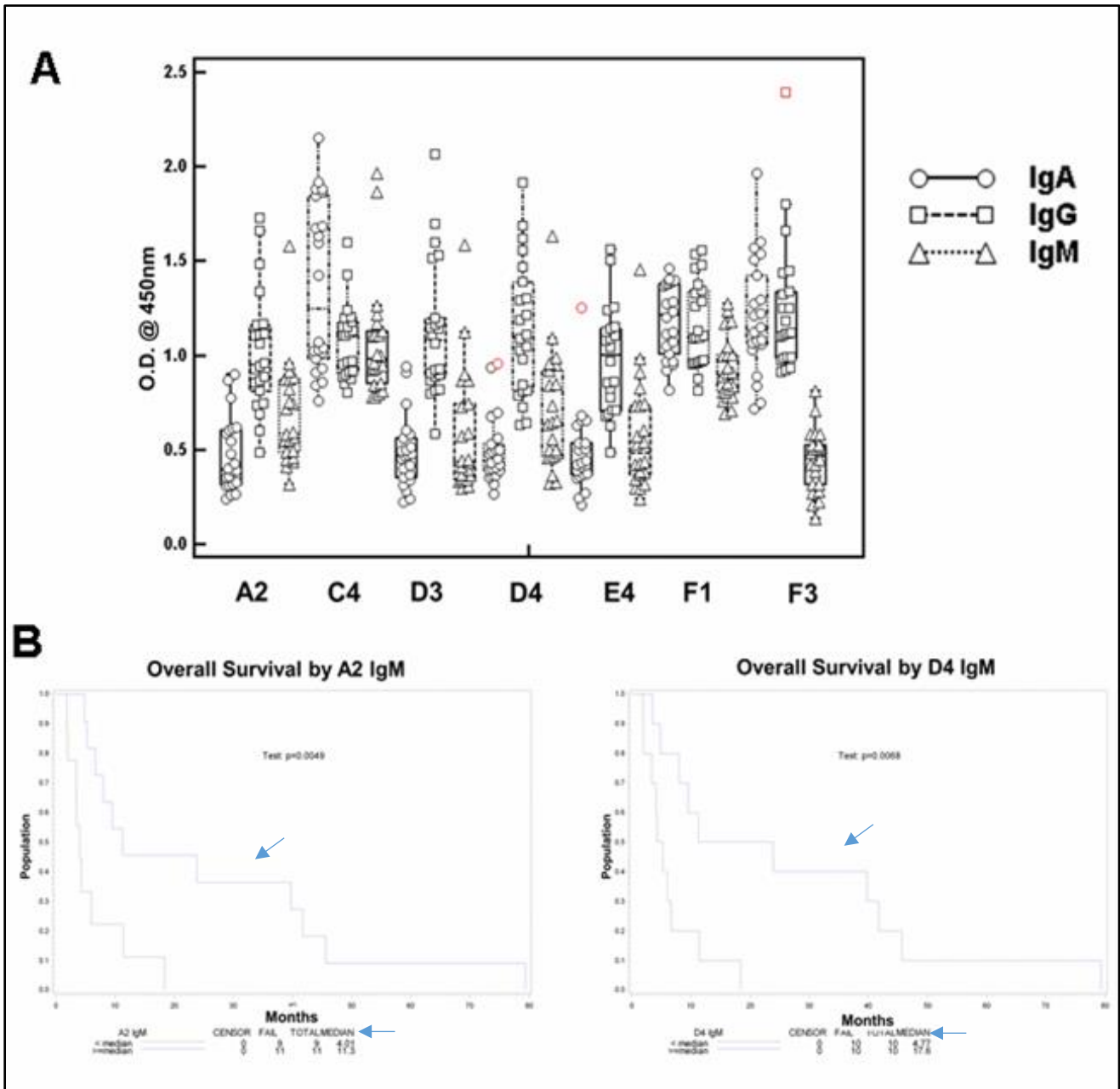
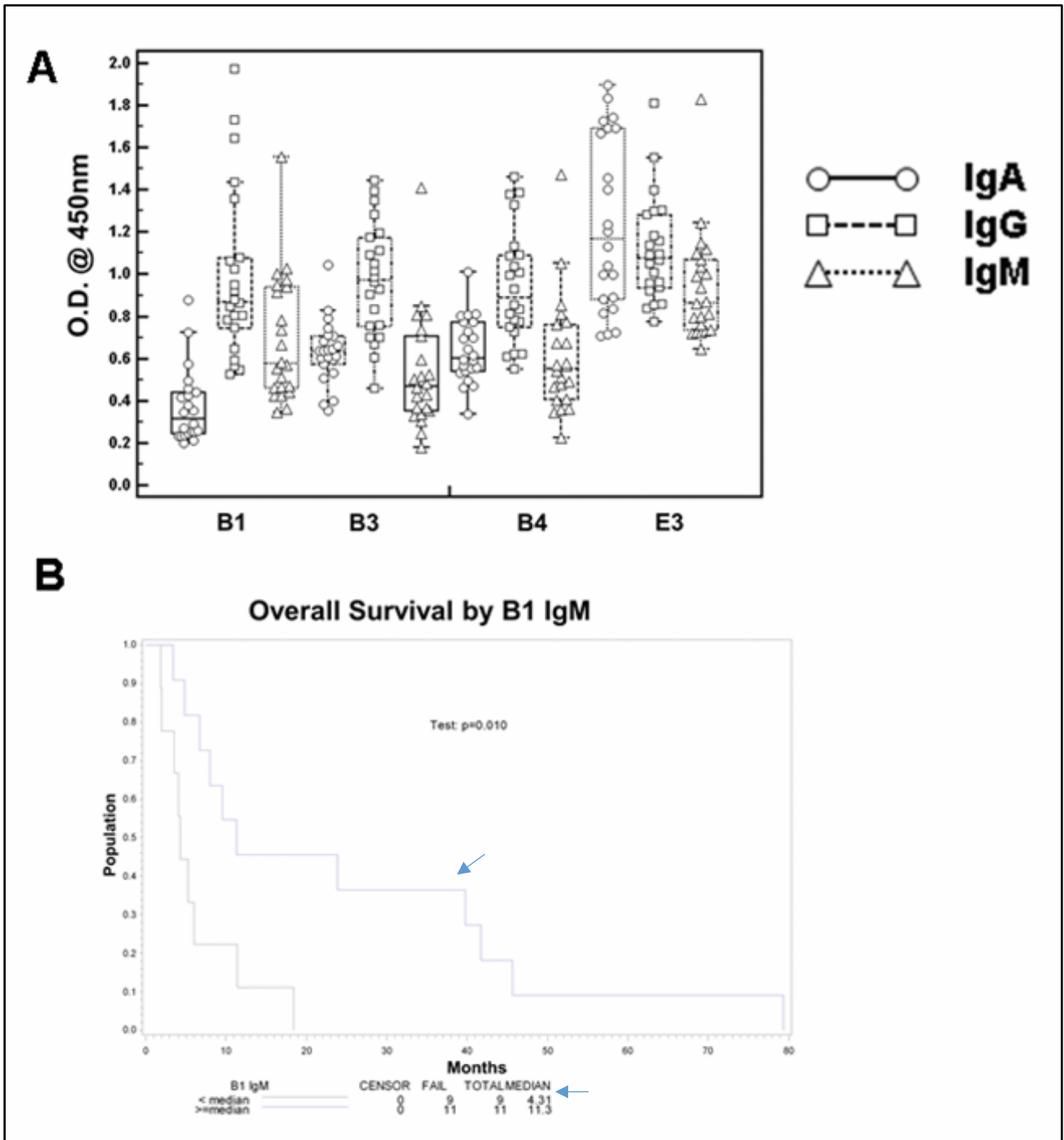


Figure 7: IgM autoantibodies to MUC4 β peptides show a better prognosis in PC patients.

Isotype analysis revealed that differential levels of all three IgM, IgG and IgA isotypes have been detected against all MUC4 β peptides (A). Kaplan-Meier graph shows that high levels of IgM autoantibodies to B1 (blue arrow) significantly correlated with PC patient survival (MSD: 344 days).

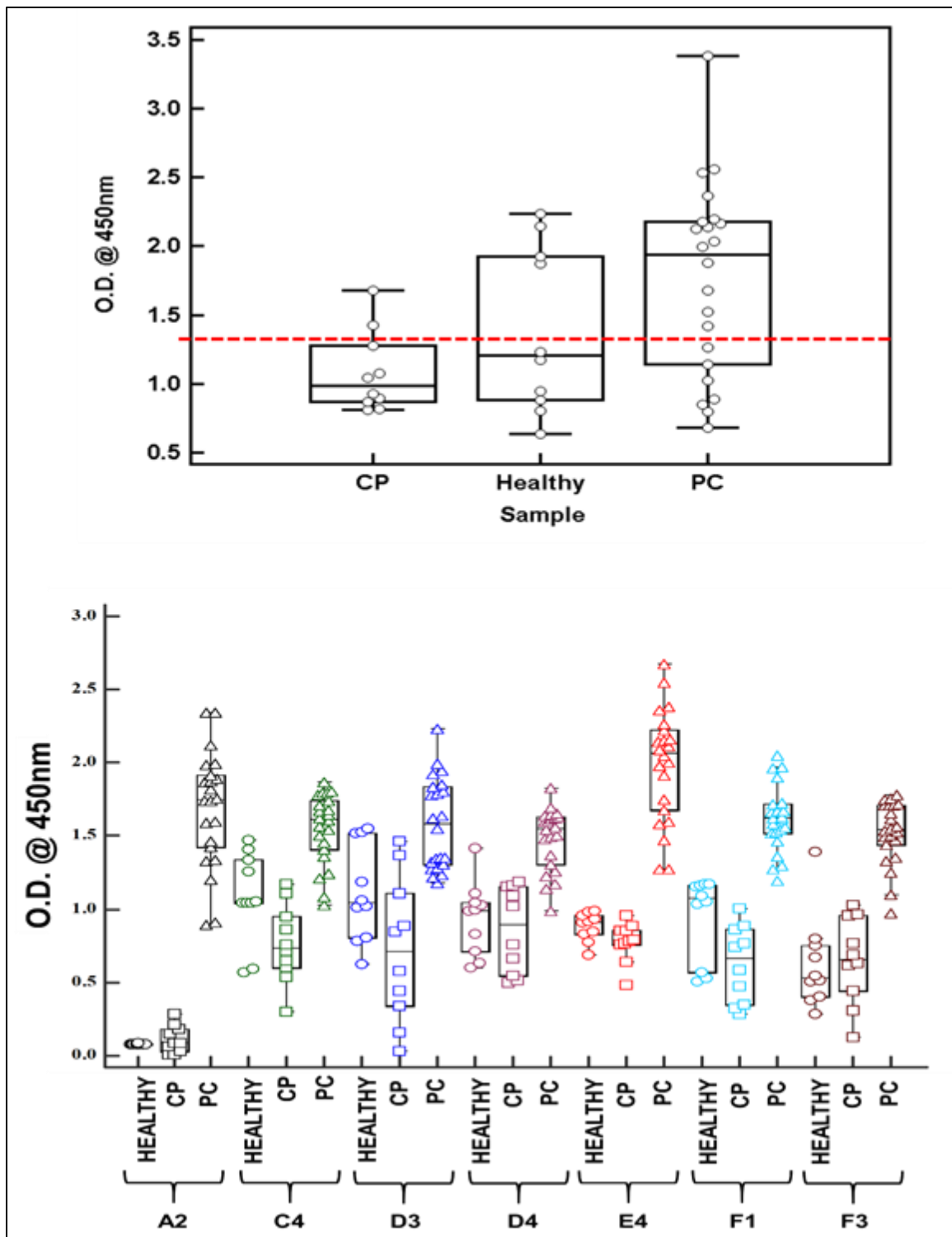
Figure 7



Supplemental Figure 1: Reactivity against predicted MUC4 α epitopes was found only in PC patients.

PC specific reactivity was found against MUC4 α peptides with significantly low reactivity in both healthy and CP controls, suggesting that these peptides could serve as unique immunodominant epitopes being immunologically recognized specifically by PC patient's immune system.

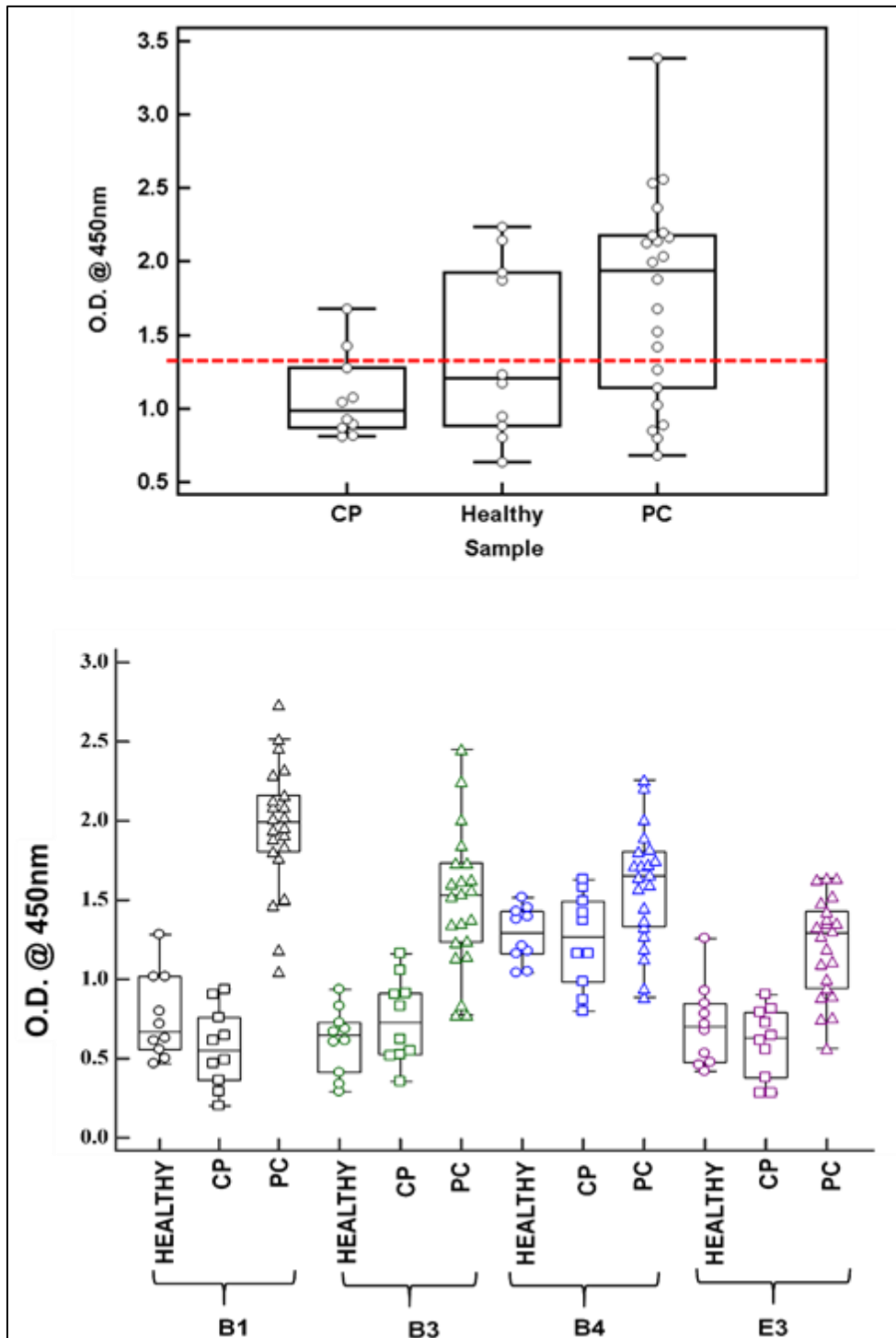
Supplemental Figure 1



Supplemental Figure 2: Reactivity against predicted MUC4 β epitopes was found only in PC patients.

PC specific reactivity was found against MUC4 β peptides with significantly low reactivity in both healthy and CP controls, suggesting that these peptides could serve as unique immunodominant epitopes being immunologically recognized specifically by PC patient's immune system.

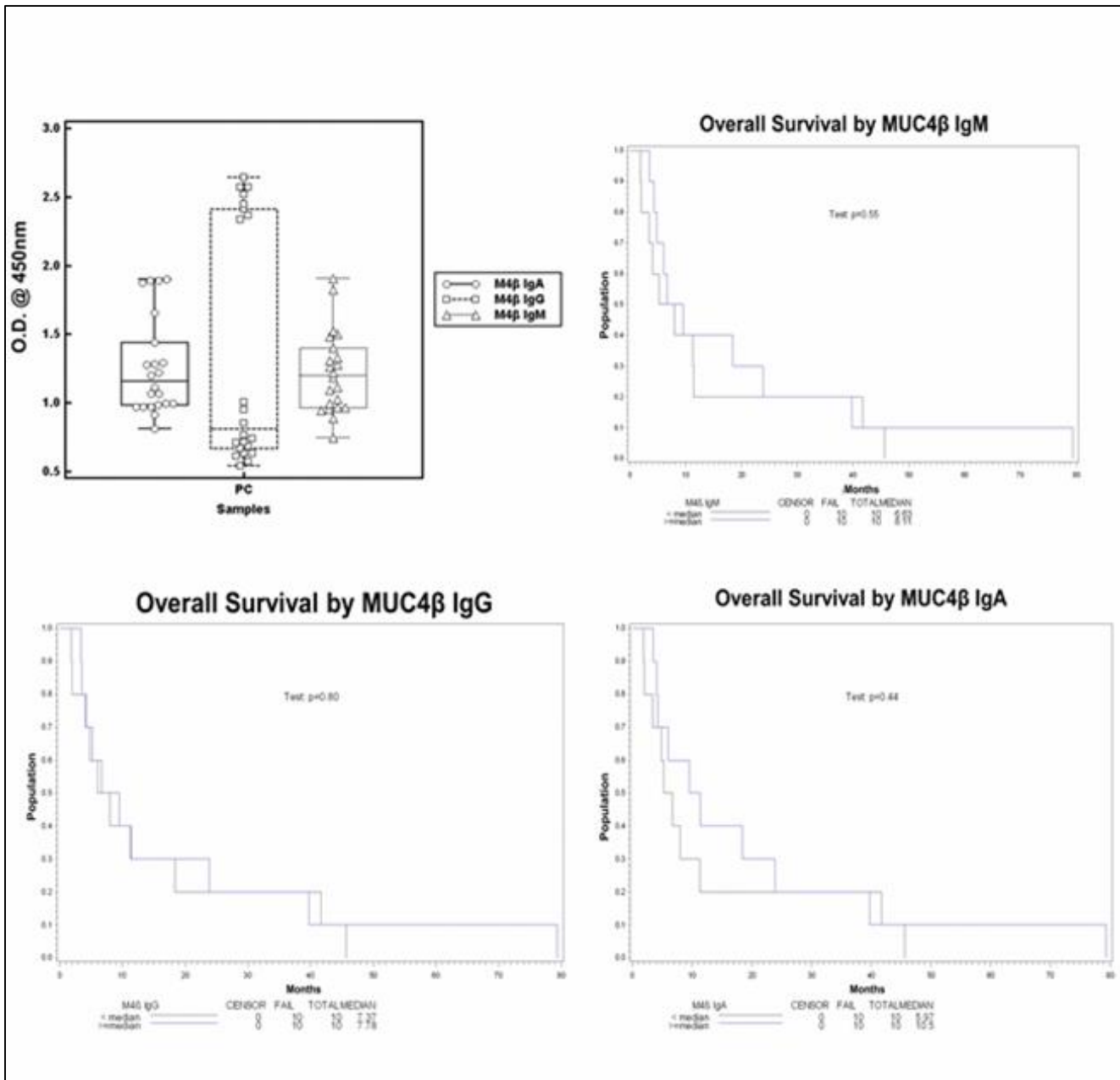
Supplemental Figure 2



Supplemental Figure 3: Isotyping of MUC4 circulating antibodies didn't show any correlation with survival status of PC patients.

Isotype analysis of autoantibodies to recombinant human MUC4 β protein showed that PC patients could be segregated on the levels of IgG autoantibodies present in PC patient serum. However, none of these isotypes correlated with PC patient survival.

Supplemental Figure 3



Supplemental Figure 4: Autoantibodies isotype to MUC4 α and MUC4 β peptides reveals a differential correlation with patient survival.

Correlation of PC patient survival with isotypes of autoantibodies to MUC4 peptides present in patient sera demonstrates that high levels of IgM autoantibodies seem to positively correlate with patient survival compared to IgG or IgA autoantibodies for all peptides.

Supplemental Figure 4

		A2			C4			D3			D4		
		IgM	IgG	IgA	IgM	IgG	IgA	IgM	IgG	IgA	IgM	IgG	IgA
Survival time (days)	N	11	10	10	10	10	11	10	10	11	10	10	11
	Median	344	246.5	174.5	316.5	266.5	204	316.5	266.5	184	535.5	266.5	204

		E4			F1			F3		
		IgM	IgG	IgA	IgM	IgG	IgA	IgM	IgG	IgA
Survival time (days)	N	11	10	10	10	11	10	9	11	9
	Median	289	266.5	194	266.5	204	236.5	244	244	204

		B1			B3			B4			E3		
		IgM	IgG	IgA	IgM	IgG	IgA	IgM	IgG	IgA	IgM	IgG	IgA
Survival time (days)	N	11	11	11	10	10	10	10	10	10	10	11	10
	Median	344	289	184	316.5	224	224	316.5	266.5	246.5	266.5	244	266.5

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CHAPTER 4: DEVELOPMENT OF MUC4 NANOVACCINE

The material covered in this chapter is the subject of a research article

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1. Synopsis

Mucin 4 (MUC4) is high molecular weight glycoprotein that is differentially overexpressed in pancreatic cancer (PC) and functionally contributes to disease progression while its expression correlates with poor survival. Further, due to its aberrant glycosylation and extensive splicing in cancer, MUC4 is a potential target for cancer immunotherapy. Our previous studies have demonstrated the utility of amphiphilic polyanhydride nanoparticles as a useful platform for the development of protein-based prophylactic and therapeutic vaccines. In present study, we encapsulated purified recombinant human MUC4 (MUC4 β) protein in 20:80 ratio of amphiphilic polyanhydride (CPTEG & CPH) adjuvants (MUC4-nanovaccine) and evaluated its ability to activate dendritic cells and induce anti-tumor immunity. Immature dendritic cells when pulsed with MUC4-nanovaccine exhibited more than 2-fold increase in surface expression of activation markers (MHC-II and MHC-I) and 1.5-fold increase in pro-inflammatory cytokines (IFN- γ , IL-6, and IL-2) levels as compared to cells exposed to MUC4 β alone or MUC4 β mixed with blank nanoparticles (MUC4 +NP). Further, the antibody analysis in the sera collected from immunized mice showed two-fold higher levels of IgG2b antibodies than IgG1 antibodies, suggesting a predominantly Th1-type of immune response in MUC4-nanovaccine group. Thus, our findings demonstrate MUC4-nanoformulation as a novel platform for PC vaccine development.

2. Background and Rationale

Pancreatic cancer (PC) has a dismal prognosis with an overall survival rate of 8%, due to the limited efficacy of existing treatment modalities like surgery, chemotherapy, and radiation [1, 2]. Furthermore, PC has an elaborate immunosuppressive tumor microenvironment comprising of high desmoplasia, immune-suppressive cells and an anti-inflammatory cytokine *milieu* [3]. Due to high level of chemotherapy-induced toxicity, PC patients seldom benefit from chemotherapy. Recent studies have shown that immunotherapy-based strategies like cancer vaccines can provide therapeutic benefit by breaking the tolerance, overcoming immunosuppression and thereby, improving the overall survival and quality of life [4, 5]. However, the development of anti-cancer vaccines are rather more arduous due to the challenges in finding optimal tumor-associated antigens (TAAs), because the majority of these antigens behave as “self”, and therefore, are immunologically ignored by the host immune system leading to development of tolerance against TAAs [4].

Mucins are high molecular-weight glycoproteins that are overexpressed on pancreatic tumor cells and have oncogenic functions in PC pathogenesis. MUC family members have emerged as TAAs for PC and are currently being exploited for cancer immunotherapy. Mucin1 (MUC1) is one of the well-studied targets for PC vaccine studies [6]. MUC1 peptide and glycopeptide vaccine studies have shown the potential of reprogramming the immune system against mucins and generating anti-tumor responses in various malignancies [7-12]. However, limited immunogenic epitopes provided by peptide-based MUC1 vaccines have achieved suboptimal clinical success in PC patients [9, 13, 14]. Unlike MUC1, Mucin4 (MUC4) is undetectable in normal pancreatic tissue and its expression progressively increases with disease progression and [15]. MUC4 is putatively cleaved at a Gly-AspPro-His (GDPH) site in an autocatalytic manner into two

subunits: a large N-terminal MUC4 α containing the tandem repeat region and a smaller membrane-tethered MUC4 β [16-18]. The membrane-tethered MUC4 β region is considered functionally important as it has 3 EGF-like domains that interact with HER-2 and promote cell proliferation. Further, targeting MUC4 β is simplified since the subunit will still be present on the cell surface of PC tumor cells post-cleavage due to its transmembrane region. In a study, mice immunized with MUC4 glycopeptides mixed with tetanus toxin induced strong immune responses and predominantly produced IgG1 antibodies [19]. Such “cherry-picked” immunodominant peptides limit the epitopes that can be employed to elicit immune response in an unbiased manner, are of limited translation values. While large size of MUC4 can potentially provide a large epitope repertoire for eliciting potent immune response, it also makes the production and purification of intact protein equally challenging. We thus investigated the utility of recombinant MUC4 β -domain for cancer vaccine development. To circumvent these, we investigated the utility of recombinant MUC4 β subunit for tumor vaccine development.

One of the major challenges of vaccine delivery vehicles is to ensure protein stability and release over a sustained period in circulation [20, 21]. Amphiphilic polyanhydride nanoparticles, composed of 1,8-bis(*p*-carboxyphenoxy)-3,6-dioxaoctane (CPTEG) and 1,6-bis(*p*-carboxyphenoxy)hexane (CPH), have been shown to stabilize the structure and activity of encapsulated proteins while providing sustained release via a surface erosion mechanism [22, 23]. Furthermore, these nanoparticles have been shown to be readily internalized by antigen presenting cells (APCs), such as dendritic cells and macrophages, leading to the upregulation of surface activation markers including major histocompatibility complexes class I and II (MHC-I & MHC-II), co-activating ligands (CD86, CD40), secretion of inflammatory cytokines and generation of humoral responses [24-27].

In the present study, we encapsulated endotoxin free recombinant human MUC4 β in 20:80 CPTEG:CPH nanoparticles (MUC4-nanovaccine). We investigated the relationships between antigen release kinetics, immunological activity in terms of APC activation, and induction of humoral responses by the MUC4-nanovaccine. Our study demonstrates that the MUC4-nanovaccine activated mature DCs, eliciting a Th1 type of immune response. We further observe that MUC4 nanovaccine-immunized mice generate more IgG2b anti-MUC4 β serum antibodies than IgG1 antibodies, suggesting that MUC4-nanovaccine elicits a predominantly Th1-type response. Therefore, recombinant human MUC4 β -based polyanhydride nanovaccine has the potential to be an effective immunotherapy against pancreatic cancer and other MUC4 overexpressing malignancies.

3. Results

3.1 Encapsulation of MUC4 β into Polyanhydride Nanoparticles Provides Sustained Antigen Release Kinetics

The 20:80 CPTEG:CPH nanoparticles loaded with 3 wt. % MUC4 β were synthesized via solid-oil-oil double emulsion. Scanning electron microscopy showed the nanoparticles to be relatively spherical with a geometric mean diameter of 147 nm (with a geometric standard deviation of 1.3) (**Figure 1A**). The release kinetics of MUC4 β from 20:80 showed a burst of ca. 20% at early time points followed by slow, sustained release with smaller amount of protein released over 30 days. The data showed that after one month, the hydrophobic 20:80 CPTEG:CPH particles released ca. 25% of the encapsulated protein in a near-zero order release profile, which was consistent with previous work on protein release kinetics from CPTEG:CPH polyanhydrides formulations [22, 23, 27-29]. Finally, the encapsulation efficiency of the MUC4 β was determined to be $32 \pm 1\%$ (**Figure 1B**).

3.2 MUC4 Nanovaccine Enhances Surface Expression of MHC and Co-Stimulatory Molecules on DCs

While the functional role of MUC4 in pancreatic cancer pathobiology has been studied extensively, the immunogenicity of MUC4 protein has not been assessed until now. To examine the antigenicity of MUC4 β protein (MUC4) and characterize the potential of MUC4 nanovaccine in activating CD86⁺ CD11c⁺ DCs (**Supplementary Figure 1**), flow cytometry was used to measure expression of cell surface markers such as major histocompatibility complex molecule class II (MHC II) and class I (MHC I), co-stimulatory molecule CD80, and C-type lectin CD205 (DEC-205: DC maturation marker). Recombinant MUC4 β protein alone or delivered with empty nanoparticles did not upregulate surface expression of MHC I and II on DCs over controls. However, a significant 4-fold increase in mean fluorescence intensity (MFI) of MHC I and II was observed on DCs cultured with MUC4 nanovaccine in contrast to unstimulated DCs. Moreover, DCs cultured with MUC4 nanovaccine expressed high levels of MHC II (2-fold higher) than LPS-stimulated DCs (**Figure 2 A & B**). Furthermore, MUC4 nanovaccine enhanced the DC surface expression of the co-stimulatory molecule CD80 by 75% when compared to MUC4 only treated DCs, but no significant difference was observed in CD40 expression (**Figure 2 C & D**). In addition, DC surface expression of CD205 was 60%, 20% and 25% higher on MUC4-nanovaccine stimulated DCs when compared to unstimulated, LPS- and MUC4 β protein-treated DCs (**Figure 2E**) suggesting a higher proliferation and maturation of DCs which is shown in **Figure 2F**. Altogether, these results demonstrate that the MUC4 nanovaccine significantly enhanced the expression of surface markers and co-stimulatory molecules involved in DC maturation and antigen presentation.

3.3 MUC4 Nanovaccine Induces Pro-Inflammatory DC Cytokine Secretion

Dendritic cells direct immune responses by not only cross-presenting antigens to effector T and B cells, but also by secreting an array of cytokines to modulate these responses. After culturing DCs with various treatment groups for 48 h, we observed that the MUC4 nanovaccine significantly enhanced DC secretion of the cytokines IL-12p40, IL-6, and IFN γ in comparison to untreated DCs and free MUC4 simulated DCs (**Figure 3**). The amounts of IL-12p40 and IL-6 in culture supernatants of DCs treated with MUC4 nanovaccine were 40% and 30% higher than that of DCs stimulated with LPS respectively (**Fig 3A & B**), and the levels of IFN γ were comparable between these two stimulants (**Fig 3C**). DCs treated with MUC4 β alone (MUC4) or MUC4 β mixed with empty nanoparticles (MUC4+NP) expressed low or undetectable levels of cytokines, which were no different from unstimulated DCs. Similar to the data obtained for surface expression of MHC II and DC co-stimulatory molecules, encapsulation of MUC4 β protein in 20:80 CPTEG:CPH nanoparticles significantly enhanced DC cytokine production.

3.4 Single Immunization with MUC4 Nanovaccine Elicit Robust Anti-MUC4 Humoral Responses

Polyanhydride nanovaccines have been shown to generate germinal centers and B cells thus leading to sustained serum antibody responses in a single dose [30]. Apart from measuring antigen-specific antibody levels (which indicates the degree of humoral stimulation), isotyping of antibodies (IgG1 and IgG2b levels) demonstrate the type of immune responses (Th1 or Th2) generated through the determination of IgG2b:IgG1 ratios. The presence of high levels of antigen-specific IgG2b antibodies over IgG1 antibodies indicates preferred isotype-switching to Th1 type responses, whereas low IgG2b:IgG1 ratio is indicative of a Th2 type response [31]. To investigate if a single immunization with MUC4 nanovaccine induced robust humoral immune responses, animals were immunized

subcutaneously with MUC4 β -loaded 20:80 CPTEG:CPH nanoparticles. Serum antibodies were detected at 1:1000 serum dilution in mice that were immunized with MUC4 β alone, MUC4 β mixed with empty nanoparticles and MUC4 nanovaccine. Antibodies against MUC4 β were detected at higher dilutions in mice administered MUC4 β only and MUC4 nanovaccine, but not when free protein was mixed with empty particles (**Figure 4A**).

Since it has been shown that MUC4 peptides induce primarily IgG1 Th2 antibodies and not IgG2b [19], we evaluated the isotypes of the antibodies in MUC4 β -immunized mice. Mice immunized with the MUC4 nanovaccine demonstrated a high IgG2b:IgG1 ratio, whereas mice immunized with MUC4 β alone had a low IgG2b:IgG1 antibody ratio (**Figure 4B**). These results indicate that encapsulation of MUC4 β protein into 20:80 CPTEG:CPH nanoparticles modulates the immune response towards a Th1 phenotype, which could likely provide anti-tumor protection.

3.5 Immunization with MUC4 Nanovaccine Enhances the Presence of Inflammatory Cytokines in Sera

Since the analysis of antibody isotypes demonstrated an induction of a Th1 immune response, we also investigated the presence of Th1 cytokines (IL-12p40, IL-6, IL1 β , and IFN γ) in sera of immunized mice. Sera from MUC4 nanovaccine immunized mice had a slightly higher increased mean amount of IL-6 compared to MUC4 and MUC4+NP immunized mice (**Figure 5A**). In addition, sera from MUC4 nanovaccine immunized mice had significant amounts (~50 fold increase) of IL1 β Th1 cytokine when compared to other treatment groups and PBS-treated mice (negative control) (**Figure 5B**) that correlates with Th1 isotype switching of antibodies in these mice. The amounts of IL-12p40 and IFN γ were below detection levels in all treatment groups (data not shown).

4. Discussion

To date vaccine studies involving mucins have been based on selected peptides which have limited repertoire of immunogenic epitopes. In pre-clinical studies, these vaccines have not shown promising results which could be attributed to studies done only with the tandem repeat regions of mucins including MUC4. Recombinant proteins could address these limitations by presenting the entire spectrum of possible epitopes present on the original antigens in an unbiased manner [3]. In this study, the β -subunit of MUC4 was expressed in a bacterial system and its immunogenicity was investigated. The data presented showed that MUC4 β (MUC4) induces a Th2 type immune responses such as low expression of MHC-I and II complexes and co-stimulatory molecule CD80 (**Fig 2 A-C**), and low levels of inflammatory cytokine generation by pulsed DCs (**Fig 3 A-C**). In addition, mice immunized with only MUC4 produced significantly high levels of Th2 IgG1 antibodies to MUC4 (**Fig 4B**). This indicated that utilizing the free protein by itself will likely not provide an effective immunotherapeutic response. Th2 immune responses have been well established to promote tumor pathogenesis and aggressiveness, whereas shifting the immune response to Th1 phenotype provides anti-tumor protection [32-34]. Thus, we encapsulated MUC4 β into 20:80 CPTEG:CPH nanoparticles (MUC4 nanovaccine) and investigated whether the MUC4 nanovaccine could enhance activation of dendritic cells and modulate Th1 type humoral responses *in vivo*.

The MUC4 nanovaccine enhanced surface expression of MHC I and MHC II in CD86⁺ CD11c⁺ DCs (**Figure 3 A-B**), which are implicated in presentation of antigen to T cells and B cells, along with CD80 (**Figure 3C**), a co-stimulatory molecule required for activation of naïve CD4 helper T cells. Proper antigen presentation followed by secondary activation signal provided by CD80 is crucial in programming effector immune cells to specifically target the cancer cells. Upregulation of these markers by MUC4-nanovaccine,

when compared to the free MUC4 pulsed DCs, suggests that the encapsulated formulation was able to reprogram the DCs from Th2 to a Th1 response phenotype. Further, we observed that only stimulation with MUC4-nanovaccine enhanced DC secretion of the cytokines IL-12p40, IL-6 and IFN γ *in vitro* when compared to free MUC4 or free protein mixed with empty nanoparticle (MUC4+NP) that corroborated with the DC activation marker expression data. These results indicate that the encapsulation of MUC4 β protein in CPTEG:CPH nanoparticles is crucial to enhancing the immunogenicity of recombinant MUC4 β . Thus, the result suggests that the encapsulation of MUC4 β protein in CPTEG:CPH polymers is important for modulating the immunogenic response of Th2-inducing recombinant MUC4 β .

Previously, it has been shown that a single immunization of polyanhydride nanovaccines can induce high antibody titers and provide protective immunity against multiple pathogens in mice [28, 35]. Additionally, it is important to consider the quality of antibody response generated by the nanovaccine, which may be characterized by the specificity, avidity and isotype profile of the antibody response [36]. It is therefore noteworthy that MUC4-nanovaccine immunized mice had the highest IgG2b:IgG1 ratio which indicates Th1 type antibodies, whereas MUC4 β alone or in combination with blank nanoparticles preferentially induced Th2 type IgG1 anti-MUC4 β antibodies in alignment with its immunogenic nature (**Figure 4B**). This observation supports our *in vitro* observation that encapsulation of MUC4 β into polyanhydride nanoparticles likely plays a crucial role in activating dendritic cells in favor of Th1 type immune responses, which was further validated with the detection of higher levels of IL-6 and IL-1 β cytokines (**Figure 5**). The data herein shows encapsulating MUC4 β in 20:80 CPTEG:CPH nanoparticles is an effective strategy to activate dendritic cells in MUC4-specific manner and modulate the response towards an anti-tumor Th1 phenotype. The presence of IgG2b antibodies could

possibly further provide immunity against MUC4-expressing tumors by inducing antibody-dependent cellular cytotoxicity (ADCC) or antigen-mediated tumor killing by NK cells that can recognize the FcR region of antibodies bound to tumor cells [37, 38]. The current studies provide a basis for investigating the use of the MUC4 nanovaccine as an immunotherapeutic strategy in cancer models that overexpress MUC4 as a tumor-associated antigen.

5. Conclusion

Our data have clearly demonstrated that the MUC4-nanovaccine enhances DC surface expression of both MHC molecules and co-stimulatory ligands, and Th1 cytokine secretion. Further, in *in vivo* studies, the MUC4 nanovaccine effectively induced production of anti-MUC4 β antibodies and isotype-switching of these antibodies to primarily IgG2b (Th1 type) isotypes, which typically correlates to anti-tumor immune responses [39]. Thus, this work demonstrates that polyanhydride MUC4 nanovaccines are a promising platform for immunotherapies against pancreatic cancer.

Acknowledgments

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Figure 1: Synthesis, encapsulation and release kinetics of MUC4 nanovaccine.

Endotoxin-free recombinant MUC4 β protein was isolated from Rosetta bacteria and purified by affinity chromatography. 20:80 CPTEG:CPH nanoparticles encapsulating 3% MUC4 were synthesized via a solid/oil/oil double emulsion flash nanoprecipitation process. SEM images of 20:80 CPTEG :CPH nanoparticle encapsulated recombinant mucin fragment: MUC4- β (**A**). Antigen release kinetics were characterized by incubating the nanoparticles in PBS and measuring MUC4 released at regular intervals with a microBCA assay. 3% MUC4-loaded 20:80 CPTEG:CPH nanoparticles exhibited an initial burst (20%) release of protein followed by sustained release. The encapsulation efficiency of protein was determined to be 32%, suggesting low affinity between the polymer and MUC4 protein.

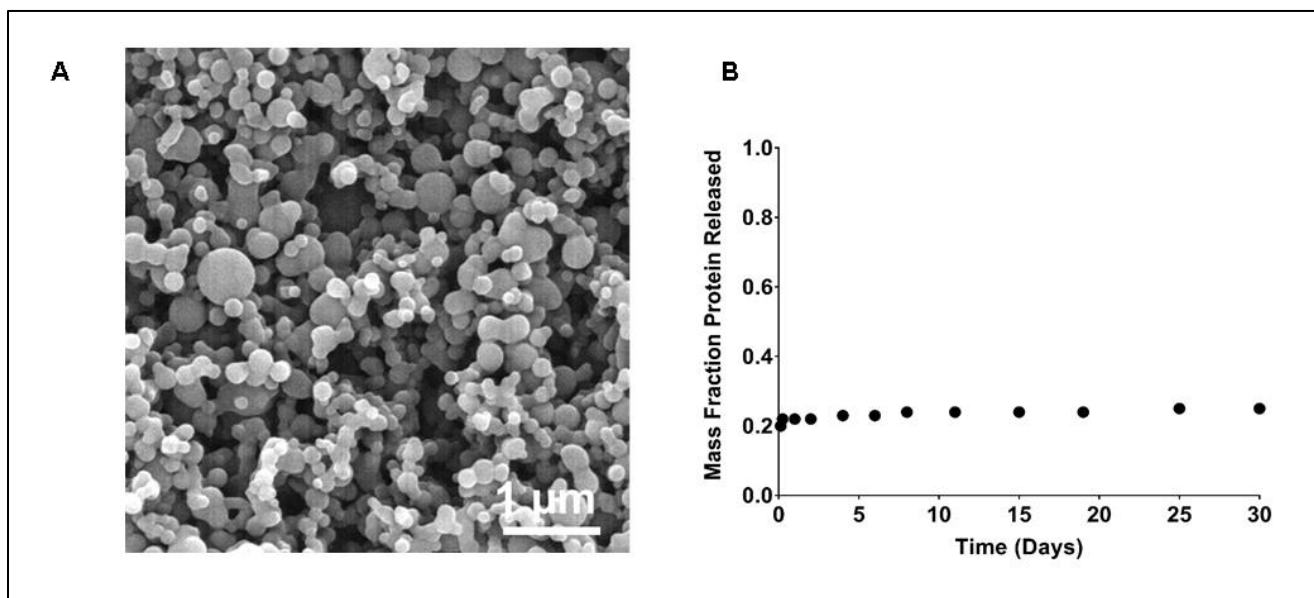
Figure 1

Figure 2: MUC4 nanovaccine activates DCs and induces expression of MHC II and co-stimulatory molecules.

Flow cytometry analysis demonstrated that MUC4 nanovaccine have activated DCs robustly like the LPS positive control. MUC4 β (MUC4) protein has Th2 immunogenic activity that abrogates DC activation and expression of MHC and co-stimulatory molecules. Free MUC4 mixed with empty nanoparticles (MUC4+NP) slightly increases the immunogenicity of MUC4 β , but only MUC4 nanovaccine could significantly activate DCs, suggesting encapsulation of MUC4 β is crucial for inducing a Th1 immune responses **(A-E)**. Increased DC proliferation was observed in MUC4 nanovaccine group compared to unstimulated DCs and other MUC4 treatment groups **(F)**. Statistical significance was set at $p < 0.05$. ANOVA analysis of the data was $p < 0.05$. Statistical comparison between MUC4, MUC4+NP & US with MUC4 nanovaccine is denoted by #, * & + respectively.

Figure 2

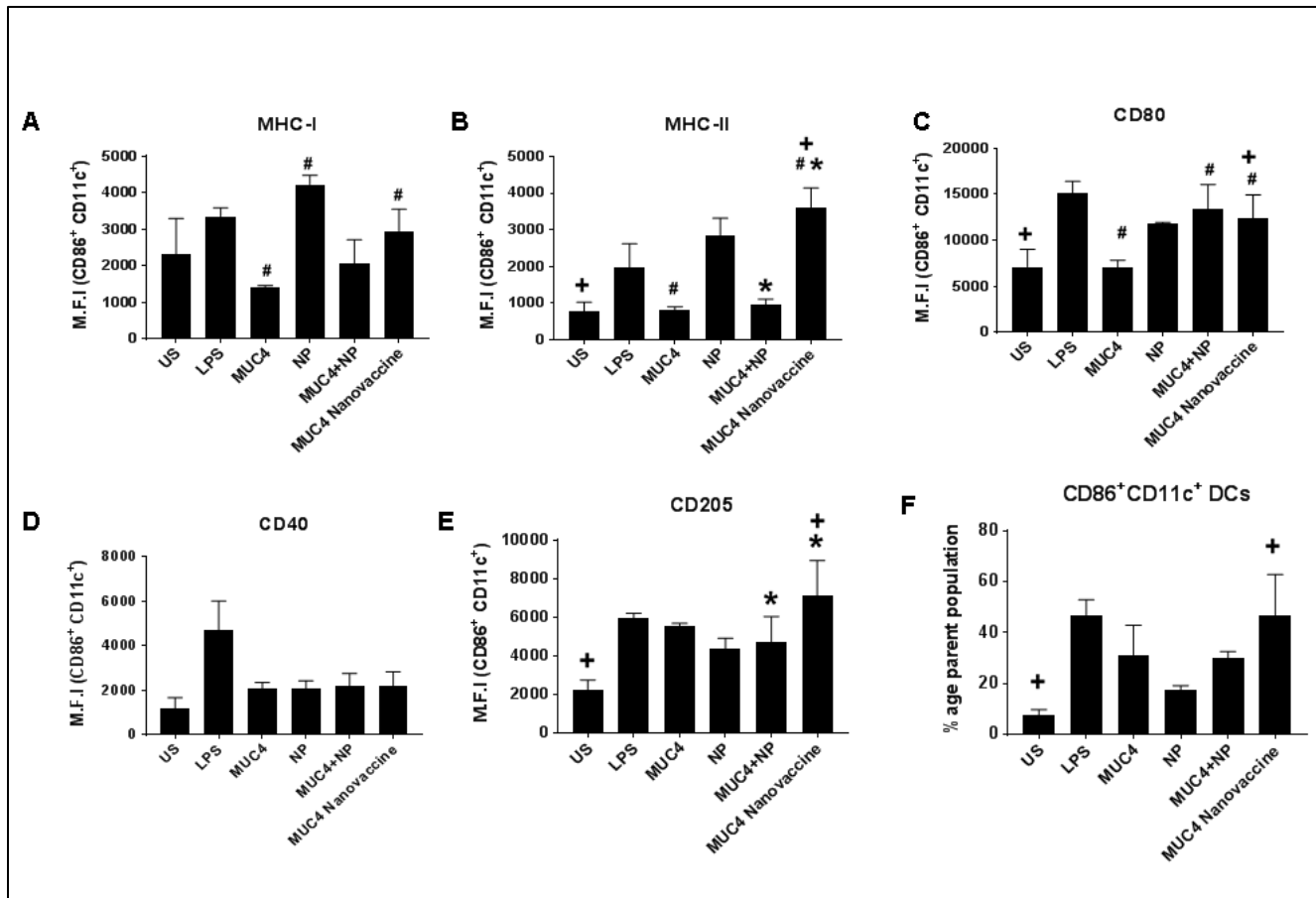


Figure 3: MUC4 nanovaccine induced robust Th1 DC cytokine secretion.

Cytokine analysis demonstrated that MUC4 nanovaccine activated DCs secrete Th1 cytokines. Free MUC4 mixed with empty nanoparticles (MUC4+NP) and MUC4 pulsed DCs have low expression of Th1 cytokines, suggesting that mixing of a strong adjuvant was not able to modulate the immune response induced by MUC4. Only encapsulation of MUC4 could modulate and reprogram DCs to secrete IL12, IL6 and IFN γ Th1 cytokines **(A-C)**. Statistical significance was set at $p < 0.05$. ANOVA analysis of the data was $p < 0.05$. Statistical comparison between MUC4, MUC4+NP & US with MUC4 nanovaccine is denoted by #, * & + respectively.

Figure 3

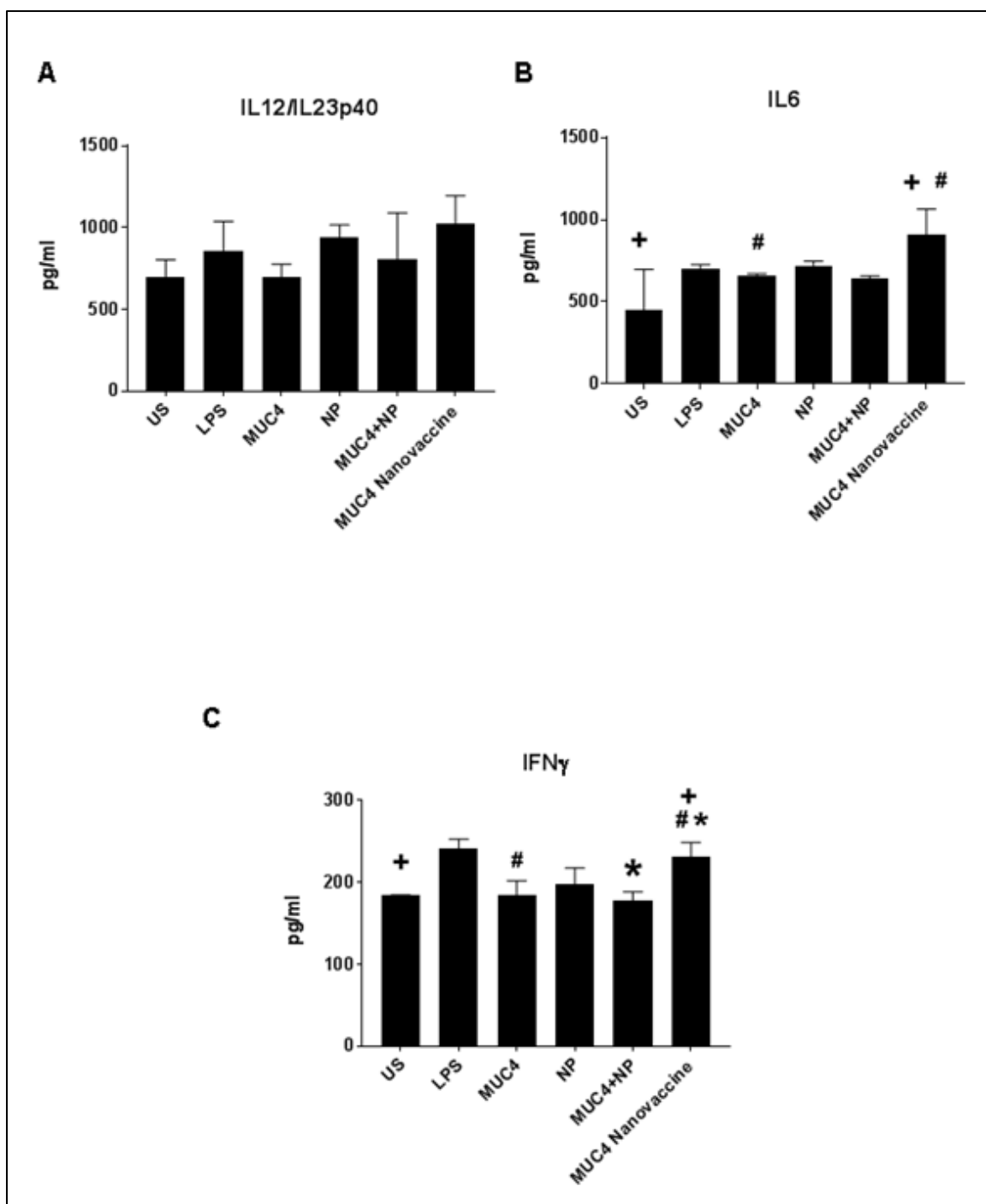


Figure 4: Mice immunized with MUC4 β nanovaccine generated anti-MUC4 β Th1 humoral response.

Eight week-old C57BL/6 mice were immunized with a single dose and serum was collected for detecting antibodies. ELISA studies showed that MUC4, MUC4+NP and MUC4 nanovaccine immunized mice serum carried MUC4 β antibodies **(A)**. Further, isotyping of MUC4 β antibodies demonstrated that isotype switching was predominantly to Th1 IgG2b in MUC4 nanovaccine immunized mice, whereas in MUC4 immunized mice it was predominantly in IgG1 Th2 isotype **(B)**. Statistical significance was set at $p < 0.05$. ANOVA analysis of the data was $p < 0.05$. Statistical comparison between MUC4, MUC4+NP and US with MUC4 nanovaccine is denoted by #, * & + respectively.

Figure 4

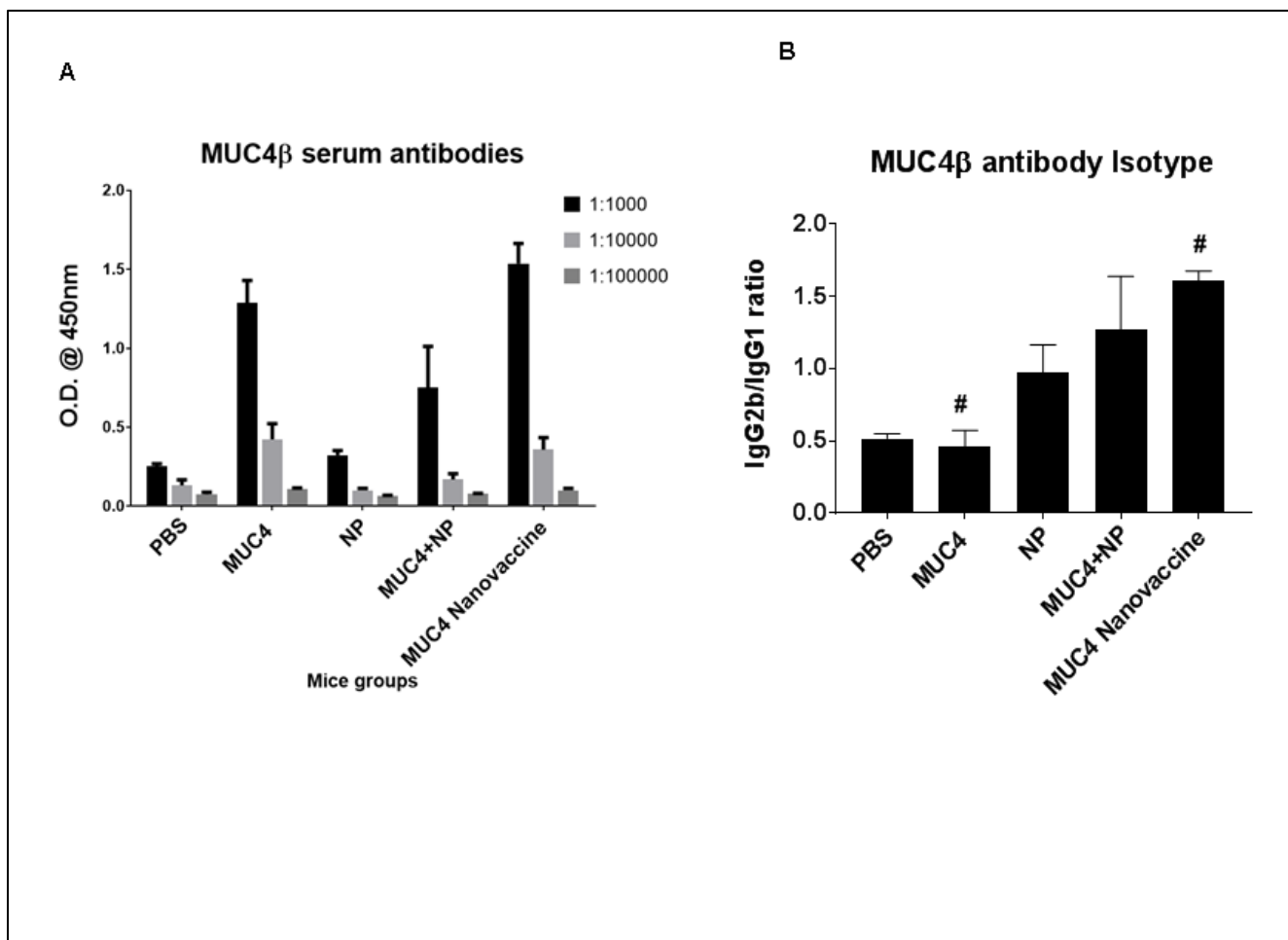
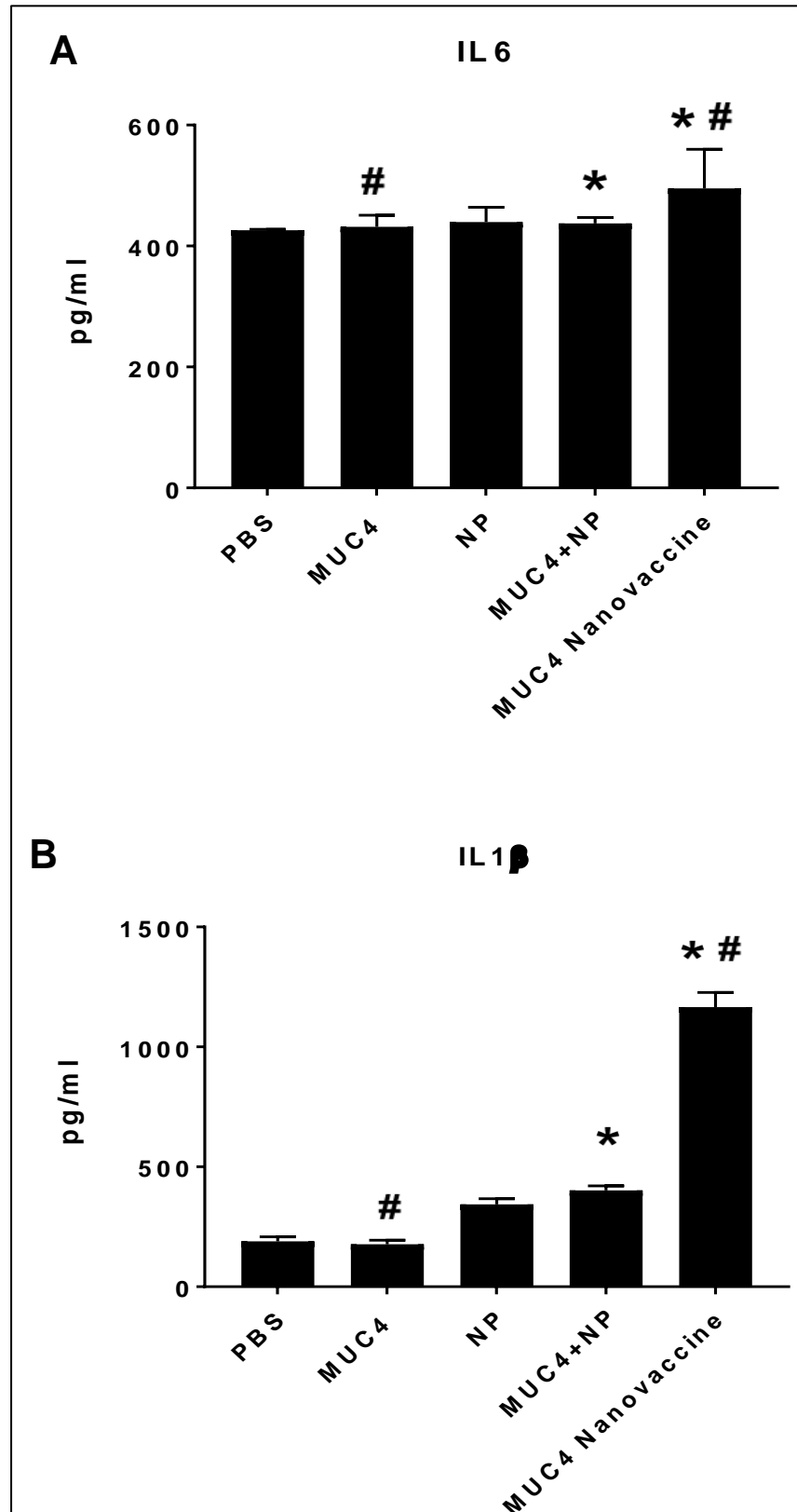


Figure 5: Presence of Th1 cytokines in mice immunized with MUC4 β nanovaccine.

MUC4 nanovaccine immunized mice serum had significantly higher levels of IL6 **(A)** and IL1 β **(B)** Th1 cytokines that correlated with the predominant Th1 IgG2b MUC4 β antibodies present in these mice. Statistical significance was set at $p < 0.05$. ANOVA analysis of the data was $p < 0.05$. Statistical comparison between MUC4, and MUC4+NP with MUC4 nanovaccine is denoted by # & * respectively.

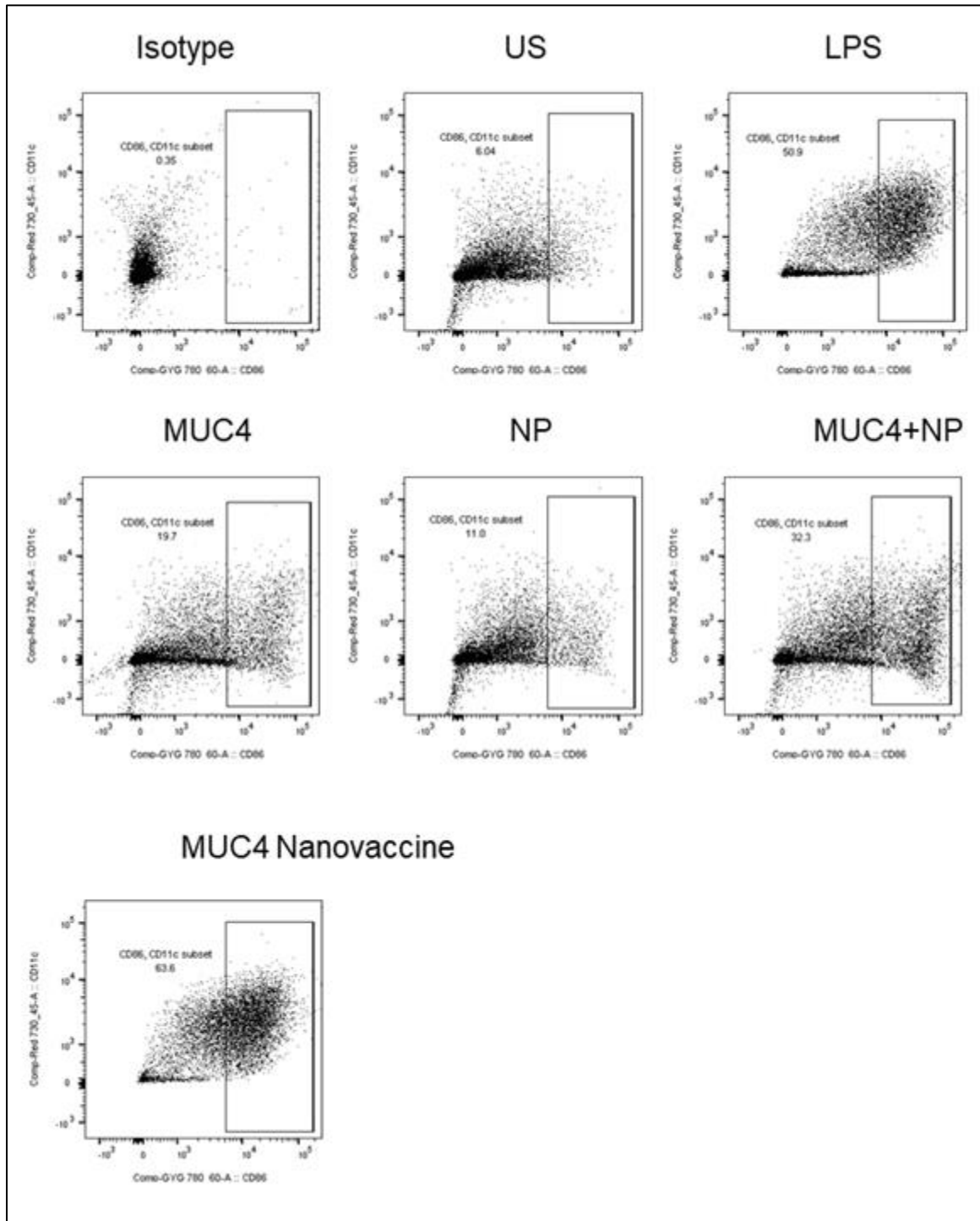
Figure 5



Supplementary Figure 1: MUC4 nanovaccine elicits robust proliferation of dendritic cells (DCs).

Total of 1×10^5 DCs were pulsed with free MUC4 β protein (MUC4), free MUC4 β protein mixed with empty nanoparticles (MUC4+NP) and MUC4-nanovaccine. Unstimulated DCs (US) and LPS-treated DCs serve as negative and positive controls. Antigen-matured DCs were characterized as CD86^{hi} CD11c positive (CD86⁺CD11c⁺) cells. MUC4 nanovaccine strongly stimulates DCs and robust 3-fold proliferation is observed when compared to MUC4-pulsed DCs.

Supplementary Figure 1



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**CHAPTER 5: COMBINING IMMUNE
CHECKPOINT BLOCKADE WITH
MUC4 NANOVACCINE FOR PC
TREATMENT**

1. Synopsis

Pancreatic cancer (PC) is a highly metastatic and therapy-resistant malignancy characterized by immunosuppressive tumor microenvironment (TME) mucin overexpression. Immunotherapy strategies for PC treatment are constantly being developed and tested in clinical trials but have achieved underwhelming outcomes. MUC4 is the most differentially overexpressed mucin and functionally contributes to PC disease aggressiveness. Our previous studies have demonstrated that MUC4 is immunogenic in PC patients and is thus an ideal candidate for targeted therapies. We developed a novel vaccine using recombinant MUC4 fragments and exploiting adjuvant-like properties of an amphiphilic polyanhydride-based nanoparticle delivery system. In addition, we showed that recombinant human MUC4 β protein encapsulated in amphiphilic polyanhydride nanoparticles (MUC4 nanovaccine) successfully activates dendritic cells and induces Th1 anti-MUC4 β antibodies in immunized C57BL/6 mice. Tumor cells are also known to express programmed death ligand 1 (PD-L1) to suppress effector immune cells activity via the PD-L1-PD-1 axis. Our preliminary study elucidated that MUC4 nanovaccine-immunized mice exhibited slower tumor growth kinetics than unimmunized control mice. However, we did not observe complete tumor regression and detected PD-L1 expression on MUC4-expressing tumors only. Based on this, we rationalized that PD-L1 expression by MUC4-expressing tumor cells suppressed and inhibited the therapeutic benefits of the nanovaccine in-vivo. The strong involvement of MUC4 in disease aggressiveness and PD-L1 in immunosuppression thus makes a compelling case for their combined targeting.

2. Background and Rationale

The PC tumor microenvironment is a complex relationship between cellular components, desmoplasia and cytokine milieu. PC's dense stroma/desmoplasia harbor immune cells such as TAMs, MDSCs, and T_{reg} cells that directly inhibit the activity of effector CD4⁺ and CD8⁺ T-cells. These pro-tumor immune cells further secrete anti-inflammatory Th2 cytokines like IL-4, IL-10 and IL-13, which negatively regulate the infiltration and functionality of effector immune cells by precluding immunological recognition [1-3]. To overcome the immune suppression elicited by the TME of pancreatic cancer, reprogramming of patient's immune system and generation of strong anti-tumor Th1 responses are necessary. Therapeutic cancer vaccines have emerged as a tool to reprogram and activate the patient's immune system in a tumor antigen-specific manner to effectively target tumor cells [4, 5]. Cancer vaccines have seen limited clinical success in some solid tumors such as lung cancer, breast cancer and renal cancer. A major factor that could improve the efficacy of these vaccines is firstly identifying TAAs that could be targeted [5]. Our previous study has identified MUC4 to be immunogenic in PC patients, suggesting that these patients have compromised peripheral tolerance.

A second major factor is a generation of robust anti-tumor responses to cancer vaccines to overcome peripheral immune tolerance and escape immune suppression of TME. To break tolerance and constrain immune suppression, selection of a strong immunological adjuvant to be combined with the antigen is required [5-7]. Adjuvants are an instrumental component of a potent vaccine that enhances the immunogenicity of the antigen and increases the antigen-specific immune response. However, developing a cancer vaccine with a successful adjuvant is not easy because the adjuvant must preserve the antigen and needs to be safe, potent and economically viable [6, 7]. We developed a novel MUC4 nanovaccine by encapsulating human MUC4 β recombinant protein in

pathogen-mimicking amphiphilic polyanhydride nanoparticles. We observed that the MUC4 nanovaccine elicited robust activation of DCs by upregulating surface expression of DC activation markers, as well as secretion of Th1 cytokines like IL12, IL6 and IFN γ . Further, we elucidated that mice immunized with MUC4 nanovaccine carried high levels of Th1 IgG2b anti-MUC4 β antibodies circulating in their serum. These data provide initial evidence that MUC4 nanovaccine could potentially serve as a potent immunotherapy strategy to treat PC.

The PD-L1 co-inhibitory molecule expressed on tumor cells binds to its receptor programmed cell death protein 1 (PD-1) present on T cells and negatively regulates T-cell signaling and effector functions [8]. PD-1 is expressed only on antigen-experienced CD4 $^+$ and CD8 $^+$ T cells, as well as B cells, and is absent on resting T and B cells [9]. Its ligand PD-L1, also known as cluster of differentiation 274 (CD274) or B7 homolog 1 (B7-H1), is expressed on activated DCs and macrophages[10], as well as non-lymphoid tissues like cancer cells upon IFN γ stimulation [11]. PD-L1 is also expressed in peripheral tissues, thus suggesting its role in peripheral tolerance against self-reactive T and B cells, and may serve in regulating inflammatory responses at these sites [12]. PD-L1, upon binding to its receptor PD-1, inhibits T cell proliferation and its effector functions by inducing apoptosis. In addition, this axis promotes differentiation of CD4 $^+$ T cells into Foxp3 $^+$ regulatory T cells [13-15].

One of the major challenges that have emerged from recent preclinical studies is the counter PD-L1 mediated-suppression elicited on infiltrating effector lymphocytes by tumor cells, therefore inhibiting the efficacy of these vaccines [16-18]. Expression of PD-L1 by tumor cells is an indicator of an active immune interaction occurring between cytotoxic CD8 $^+$ T-cells and tumor cells. Blockade of PD-1 receptor on effector T-cells to its ligand PD-L1 by PD-1 inhibitors have shown some promise in rescuing anti-tumor effects

as been discussed in the following review [1]. However, PC patients show differential expression of PD-L1 in tumors due to the various degree of effector T-cells infiltration found in the TME [19]. Combined high percentage of infiltrating CD8⁺ T-cells and high PD-L1 expression is correlated with lymph node metastasis and poor survival of PC patients [20-22].

Based on the data, our working goal was to investigate whether MUC4 nanovaccine could provide immunity against MUC4 tumor-bearing mice. Our preliminary data suggest that MUC4 nanovaccine has the potential to induce anti-tumor responses. We observed a positive correlation between TILs and tumor regression. Accumulation of infiltrating CD8⁺ and CD4⁺ T cells was greater in mice receiving the MUC4 nanovaccine compared to soluble MUC4 delivered with blank nanoparticles (MUC4+NP), indicating the benefit of sustained availability of antigen via encapsulation. However, we didn't achieve complete clearing of the MUC4-expressing tumor and thus investigated the PD-L1 expression in them. We did observe that MUC4 tumors had high expression of PD-L1 on tumor cells whereas only low expression of PD-L1 was seen in vector control tumors. These results suggest that MUC4 nanovaccine combined with checkpoint blockade could have enhanced therapeutic potential.

3. Results

3.1 The Cytotoxic Killing of MUC4-Expressing Cell Lines by CTLs

miniMUC4 construct captures the entire structure of original human MUC4 protein but has approximately 90% of the VNTR region missing due to which it runs at a lower molecular weight of 250-300 kDa in 2% agarose SDS gel [23]. Our cytotoxicity assay demonstrated that T-cells activated *ex vivo* by MUC4 nanovaccine-pulsed DCs mediated 25% higher antigenic-specific killing of miniMUC4 expressing-KPC960 cells when

compared to T-cells activated by MUC4+NP-pulsed DCs (**Figure 1**). Based on this, we proposed to elucidate the therapeutic potential of MUC4 nanovaccine in *in vivo* model.

3.2 MUC4 Nanovaccine Immunized Mice have Slower Tumor Growth Kinetics

To fully understand the preventive and therapeutic role of MUC4 nanovaccine, we pre-immunized the mice with primary dose and first booster. After tumor cells implantation, once the tumor volume reached 100 mm³, we immunized the mice with a lower second booster dose (**Figure 2A**). We observed that miniMUC4 tumors had slower growth kinetics with respect to its contralateral vector control KPC960 tumors. In addition, miniMUC4 tumors in MUC4 nanovaccine-immunized mice had overall lower tumor volume when compared to the unimmunized control mice (**Figure 2B & C**).

3.3 MUC4 Nanovaccine Enhanced Immune Cell Infiltration in miniMUC4 Tumors

Our observation of slower tumor growth kinetics and overall smaller tumor volume of miniMUC4 tumors, when compared to vector control tumors, in MUC4 nanovaccine-immunized mice, made us curious to investigate the immune cell infiltration in miniMUC4 tumors. Hematoxylin and eosin staining of tumor tissue sections showed a MUC4-specific immune cell infiltration in miniMUC4 tumor cells in the MUC4 nanovaccine immunized mice (**Figure 3A**). We did observe some antigen-specific immune cell infiltration in miniMUC4 tumors of free soluble MUC4-immunized mice, but the percentage of infiltration was significantly lower than MUC4 nanovaccine-immunized mice. The pathologist noticed an area of necrosis proximal to the immune infiltrate region, which were quantified. We found a positive correlation between tumor necrosis and the degree of immune cell infiltration in miniMUC4 tumors. MUC4 nanovaccine-immunized mice had 5% higher tumor necrosis than other groups where the MUC4 antigen was provided (**Figure 3B**).

Further, we investigated the percentage of CD8⁺ and CD4⁺ T-cells infiltrated in miniMUC4 tumor cells by immunofluorescence and confocal microscopy (**Figure 4 A-B**). We observed that in MUC4 nanovaccine-immunized mice, 65% CD8⁺ and 25% CD4⁺ of infiltrating T-cells were present in miniMUC4 tumors, whereas only 33% CD8⁺ and 9% CD4⁺ of infiltrating T-cells were detected in miniMUC4 tumors of MUC4+NP-immunized mice (**Figure 4 C-D**). These data provide a rationale for the slower miniMUC4 tumor growth kinetics observed in MUC4 nanovaccine-immunized mice.

3.4 MUC4 Nanovaccine Induces IFN γ Mediated PD-L1 Expression on miniMUC4 Tumor Cells

Our data suggest that MUC4 nanovaccine is inducing immunological targeting of miniMUC4 tumors; however we did not achieve complete tumor regression. To overcome immunosuppression of PC tumor cells, it is well understood that cancer vaccines need to generate robust cellular responses as well as high levels of Th1 cytokines. To evaluate whether MUC4 nanovaccine was able to induce strong cytokine secretion, we analyzed through flow cytometry the levels of Th1 cytokines produced by effector T cells. Our analysis demonstrated that significantly high levels ($P < 0.01$) of Th1 CD4⁺ (Tbet-positive) T-cells were present in MUC4 nanovaccine-immunized mice. In addition, both CD8⁺ and CD4⁺ T-cells produced robust levels of IFN γ cytokine in contrast to unimmunized mice (**Figure 5 A-B**).

Survey of literature has elucidated that IFN γ secreted by effector T-cells could potentially induce PD-L1 expression by tumor cell as a counterattack mechanism and inactivate the functionality of cytotoxic T-cells [11]. Since our data showed that significantly higher levels of IFN γ are produced by effector T-cells, we were curious to understand whether IFN γ was inducing PD-L1 expression on miniMUC4 tumor cells that lead to the limited success of MUC4 nanovaccine in our *in vivo* mouse model. IHC staining of PD-L1

revealed that PD-L1 was expressed on the surface of miniMUC4 tumor cells and stroma from MUC4 nanovaccine-immunized mice (**Figure 5C**). There was some low-intensity stromal staining and no tumor cell staining was observed in vector control tumors, whereas cellular surface staining of PD-L1 was observed on miniMUC4 tumor cells, suggesting an active cross-talk between effector T-cells and tumor cells, which corroborates with the high degree of immune infiltration detected in miniMUC4 tumors removed from MUC4 nanovaccine-immunized mice.

4. Discussion

The present study investigated whether MUC4 nanovaccine could elicit anti-tumor responses and abrogate tumor growth. Studies from our lab have shown that MUC4 nanovaccine could potentially activate dendritic cells and humoral responses. But whether the cancer vaccine could provide cellular immunity against MUC4 expressing tumor cells needed to be investigated. Thus, we did a preliminary study to investigate the preventive and therapeutic potential of MUC4 nanovaccine in a subcutaneous PC tumor mice model. Our *ex vivo* experiments provided evidence that MUC4 nanovaccine could activate effector T-cells in MUC4-specific manner and induce robust cytotoxic killing of MUC4 expressing KPC960 cells. Based on this data, we investigated whether MUC4 nanovaccine could inhibit the growth of MUC4 expressing tumors and lead to complete tumor regression. Our results suggest that even though MUC4 nanovaccine could significantly reduce the tumor growth kinetics and tumor volume along with increased immune cells infiltration and Th1 cytokine production, complete tumor regression was not attained in these immunized mice.

One of the reasons for not being able to attain a complete regression was that this study was done in C57BL/6-FBP mixed background mice, due to which different HLA

haplotypes could lead to limited activation of cytotoxic T cells and recognition of tumor cells by CTLs. Another reason for the limited success was due to the immune suppression exhibited by the PD-L1 expressed by tumor cells on CTLs. We propose that IFN γ secreted by effector T-cells induced PD-L1 immunoinhibitory ligand expression on miniMUC4 tumor cell surface as an immunosuppressive response and thus induced inactivation of CTLs that resulted in reduced tumor volume but no complete regression.

To evaluate and elucidate the full potential of MUC4 nanovaccine as a potent immunotherapy strategy for PC, we need to study MUC4 nanovaccine in a syngeneic PC mice model. Further to fully replicate clinical settings, we are required to study the nanovaccine in human MUC4 transgenic genetically engineered PC mice model. Our data suggest that combining checkpoint blockade therapy with MUC4 nanovaccine could potentially alleviate the immunosuppression exhibited by PC tumor, which needs to be investigated in the future.

Figure 1: MUC4 nanovaccine pulsed DCs activated CTLs in an antigen-specific manner and induced cytotoxic killing of miniMUC4-expressing murine pancreatic cancer cell lines.

Spontaneous pancreatic cancer mouse model (KPC)-derived pancreatic cell line (KCT-960) was transfected with the Mini-MUC4 construct and was used for LDH cytotoxicity assay. The assay showed the specific killing of Mini-MUC4 expressing cells compared to vector control **(A)**.

Figure 1

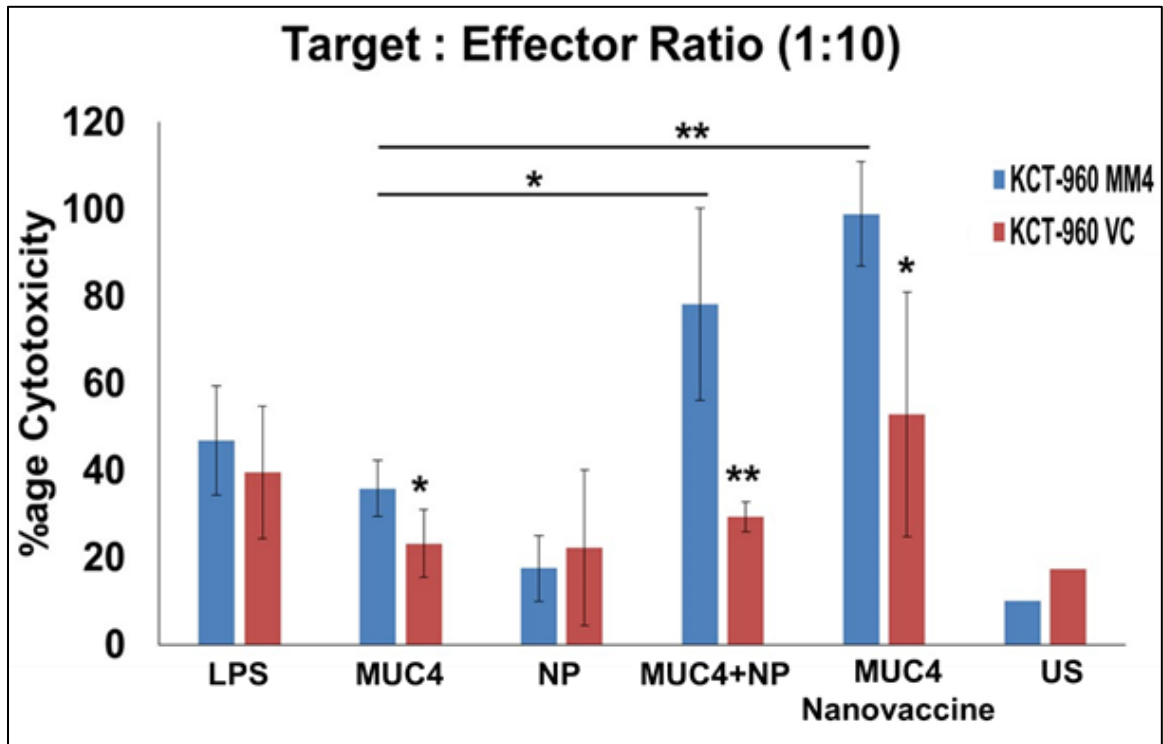


Figure 2: Schematic representation of tumor implantation and treatment by MUC4 nanovaccine. MUC4 nanovaccine-immunized mice showed a decrease in tumor growth.

Immunization of mice was done as indicated. Mice were injected with 1×10^6 cells of miniMUC4 or Vector expressing into right and left flanks of mixed background mice (n=5), respectively. Immunization of tumor-bearing mice to analyze the efficacy of the MUC4 nanovaccine **(A)**. Tumors were harvested at day 23. MUC4 nanovaccine-treated mice (inverted green triangle) showed tumor size reduction (below the red median line) in mini-MUC4 tumors with respect to contralateral vector control tumor, in comparison to unimmunized control mice (orange sphere) **(B)**. Representative photograph of tumors isolated from Control and MUC4 nanovaccine immunized mice group **(C)**.

Figure 2

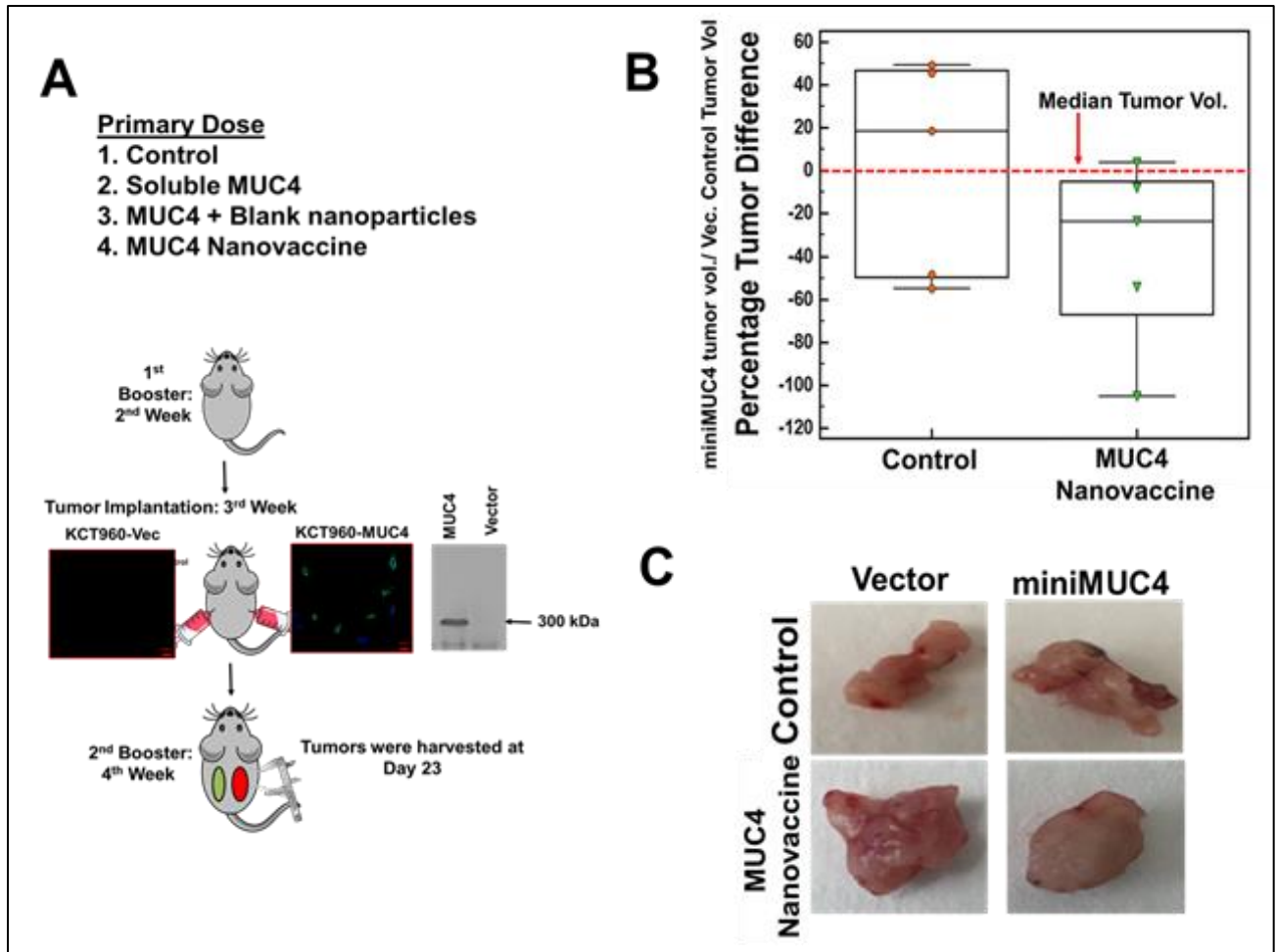


Figure 3: MUC4 nanovaccine enhanced immune infiltration and corresponding necrosis in the miniMUC4 tumor.

The miniMUC4- or vector-expressing KCT-960 cells were implanted into contralateral right and left flank of the same mouse, respectively. These mice were pre-immunized and received second booster post-tumor implantation, with MUC4 free protein mixed with empty CPTEG: CPH 20:80 nanoparticles (MUC4+NP) in PBS, MUC4 free protein in PBS and MUC4 nanovaccine in PBS as indicated in **Figure 2A**. Mice immunized with the MUC4 nanovaccine formulation demonstrated increased infiltration of immune cells **(A)**. Pathological analysis of tumor tissues showed that encapsulated MUC4 induced greater necrosis in MUC4-expressing tumors compared to all other treatment groups **(B)**.

Figure 3

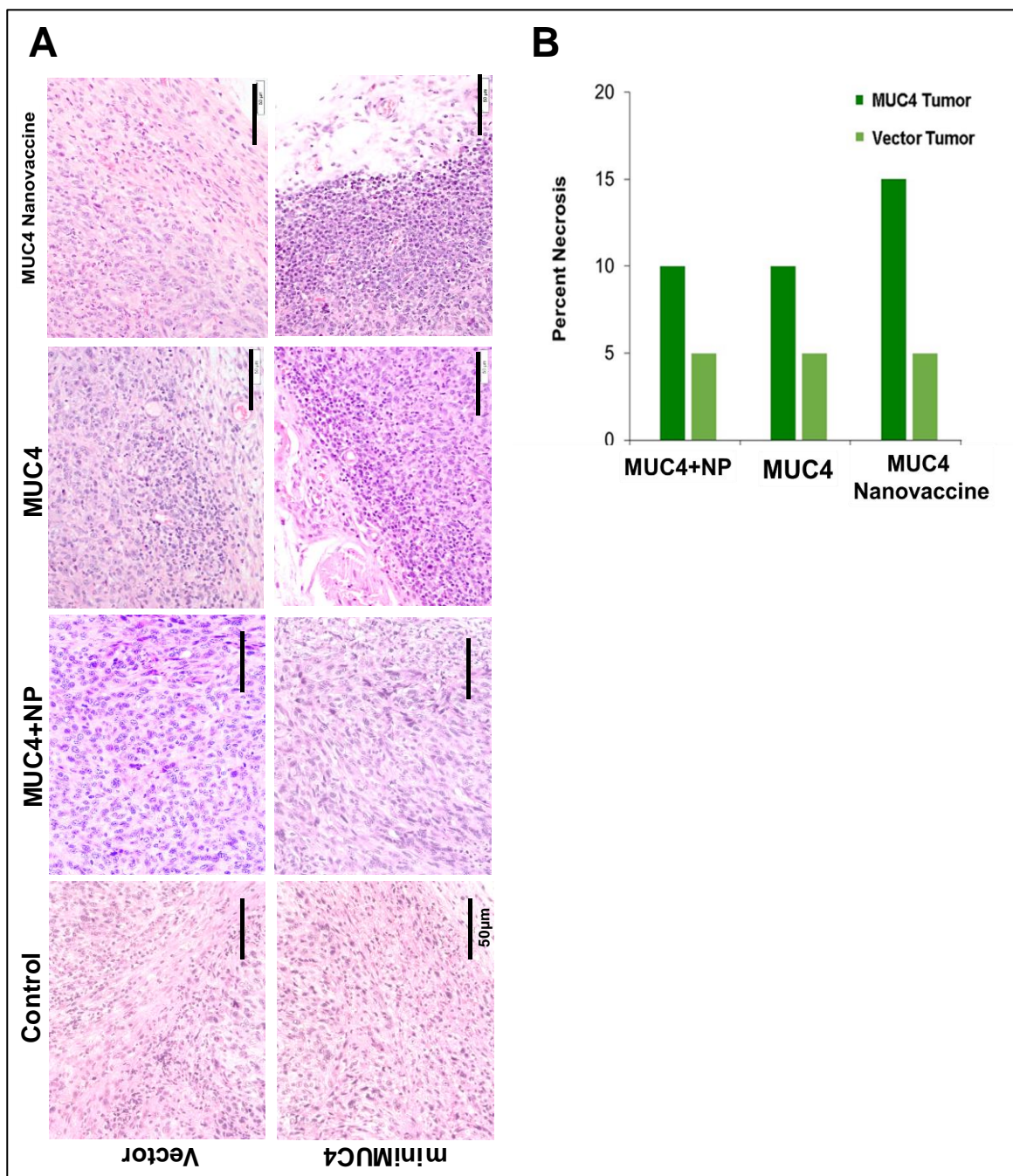


Figure 4: CD4⁺ and CD8⁺ T cell populations in miniMUC4 or Vector expressing mouse pancreatic ductal (PDAC) tumors.

Mice were pre-immunized and received second booster post-tumor implantation, with MUC4 free protein mixed with empty CPTEG: CPH 20:80 nanoparticles (MUC4+NP) in PBS, MUC4 free protein in PBS and MUC4 nanovaccine in PBS as indicated in **Figure 2A**. Tumor tissues were stained with CD8 (**A and B**) CD4 (**C and D**) T cells surface marker antibodies and were subjected to Immunofluorescence (IF) analysis by EVOS microscope. Quantification of infiltrating CD8⁺ and CD4⁺T cells in miniMUC4 or Vector expressing PDAC tumors in MUC4 nanovaccine- or miniMUC4+NP-immunized mice ($p < 0.01$ by Tukey's t-Test) (**B and D**).

Figure 4

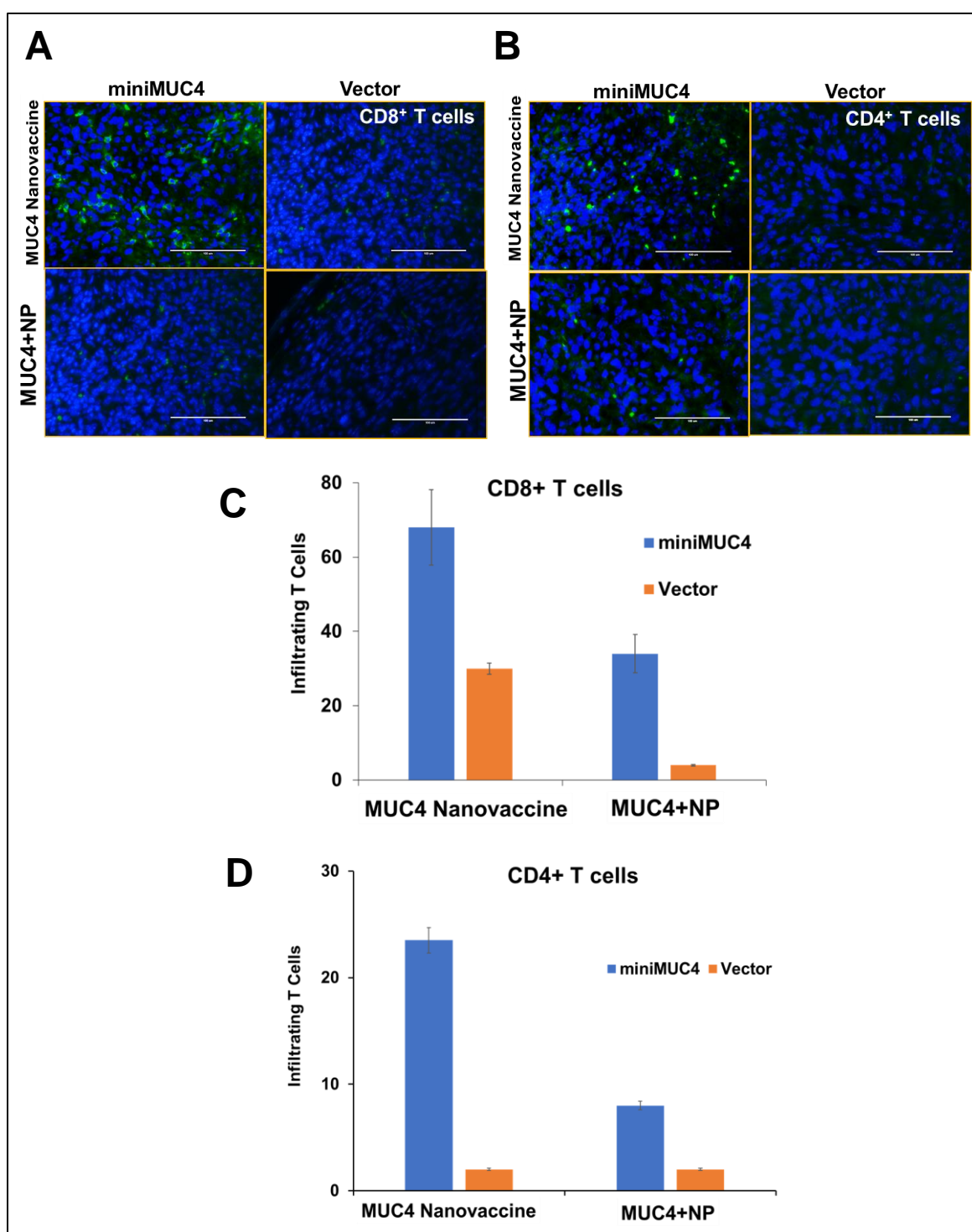
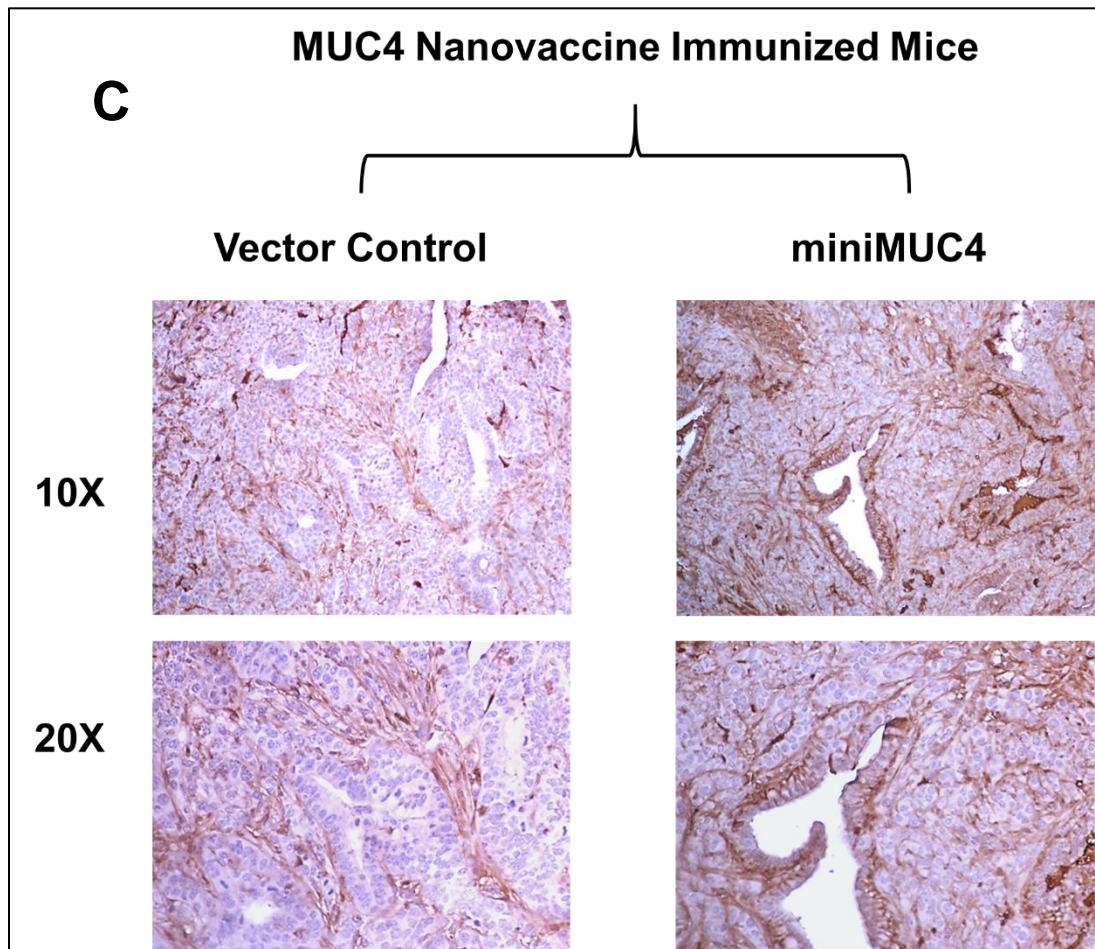
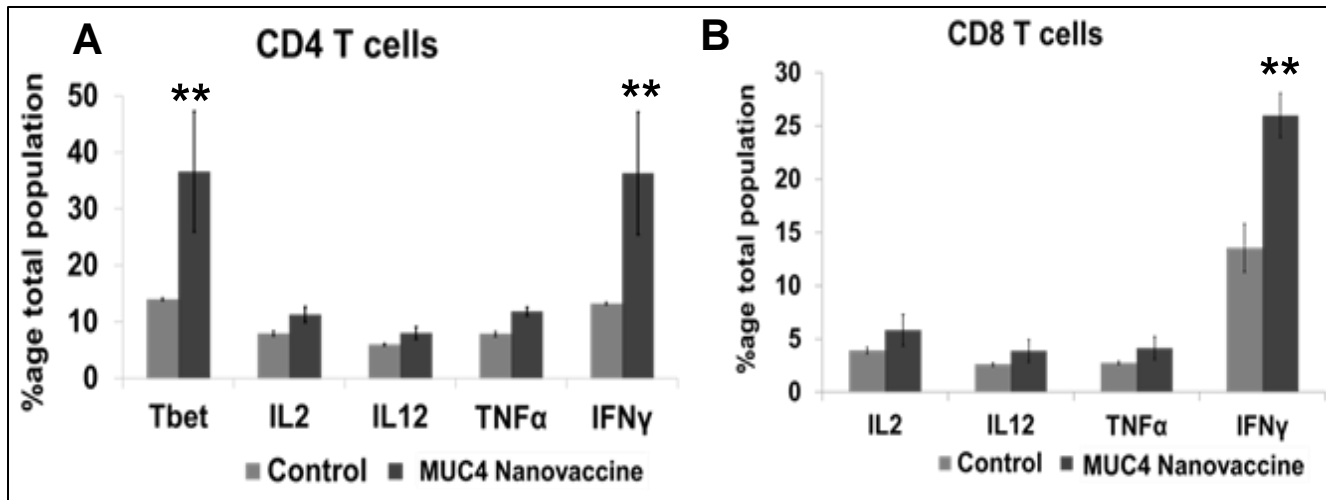


Figure 5: MUC4 nanovaccine immunized mice activates T-cells in Th1 phenotype that induces the corresponding expression of PD-L1 by miniMUC4 tumor cells.

Mice pre-immunized with MUC4 nanovaccine showed that MUC4-delivered through nanoparticles induced Th1 immune responses of both CD4⁺ and CD8⁺ T-cells in immunized mice compared to unimmunized control (** signifies $p < 0.01$ by Tukey's t-Test) **(A & B)**. IHC analysis for PD-L1 on MiniMUC4 tumors and vector control tumors treated with MUC4 nanovaccine showed an upregulation in PD-L1 expression only in MiniMUC4 tumors, indicating an immunosuppressive response by MiniMUC4 tumors **(C)**.

Figure 5



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**CHAPTER 6: SUMMARY,
CONCLUSIONS AND FUTURE
DIRECTIONS**

1. Summary and Conclusion

PC has an elaborate immunosuppressive microenvironment and immunotherapy has emerged as a tool to effectively target PC and its microenvironment [1]. Developing immunotherapy strategies for PC treatment comes with its own challenges that are: i) identifying a tumor-associated antigen that could be targeted, ii) identifying biodegradable adjuvants that could efficiently deliver antigens while preserving their antigenicity, iii) eliciting robust anti-tumor responses on the face of immunosuppression and iv) overcoming peripheral tolerance and immunosuppression elicited by the tumor.

Mucins are overexpressed in PC tumor cells and, due to their overexpression and aberrant glycosylation, they have emerged as potential candidates for targeted immunotherapeutic strategies for PC treatment [2]. MUC1 is one of the well-studied candidates for PC immunotherapy strategies but have attained only limited success in clinical trials. Most of these immunotherapies are MUC1 peptide-based vaccines that capture limited epitopes and excludes possible antigen epitopes present on the entirety of the protein. In addition, MUC1 is expressed at low levels in all normal tissues thus it doesn't provide tumor specificity [1-3]. Over the past several years, various aspects of MUC4 function and regulation in PC has been investigated [4-6]. In contrast to MUC1, MUC4 has restricted expression on normal tissues and is undetectable in normal pancreas [7, 8], whereas it is differentially overexpressed in PC tumor and its expression gradually increases with the disease progression [4, 8], thus MUC4 overexpression provides a spatiotemporal specificity to PC tumors that could be immunogenic. Further, MUC4 has been extensively studied and reported to be instrumental in PC pathogenesis [4], therefore we proposed that MUC4 could serve as a potential target for PC immunotherapy.

1.1 Pancreatic cancer patients have compromised peripheral tolerance against MUC4.

Here we focused on understanding whether the tolerance against MUC4 in PC patients is compromised. Peripheral tolerance is comprised of CD4 T helper cells and B-cells and identification of tumor-associated antigens activate B-cells to generate autoantibodies against those antigens. Thus detection of autoantibodies against MUC4 in PC patients will be an indicator of compromised peripheral tolerance in those patients. Our data demonstrated that autoantibodies are present against the recombinant MUC4 β protein as well as to randomly selected MUC4 peptides derived from the entirety of MUC4 protein. Further our analysis showed that the autoantibodies to MUC4 peptides are generated in a PC-specific manner and negligible in healthy or chronic pancreatitis individuals. Additionally, IgM autoantibodies to MUC4 peptides (A2, D3 and B1) significantly correlated with PC patient overall survival. Thus suggesting that immunologically targeting MUC4 through a unique immunotherapy platform could exploit this compromised tolerance and generate efficacious anti-tumor responses potentially contributing to clinical success.

1.2 Encapsulation of MUC4 β protein in novel amphiphilic polyanhydrides nanoparticles shifts its immunogenicity from Th2 to Th1 phenotype, and induces robust DC activation and humoral responses.

Most of the cancer vaccines developed are peptide-based due to the difficulty of encapsulating proteins while preserving their antigenicity and allowing sustained release in the circulation for activation of immune cells [9, 10]. Prior to our study, isolation and purification of MUC4 recombinant proteins and its characterization have not been achieved. Here we successfully purified the MUC4 β protein from bacterial expression

system, and encapsulated the protein in novel amphiphilic polyanhydrides nanoparticles to develop MUC4 nanovaccine. Polyanhydride nanoparticles is a suitable platform due to its pH-neutral core that stabilizes protein, pathogen-mimicking properties and tunable release kinetics that enable immune-modulation [11-16]. In addition, MUC4 recombinant protein provides a wide repertoire of potential epitopes when compared to single epitope peptide-based immunotherapy strategies thus potentially eliciting a robust and sustained anti-tumor Th1 immune responses.

We demonstrated that soluble MUC4 β protein alone activates DCs in the Th2 pathway. It is well understood that Th1 immune responses have anti-tumor outcomes, therefore the protein by itself is not suitable for PC treatment. However, encapsulation of MUC4 β protein shifted the overall response to Th1 phenotype. MUC4 nanovaccine robustly activated DCs and induced both surface expression of activation markers and secretion of Th1 cytokines. Further, in the immunized mice MUC4 nanovaccine induced generation of Th1 IgG2b humoral responses, which was not observed in mice immunized with soluble MUC4 or MUC4+NP groups. Thus indicating that the encapsulation of MUC4 β in the nanoparticles is crucial for reprogramming the immune responses from Th2 phenotype to Th1 phenotype.

1.3 MUC4 nanovaccine induces immune infiltration in miniMUC4 tumors in an antigen-specific pathway.

Cancer vaccines have been shown to induce effector immune cell infiltration in tumors in an antigen-specific pathway that affects the overall tumor growth [17]. Here we tried to elucidate the therapeutic potential of a MUC4 nanovaccine in eliciting robust anti-tumor responses in a PC subcutaneous tumor mouse model. Our study showed that miniMUC4 tumors in mice immunized with MUC4 nanovaccine had slower tumor growth and lower tumor volume with respect to vector control tumors on the contralateral flank of

the same mice. These data suggested a MUC4-specific immune response in these vaccinated mice. Upon further evaluation, we observed that only in MUC4 nanovaccine immunized mice there is enhanced total immune cell infiltration and also effector T cell infiltration which correlated with high levels of necrosis seen in miniMUC4 tumors. Despite high T-cell infiltration, we were unable to attain complete regression in miniMUC4 tumors. Due to the high levels of IFN γ generated by T-cells isolated from MUC4 nanovaccine-immunized mice, we hypothesized that tumor cells are expressing PD-L1 upon IFN γ induction on the cell surface and inhibiting effector T-cells, cytotoxic activity. Our IHC analysis validated our hypothesis and we observed PD-L1 surface expression on miniMUC4 tumor cells. Thus, in conclusion, MUC4 nanovaccine could potentially be exploited as a PC treatment strategy.

2. Future Directions

While our studies elucidated the potential of MUC4 nanovaccine as a potent strategy for treating pancreatic cancer, the fact that our *in vivo* tumor model study was done in mixed background mice limits the clinical translation of our work. To fully understand the anti-tumor potential of MUC4 nanovaccine we are required to perform our studies in a syngeneic mice model. Our lab has recently developed murine PC cell lines derived from the C57BL/6 pure background KPC mice model. We will be transfecting these cells with miniMUC4 plasmid construct and test the cytotoxic killing of this human MUC4-expressing syngeneic mouse PC cell lines by MUC4 nanovaccine-activated T-cells. This study will give us a better understanding of the potential of MUC4 nanovaccine activating T-cells in a MUC4 antigen-specific manner.

Even though our *in vitro* and subcutaneous study with human MUC4 expressing syngeneic mice model will elucidate the therapeutic potential of MUC4 nanovaccine, the

model does not recapitulate PC patient's immune system where immune cells might recognize MUC4 on tumor cells as self-antigen and are tolerant towards MUC4. Our autoantibodies study suggests that MUC4 is immunogenic in PC patients and based on serum reactivity some PC patients might respond well to MUC4 based immunotherapies. To recapitulate this clinical setting and validate our hypothesis, our lab (with the help of Dr. Satyanarayana Rachagani) has recently developed a human MUC4 transgenic mice model. We will utilize this model to primarily address two questions: i) whether MUC4 nanovaccine could break the immunological tolerance in MUC4 transgenic PC mice model and effectively kill MUC4-expressing PC tumor cells, and ii) whether MUC4 nanovaccine will lead to severe side-effects due to the non-specific recognition of MUC4 on other normal tissues. Further we have also developed a human MUC4 transgenic Kras^{G12D} Pdx-1 Cre spontaneous PC mice model that expresses MUC4 only on pancreatic tumor cells. This mice model will be used to study whether MUC4 nanovaccine could either stabilize or regress human MUC4 expressing spontaneous PC tumor. A culmination of all these studies will provide us with a clearer understanding of the efficacy of MUC4 nanovaccine as a therapy strategy.

There is very limited knowledge about MUC4's contribution to the PC immunosuppressive TME. We are curious to investigate the correlation of MUC4 expression with the immune-phenotype of resected tumors, and cross-refer the observation in our MUC4 transgenic Kras^{G12D} Pdx-1 Cre spontaneous PC mice model system. Further, PC tumors are fairly desmoplastic that restricts the infiltration of effector T-cells. Our *in vivo* work provided evidence that PD-L1 is expressed on miniMUC4 tumor cells as a counterattack to escape cytotoxic killing by cytotoxic CD8⁺ T cells. Thus, based on the data that we will collect from MUC4 transgenic mice models, combining immune-checkpoint blockade agents such as anti PD-1 antibody and stroma cell depletion agents

with MUC4 nanovaccine to enhance the therapeutic efficacy of the vaccine needs to be tested.

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