Endocrine-exocrine signaling drives pancreatic cancer progression Cathy Garcia (1,2), Lauren Lawres (1,2), Jaffarguriqbal Singh (1,2), Sherry Agabiti (1,2), Alex Tong (1), Smita Krishnaswamy (1), Pamela Kunz (3,4), Mandar Deepak Muzumdar (1,2,3,4) (1) Department of Genetics,, (2) Cancer Biology Institute, (3) Department of Medical Oncology, (4) Smilow Cancer Hospital, Yale University School of Medicine

INTRODUCTION

The predominant histologic subtypes of pancreatic cancer, pancreatic ductal adenocarcinoma (PDAC) and pancreatic neuroendocrine tumors (PNET) are thought to originate from independent cellular compartments comprised of the exocrine (acinar cells/ducts) and endocrine (islets) pancreas, respectively. Recent ultrastructural and perfusion studies have identified bidirectional endocrineexocrine blood flow, but whether and how intra-pancreatic endocrine-exocrine signaling contributes to PNET and PDAC progression is not known. Understanding these mechanisms could reveal novel strategies for PNET prevention and therapy.

MATERIAL & METHODS

We generated a new obesity-driven mouse model of pancreatic cancer, since obesity is a major host risk factor for both PDAC and PNET and places a significant physiologic stress on both the endocrine and exocrine pancreas. We crossed the leptin-deficient (ob/ob) model of obesity with an oncogenic Kras-driven model (KC: Pdx1-Cre; Kras^{LSL-G12D}) of PDAC, which faithfully mimics the genetic and histologic features of the human disease. We induced weight loss using adeno-associated virusmediated leptin gene therapy (AAV-Leptin) or caloric restriction (CR) and analyzed tumor progression by histology. We performed molecular analyses of mouse and human PDAC tumors and sera using exome sequencing, RNA-sequencing, immunohistochemistry, and ELISA and assessed islet gene expression in obese and non-obese conditions by single-cell RNA-sequencing (scRNA-seq).





