

ABSTRACT

Cancer cachexia is a multifaceted condition affecting a large majority of cancer patients, with pancreatic ductal adenocarcinoma (PDAC) patients presenting the highest prevalence. Cachexia is characterized by progressive whole-body and skeletal muscle wasting, including vital cardiorespiratory muscles. There are currently no medical treatments for cachexia, largely due to a limited understanding of the underlying mechanisms leading to the progression of the disease, especially with respect to cardiorespiratory muscles. The purpose of this study was therefore to characterize the time course of diaphragm muscle pathology and dysfunction in a preclinical model of PDAC. C57BI6/J mice (male, 10 weeks old) received orthotopic injections into the pancreas of either PBS (SHAM, n=6) or mouse pancreatic cancer cells (KPC, n=30). Mice were monitored daily and euthanized at predetermined time points corresponding to 8 (D8), 10 (D10), 12 (D12), and 14 (D14) days post cancer cell inoculation or when mice met IACUC-mandated tumor endpoint (i.e. (i.e. 15-17 days post tumor cell inoculation (END), body condition score < 2). In-vivo diaphragm function was assessed via M-mode ultrasonography immediately prior to euthanasia. Hemispheres of the diaphragm were harvested and subjected to immunohistochemical analyses. In-vivo diaphragm (excursion amplitude) and respiratory function (respiratory rate, minute ventilation) were significantly impaired beginning at D12 versus Sham. Additionally, significant diaphragm muscle fiber atrophy occurred at D12, D14, and END. Collagen remodeling, indicated by collagen hybridizing peptide reactivity, was increased by 113% at D12 versus Sham. Muscle area occupied by extracellular matrix was significantly increased by 52% D12 and persisted through END. Infiltration of CD45+ leukocytes significantly increased versus Sham at D8 and steadily increased through END. The abundance of infiltrating leukocytes significantly correlated to minute ventilation, muscle fiber size, collagen remodeling, and muscle area occupied by extracellular matrix. Overall, this is the first study to characterize the time course of cachexia development and progression in the orthotopic KPC model. The current findings demonstrate an early and profound pathological remodeling of the diaphragm in response to pancreatic tumor burden, characterized by muscle fiber atrophy, extracellular matrix remodeling, immune cell infiltration, and ultimately respiratory dysfunction. These data help define the early cellular events underlying the development of muscle pathology and the associated respiratory dysfunction, providing an important translation tool for therapeutic investigations.

BACKGROUND

- Cachexia is a devastating consequence of cancer, characterized by whole-body, fat, and skeletal muscle wasting
- Cachexia is most prevalent in patients diagnosed with pancreatic ductal adenocarcinoma (PDAC)
- The loss of skeletal muscle impacts both limb muscles and cardiorespiratory muscles, which may mediate respiratory dysfunction commonly reported in patients with advanced cancers
- No approved treatments for cachexia currently exist, partly due to a limited mechanistic understanding of its etiology

OBJECTIVE

Characterize the time course of diaphragm muscle pathology and dysfunction in a preclinical, orthotopic model of pancreatic cancer

METHODOLOGY

- C57BI6/J mice (10 weeks old, male) were randomly assigned to receive an orthotopic injection into the pancreas of either
- Sterile PBS (SHAM) or
- Pancreatic cancer cells (KPC), originally derived from Kras(G12D);Trp53(R172H);Pdx1-Cre (KPC) mice
- <u>KPC</u>: 2.5 x 10⁵ KPC cells in 50 μl PBS; <u>SHAM</u>: 50 μl PBS
- Mice were monitored daily until predetermined time points or until IACUC-mandated endpoint
- Skeletal muscle and tumor were dissected upon euthanasia



injections. * = different from Sham.

Pancreatic Tumor Burden Elicits Progressive Respiratory Muscle Pathology and Dysfunction

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pancreatic tumors of



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refractory/severe cachexia)



Collectively, these findings demonstrate a rapid and robust model to investigate mechanisms involved at different stages of cachexia (i.e. pre-cachectic, cachectic,