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Context-Dependent Transformation of Adult Pancreatic Cells by Oncogenic K-Ras

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SUMMARY
Pancreatic ductal adenocarcinoma (PDAC) is one of the most lethal human malignancies. To investigate the cellular origin(s) of this cancer, we determined the effect of PDAC-relevant gene mutations in distinct cell types of the adult pancreas. We show that a subpopulation of Pdx1-expressing cells is susceptible to oncogenic K-Ras-induced transformation without tissue injury, whereas insulin-expressing endocrine cells are completely refractory to transformation under these conditions. However, chronic pancreatic injury can alter their endocrine fate and allow them to serve as the cell of origin for exocrine neoplasia. These results suggest that one mechanism by which inflammation and/or tissue damage can promote neoplasia is by altering the fate of differentiated cells that are normally refractory to oncogenic stimulation.

INTRODUCTION
Pancreatic ductal adenocarcinoma (PDAC) is the fourth leading cause of cancer death in the United States (Li et al., 2004). Identification of the cell(s) of origin of PDAC and defining the biochemical and biological changes that accompany their transformation are essential for the development of early detection and treatment tools. The pancreas is composed of four parenchymal cell types: acinar, ductal, centroacinar (exocrine), and islet (endocrine) cells. The acinar cells synthesize and secrete zymogens into the ductal lumen. The ducts carry the enzymes to the duodenum and generate bicarbonate. The centroacinar cells (CACs) have been suggested to be progenitor cells in the adult pancreas. The islets of Langerhans are composed of endocrine cell types, which secrete insulin and other proteins that regulate glucose homeostasis (Bardeesy and DePinho, 2002). Rare cells expressing endocrine markers such as insulin can also be found associated with the acini and the ductal epithelium (Mills, 2007).

PDAC arises from in situ precursor lesions termed pancreatic intraepithelial neoplasia (PanIN) (Hruban et al., 2001). PanIN lesions include a spectrum of abnormal proliferative ductal structures (termed PanIN 1–3) that are recognized by transformation from a cuboidal to columnar epithelium, mucin production, and nuclear atypia. Mutations that result in a constitutively active K-RAS are found in over 95% of PDACs and are believed to be an initiating event for this type of cancer (Jones et al., 2008; Rozenblum et al., 1997). KRAS encodes a small GTPase that in its active GTP-bound form promotes a wide range of cellular responses including proliferation, survival, migration, and metabolism (Shields et al., 2000). The CDKN2A locus encodes the Ink4A and ARF tumor suppressor genes and is also frequently mutated in PDAC. INK4A inactivation mutations are present in virtually all

SIGNIFICANCE
The identification of the cell type(s) from which PDAC originates and is maintained is essential for the development of new methods for PDAC diagnosis and treatment. In this paper, we demonstrate that oncogenic K-Ras can transform Pdx1-expressing cells in the adult pancreas of the mouse in the absence of tissue injury. In contrast, insulin-expressing cells require injury in combination with K-Ras activation for transformation. Our study emphasizes the role of both genetic and nongenetic events in PDAC initiation and provides additional mouse models to aid in the development of new treatment and diagnostic strategies.
sporadic PDAC, suggesting that INK4A/ARF normally constrains the malignant potential of mutant K-RAS. Other frequent alterations include loss-of-function mutations of the SMAD4 and p53 tumor suppressor genes (Jaffee et al., 2002). Moreover, developmental regulatory pathways, in particular the Notch, Sonic Hedgehog, and Wnt pathways, are often activated in PDACs (Berman et al., 2003; Miyamoto et al., 2003; Thayer et al., 2003; Wang et al., 2009).

PDAC is commonly believed to arise from transformation of ductal cells, although experimental studies have suggested other differentiated cells or progenitor/stem cells as potential origin(s) for this type of cancer. For instance, mouse models expressing various oncogenes in acinar cells during development, such as the Elastase-Tgfα (Wagner et al., 1998) and Elastase-KrasG12D (Grippo et al., 2003) transgenic strains and the knockin Mist1-Kras4B transgenic animals (Tuveson et al., 2006), have implicated acinar cells as being susceptible to transformation and giving rise to tumors containing ductal elements. Transplantation of mouse islet cell cultures expressing the polyoma virus middle T oncogene into histocompatible mice resulted in the induction of pancreatic cancer with ductal histology (Yoshida and Hanahan, 1994), suggesting a potential endocrine origin for PDAC.

A role of CACs has been proposed based on studies of mice that harbor a pancreas-specific deletion of the PTEN gene; these animals exhibit a proliferative expansion of CACs that can progress to carcinoma (Stanger et al., 2005). Nestin+ exocrine progenitor cells have been shown to be highly responsive to K-Ras oncogenic activation and were suggested to represent a progenitor origin for PDAC (Carriere et al., 2007).

The close developmental relationships of the different pancreatic acinar cell types and the capacity of endodermal lineages to transdifferentiate (Slack, 1995) add complexity to the PDAC cell-of-origin question. PanIN lesions can express markers of gastric and foregut differentiation (Prasad et al., 2005), suggesting that some degree of transdifferentiation occurs routinely during PDAC progression. This developmental plasticity is evident in some human PDACs that show focal expression of nonductal lineage markers, including endocrine and exocrine factors. More recently, Habbe et al. (2008) and De La O et al. (2008) have shown that activation of oncogenic K-Ras in acinar cells of the adult mouse leads to efficient mPanIN formation. These groups have also demonstrated in vivo acinar-to-ductal transdifferentiation by lineage tracing studies in these models (De La O et al., 2008; Habbe et al., 2008). Given this plasticity, it is possible that there is no exclusive cell of origin for PDAC and that specific genetic alteration(s) define the resulting malignant phenotype regardless of the cell in which they occur. Moreover, the physiological context, including tissue damage and inflammation, may alter a given cell's susceptibility to transformation. For example, Guerra et al. (2007) have reported that K-Ras activation in acinar cells (using the K-Ras<sup>LSL-G12Dgeo;Ela1-Tα1/tetO-Cre</sup> system) resulted in adult onset malignancy only following treatment with caerulein, which causes tissue injury and inflammation (Guerra et al., 2007).

In this study, we investigate whether PDAC arises from an exclusive cell of origin and whether a combination of genetic and nongenetic events can collaborate to induce neoplasia in a range of cell types. We describe the use of genetic manipulation of the mouse to systematically determine the effects of oncogenic K-Ras in distinct subsets of pancreatic cells of the adult mouse in a context-dependent manner.

**RESULTS**

**Temporally Restricted Activation of Oncogenic K-Ras in Distinct Cell Populations of the Adult Pancreas**

Activation of oncogenic K-Ras in the common pancreatic embryonic progenitor in the Pdx1Cre;LSL-Kras<sup>G12D</sup> model results in mPanIN formation and spontaneous progression to PDAC (Aguirre et al., 2003; Hingorani et al., 2003). Because human PDAC is an adult-onset malignancy, we sought to engineer PDAC mouse models in which oncogenic K-Ras (Kras<sup>G12D</sup>) is activated in the adult animal. We crossed the LSL-Kras<sup>G12D</sup> strain to different cell-specific CreER lines and compared the phenotypic effect of Kras<sup>G12D</sup> activation on these distinct cell populations in the adult pancreas (Figure 1A). The CreER lines used were: (1) the Pdx1CreER<sup>TM</sup> transgenic strain (Gu et al., 2002), which marks the earliest pancreatic progenitor cell during development as well as adult endocrine β cells, some ductal, acinar cells, and possibly adult progenitor/stem cells (Koizumi et al., 2003; Stoffers et al., 1999; Swift et al., 1998; Wu et al., 1997); (2) the RipCreER<sup>TM</sup> transgenic strain (Dor et al., 2004), which is selective for insulin<sup>+</sup> cells; and (3) the acinar procarboxypeptidase A1 CreER<sup>T2</sup> (proCPA1CreER<sup>T2</sup>) knockin strain (Zhou et al., 2007). The recombination specificity and efficiency of the different CreER strains was determined by crossing them to the LSL-LacZ reporter mouse and analyzing the X-gal-stained pancreata of their Tamoxifen (TM)-treated compound progeny (Figures 1B–1E and Table S1 available online). As shown in Figure 1B, in the Pdx1CreER<sup>TM</sup> transgenic strain, recombination was achieved in a mosaic fashion in the islets, acinar, and ductal cells. The proCPA1CreER<sup>T2</sup> strain exhibited recombination mainly in acinar cells and possibly CACs; however, recombination was evident also in a subset of ductal and islet cells. Recombination in this strain was variable, with half of the treated mice showing no evidence of recombination and the other half ranging from 0% to 49% in acinar cells (Figure 1C, Table S1, and data not shown). With TM administration, the RipCreER<sup>TM</sup> transgenic strain was very specific and efficient in causing recombination in both insulin-producing β cells located in the islets of Langerhans and in single insulin<sup>+</sup> cells found scattered throughout the pancreas parenchyma, as shown by X-gal and communofluorescent (ColF) staining (Figures 1D and 1E, respectively, and Table S1).

Expression of Kras<sup>G12D</sup> in different cell types of the adult pancreas was achieved by systemic TM administration in mice at 14–56 days of age. Because newly transformed cells would be expected to resemble their cell of origin both genetically and morphologically, we sought to define the earliest time point when transformed cells could be detected after TM administration. Therefore, mice were sacrificed at different times following the last dose, ranging from 4 to 240 days.

**Kras<sup>G12D</sup> Induces Transformation in Pdx1<sup>+</sup> Cells of the Adult Pancreas**

Pdx1CreER<sup>TM</sup>;LSL–Kras<sup>G12D</sup> mice were treated with TM at day 14, 21, 24, 27, or 56 after birth and sacrificed after 4–120 days. mPanINs and ductal metaplasia were observed in all age groups
(Figure 2 and Table S2). Low-grade mPanIN 1A lesions were the most prevalent and were observed to a comparable extent across the three main age groups, whereas mPanIN 1B were less frequent in the 56-day-old group. Although mPanIN lesions were predominantly grade 1, grade 2 lesions were also identified, mostly at 120 days after TM administration and only in the 14- and 27-day-old treated mice (Table S2 and Figure 2E). These results suggest that the Pdx1+ target cell for transformation is either less susceptible or less abundant in 56-day-old mice. mPanIN 3 was found only in one mouse that was analyzed 188 days after TM administration, and it was also the only mouse that developed PDAC. This result is consistent with observations in humans that it is unusual to find PanIN 3 in pancreata lacking invasive carcinoma (Mills, 2007).
Importantly, no lesions arose in vehicle-treated mice (n = 15) (Table S2). These results exclude the possibility of leaky Cre recombinase activity and TM-independent K-Ras activation during development and/or adulthood. In addition, we observed no evidence of chronic pancreatitis in treated mice. Since TM is dissolved in corn oil, which by itself causes a mild form of peritonitis (Figure S1E), we treated Pdx1CreERTM;LSL-KrasG12D mice with TM dissolved in carboxymethyl cellulose (CMC) by oral gavage. Dissolving TM in corn oil or CMC did not affect the specificity and efficiency of Cre-induced recombination as confirmed by X-gal staining in mice harboring a LacZ reporter (data not shown). As shown in Figure S1F, pancreata of mice treated with TM dissolved in CMC were free of peritonitis yet still developed grade 1 mPanINs (n = 5). This result precludes the possibility that the corn oil-induced peritonitis was essential for formation of mPanINs and ductal metaplasia in the Pdx1CreERTM;LSL-KrasG12D model.

Activation of KrasG12D in combination with mutations of Ink4A/Arf (Aguirre et al., 2003) or Trp53 (Jonkers et al., 2001) in Pdx1+ cells in the adult mouse resulted in formation of moderately to poorly differentiated PDAC in 10/13 mice tested (Figure 4 and Table S2). Many of the tumors contained undifferentiated areas with sarcomatoid or anaplastic features, and extension beyond the pancreas was common. We observed direct invasion into the stomach and small intestine as well as metastasis to mediastinal lymph nodes, diaphragm, and peritoneal adipose tissue (Figure 4 and data not shown). We also identified a full range of mPanIN lesions in these mice, including grade 3 (Table S2 and Figures 4A and 4B). Thus, Pdx1+ cells, or at least a subpopulation of Pdx1+ cells, represent an excellent candidate to be the cell of origin for PDAC in the mouse and in humans.

**Targeted Expression of KrasG12D in proCPA1+ Cells**

Targeting the expression of KrasG12D to pancreatic acinar cells and other cell types by treating proCPA1CreER72;LSL-KrasG12D mice with TM resulted in low grade mPanIN-1A formation in only 2/20 mice (Table S3 and Figure S2A). All mice were tested and confirmed for LSL-KrasG12D recombination by PCR analysis of...
DNA extracted from tissue sections (Table S3). These results suggest that proCPA1+ cells are not efficiently transformed by KrasG12D into mPanIN lesions in the adult mouse. It has been recently shown that KrasG12D activation in acinar cells of the adult mouse leads to efficient mPanIN formation in the absence of tissue injury in the Ela-CreERT2Tg/+; Mist1CreERT2; and ElaCreERT strains (De La O et al., 2008; Habbe et al., 2008). The phenotypic diversity observed between these models is discussed below (see also Table S7).

To determine whether loss of p53 or Ink4A/Arf could affect the neoplastic potential of proCPA1+ cells, we combined the LSL-KrasG12D allele with the Trp53lox/lox or Ink4A/Arflox/fox alleles on the background of the proCPA1CreERT2 strain. The pancreata derived from TM-treated proCPA1CreERT2;LSL-KrasG12D;Ink4A/Arflox/fox mice (n = 7) showed no overt cancer phenotype when examined at 50, 83, 113, 158, 188, and 309 days after TM administration. However, one proCPA1CreERT2;LSL-KrasG12D;Trp53fox/fox mouse (n = 5) developed a moderately to poorly differentiated PDAC 81 days after TM injection at 53 days of age. The tumor contained undifferentiated areas, was locally invasive, and metastasized to the liver and diaphragm (Table S3 and Figure S2B). As described earlier, all mice in this study were analyzed for DNA recombination by PCR (Table S3). We conclude that highly aggressive PDAC can be initiated in proCPA1+ cells in adult mice without chronic pancreatitis, albeit with low penetrance in this system.

In a previous study, Guerra et al. (2007) reported that selective expression of an endogenous K-RasG12V oncogene in adult cells of the acinar/centroacinar lineage using the K-Rasv12LSL12VCre; Ela-tTA/tetO-Cre mouse model resulted in no phenotype unless the mice were pretreated with caerulein (Guerra et al., 2007). Caerulein induces chronic pancreatitis that results in regeneration of normal tissue and formation of metaplastic lesions of the ductal phenotype in the exocrine compartment (Strobel et al., 2007). The authors concluded that chronic pancreatitis can be a cofactor in the induction of PDAC by the K-Rasv12LSL12VCre; Ela-tTA/tetO-Cre mouse model. None of the mice described above were targeted for transformation by Kras G12D, or Ink4A/Arf alleles on the background of the proCPA1CreERT2; LSL-KrasG12D mice described above were largely refractory to transformation by KrasG12D-induced transformation, we set out to test whether chronic pancreatitis could facilitate PDAC formation in these mice. We treated 34- to 60-day-old mice (n = 9) with caerulein for a total of 44–193 days. One month after caerulein treatment initiation, the mice were treated with TM to activate Cre. Interestingly, grade 1 mPanINs were detected in one of these mice. As described above, recombination of LSL-KrasG12D was confirmed by PCR analysis of DNA extracted from tissue sections of all TM-treated mice in this study (Table S3). These results strongly suggest that insulin+ cells are highly refractory to transformation by multiple oncogenic mutations.

We next set out to test the effect of chronic pancreatitis on PDAC formation in the RipCreERTM;LSL-KrasG12D mice. Two RipCreERTM;LSL-KrasG12D;LacZ mice were treated with caerulein for a total of 84 and 89 days. One month after caerulein treatment initiation, the mice were treated with TM to activate Cre. Interestingly, grade 1 mPanINs were detected in one of these mice (Figure 5B and Table S4). We confirmed LSL-KrasG12D recombination in the pancreas of the mouse that did not show mPanIN formation as well as in mPanINs derived from the second mouse by PCR analysis of DNA extracted from tissue sections and of DNA extracted from LCM lesions, respectively (Figure 5C and Table S4). Hence, the mPanINs observed arose from cells in which the transgenic rat insulin promoter was active. Importantly, all three RipCreERTM;LSL-KrasG12D;p53fox/fox;LacZ mice developed poorly and undifferentiated PDAC (n = 2 and n = 1, respectively) when treated first with caerulein and then with TM (Figure S3D and Table S4). As in the Pdx1CreERTM;LSL-KrasG12D;p53fox/fox mutant mice, these were highly aggressive tumors that exhibited a capacity for local invasion and distant metastasis.

One interpretation of these results (as well as those of Guerra et al. [2007]) is that caerulein treatment or inflammation caused a change in differentiated cells that made them susceptible to K-Ras activation. To investigate this possibility, we crossed the RipCreERTM;LSL-KrasG12D mice to the LacZ reporter strain and then with caerulein developed PDAC (Figure S3C and Table S4). To rule out false-positive results due to nonspecific Cre recombinase activity upon caerulein-induced injury, we treated proCPA1CreERT2;LSL-KrasG12D; LSL-LacZ mice (n = 3) with caerulein without prior TM administration for 8, 15, and 22 days. LacZ+ cells were not observed in these mice (data not shown). Thus, the proCPACreERT2 allele remains tightly controlled under chronic pancreatitis conditions. Immunofluorescence (IF) for CPA expression of normal tissue and formation of metaplastic lesions of the ductal phenotype in the exocrine compartment (Strobel et al., 2007).

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One interpretation of these results (as well as those of Guerra et al. [2007]) is that caerulein treatment or inflammation caused a change in differentiated cells that made them susceptible to K-Ras activation. To investigate this possibility, we crossed the RipCreERTM;LSL-KrasG12D mice to the LacZ reporter strain and
pulsed the triple compound progeny of these mice with TM 7–14 days prior to caerulein treatment initiation (n = 6) (Table S4). In this way, differentiated insulin+ cells and their progeny would be marked by the LacZ expression and their fate could be identified after caerulein treatment by X-gal staining (Figure 5D).

Strikingly, LacZ+ mPanINs were found in 4/6 pancreata of these mice (Figure 6A and Table S4). In the remaining two mice mPanINs were not observed, which might be explained by the small pancreatic area analyzed (less than 2 mm²; Table S6). In addition, two mice developed LacZ+ poorly to undifferentiated tumors with local invasion to adjacent tissues. These results suggest that KrasG12D-expressing insulin+ cells can be transformed and give rise to mPanINs and PDAC specifically in the context of chronic pancreatitis.

To explore the possibility of nonspecific Cre recombinase activity upon caerulein-induced injury, we treated RipCreERTM; LSL-KrasG12D; LSL-LacZ mice (n = 5) with caerulein without prior TM administration. Surprisingly, despite the absence of TM, LacZ+ cells were observed in the islets of Langerhans and in scattered cells throughout the parenchyma as early as 2 days after caerulein treatment initiation (data not shown). Thus, although the RipCreERTM strain is tightly controlled and does not normally show leaky expression of the Rat insulin promoter in insulin+/C0 cells or TM-independent activation of Cre recombinase (Table S1), Cre activity was more promiscuous under chronic pancreatitis conditions. This result raised the possibility that the mPanIN lesions observed in the experiment described above arose in non-insulin-expressing cells following inappropriate Cre activation during the caerulein treatment. To address this possibility directly, we treated RipCreERTM; LSL-LacZ mice (n = 5) with TM for 5–7 days, treated them with caerulein for 65 days, and subsequently analyzed them for coexpression of LacZ and insulin by CoIF (Table S5). In all cases, LacZ+ insulin+ cells were observed within the islet of Langerhans. In addition,
we scored a total of 83 LacZ+/insulin+ and 6 LacZ+/insulin-/C0 single cells scattered throughout the pancreata. We then calculated the frequency of mPanIN lesions per mm² in the TM/caerulein-treated RipCreERTM;LSL-KrasG12D;LSL-LacZ mice. Importantly, the frequency of mPanIN lesions in these mice was 6-fold higher than the frequency of LacZ+/insulin-/C0 cells in TM/caerulein-treated RipCreERTM;LSL-KrasG12D mice. Therefore, the frequency of mPanIN lesions was 6-fold higher than the frequency of LacZ+/insulin-/C0 cells in TM/caerulein-treated RipCreERTM;LSL-KrasG12D mice (p = 0.04 by Wilcoxon Rank Sum Test) (Tables S5 and S6). Thus, our results provide evidence that insulin+ cells can be made susceptible to KrasG12D and give rise to mPanIN and PDAC formation following tissue injury and inflammation. Interestingly, immunohistochemistry (IHC) revealed that most mPanIN cells in this model failed to express insulin (Figures 6B and 6C). Although the mechanism through which mature insulin-producing cells undergo this fate change is unclear, it is notable that these mPanINs harbored Pdx1+ cells and cells that positively stain for the Notch intracellular domain (Figure 6E and data not shown).

To further study transformation of insulin+ cells, we treated four RipCreERTM;LSL-KrasG12D;Trp53flox/flox;LSL-LacZ mice with TM and 14–40 days later with caerulein for additional 42–52 days. These mice developed high-grade mPanIN and poorly differentiated to undifferentiated carcinoma capable of local invasion and distant metastasis (Figures 7A and 7B and Table S4). The undifferentiated areas were morphologically identical to those observed in Pdx1CreERTM;LSL-KrasG12D;Trp53flox/flox mice and also resembled human undifferentiated pancreatic carcinoma (Hoorens et al., 1998). Tumors in these mice were negative for insulin as well as the neuroendocrine marker synaptophysin by IHC (Figures 7C and 7D). Furthermore, no preneoplastic lesions were observed in the islets of these mice (data not shown). In conclusion, our results provide evidence that in combination with pancreatic injury, KrasG12D-expressing insulin+ cells of the endocrine lineage, which are refractory to transformation under normal conditions, can also serve as a cell of origin of PDAC, a malignancy with an exocrine phenotype.

DISCUSSION

We have studied the susceptibility of adult pancreatic cells to transformation by KrasG12D. We show that Pdx1+ cells can efficiently give rise to mPanIN and ductal metaplasia and, in combination with p53 or Ink4A/Arf deletion, can develop into PDAC. In
contrast, in the absence of tissue injury and inflammation, insulin+ cells (which also express Pdx1) did not show any overt phenotype, even in the context of tumor suppressor gene mutations. Likewise, proCPA1+ cells were inefficiently transformed. Thus, we conclude that under noninflammatory conditions, a subpopulation of Pdx1+ cells of the adult pancreas have heightened sensitivity to the Kas(12D) oncogenic effects and represent a cell of origin of PDAC. Of note, Habbe et al. (2008) have reported that Pdx1+ cells are relatively resistant to Kas(12D)-induced transformation in 42-day-old mice. Our data suggest that younger mice are more susceptible to transformation of this cell type. It is also possible that the different observations are attributable to background differences between the mice used in both studies.

Four Cre lines have been used to target the expression of oncogenic K-Ras to adult acinar cells: K-RasLVSLGI12Vgeo;Kras(12D); Elas-1TA/tetO-Cre (Guerra et al., 2007); Ela-CreERT2G12D and Mist1CreERT2/+. (Habbe et al., 2008); ElaCreERT (De La O et al., 2008); and the proCPA1CreERT2 studied here. Different phenotypes, ranging from no lesion to high-grade mPanIN-3 formation, were observed in these strains (Table S7). The phenotypic diversity might be explained by several factors, including: (1) the K-RasLVSLGI12Vgeo strain lacks the 3’ UTR element, which contains several regulatory sequences, including binding sites for the let-7 miRNA family that regulates both N-Ras and K-Ras activation. (2) The models utilize distinct K-ras activating mutations (G12V versus G12D), which might have an effect on the oncogenic potency of K-Ras (Bardeesy and Sharpless, 2006; Collado et al., 2005; Seeburg et al., 1984). (3) Recombination efficiency differences between the different strains may exist, either because of differences between Cre-driving promoters, per se, mouse background, or different TM and doxycycline administration protocols. (4) The different mouse models may reflect a previously uncharacterized heterogeneity of acinar cells with respect to function and/or susceptibility to transformation. Nevertheless, the collective observations provide evidence that there are acinar cells in the adult pancreas sensitive to oncogenic K-Ras activation that can progress to at least mPanIN in the absence of overt tissue injury and that highly aggressive PDAC can be initiated from proCPA1+ cells, albeit with low penetrance.

When injury was induced before recombination, the proCPA1 CreERT2;LSL-Kras(12D) and RipCreERTM;LSL-Kras(12D) mice became sensitive to Kas(12D) activation. Importantly, when we activated Kas(12D) first and then induced chronic pancreatitis, RipCreERTM;LSL-Kras(12D) mice still developed mPanINs. We found that under caerulein-only treatment conditions Cre recombination activity was not detectable in the proCPA1CreERT2 strain, although, surprisingly, it could be detected in insulin+ and rare insulin+ cells in the RipCreERT2 strain. This result might reflect a differential sensitivity to caerulein of CreERT2 (Indra et al., 1999) in the proCPA1CreERT2 versus CreERTM (Danielian et al., 1998) in the RipCreERTM strain. Alternatively, an increased amount of Cre protein might be produced by the RipCreERTM transgene in comparison with the proCPA1CreERT2 knockin strain that might result in its greater sensitivity to caerulein. Regardless of the effects of caerulein, the vast majority of the mPanINs can be attributed to transformation of insulin+ cells, although it is possible that a few mPanINs are derived from insulin- cells as well. Furthermore, 2/6 of the RipCreERTM; LSL-Kras(12D);LSL-LacZ and all RipCreERTM;LSL-Kras(12D); p53foxfox;LSL-LacZ mice (n = 4) developed high-grade mPanIN and tumors that resembled the most aggressive PDAC that arose in Pdx1CreERTM and proCPA1CreERT2 compound mice. These tumors exhibited no morphologic or immunohistochemical evidence of neuroendocrine differentiation. In addition, the preinvasive changes in these mice were of an exocrine phenotype (i.e., mPanIN) rather than endocrine. Taken together, these results show that insulin+ cells of the endocrine lineage of the adult pancreas can transdifferentiate and give rise to highly aggressive exocrine neoplasia (PDAC). Of note, LacZ+ cells were not detected in ducal metaplasia in caerulein/TM-treated RipCreERTM;LSL-LacZ mice (data not shown), suggesting that the transdifferentiation of the insulin+ cells requires both injury and oncogenic K-Ras. Similar results were observed by Strobel et al. (2007) in RipCreERTM/Z/AP mice. Interestingly, oncogenic K-Ras activation is rarely observed in human endocrine tumors (Jonkers et al., 2007). This may in part be explained by reprogramming of endocrine cells to an exocrine fate upon injury and K-Ras activation.

The transdifferentiated cells in the RipCreERTM;LSL-Kras(12D) mice may be insulin+ cells that reside either in and/or outside the islets of Langerhans. The physiological role of the insulin+ cells that reside outside of the islets is largely unknown. However, the mPanINs in the caerulein/TM-treated mice were always observed outside the islets, suggesting that these cells may be more susceptible to Kas(12D) than those that reside in the islets. Interestingly, neoplastic ducts were found within the islets of TM-treated Pdx1CreERTM;LSL-Kras(12D) mice but not when Kas(12D) was activated during development with the nonconditional Pdx1Cre strain (Hingorani et al., 2003). This may reflect a developmental compensation mechanism or result from mouse background differences.

Caerulein might promote Kas(12D)-induced transformation by several mechanisms. It is possible that a rare population of proCPA1- and/or insulin-expressing progenitor cells that are normally susceptible to Kas(12D) oncogenic effect needs to be amplified by tissue injury for the appearance of the phenotype. Another possibility is that injury-induced regeneration or the secretion of inflammatory cytokines leads to the proliferation of stem cell/facultative stem cells that are most probably Pdx1+. Upon the initiation of differentiation of such cells, a subset might transiently express either proCPA1CreERT2 or RipCreERT2 and, thus, activate Kas(12D) upon TM administration. However, our results provide evidence that proCPA1- and insulin-expressing cells can form mPanINs, favoring the hypothesis that inflammation and tissue injury facilitate PDAC development by promoting reprogramming of differentiated cells rather than by stem cell mobilization. The mechanism through which such a reprogramming event might occur requires further investigation. However, our study and those of others suggest a role for the Notch signaling pathway in this process. Caerulein-derived acinar-ducat
metaplasia is associated with Notch pathway upregulation (Gomez et al., 2004; Jensen et al., 2005; Siveke et al., 2008). We also found Notch to be activated in mPanINs derived from caerulein/TM RipCreERTm1;LSL-KrasG12D mice. Reactivation of the Notch signaling pathway, which is important for cell-fate decisions and maintenance of undifferentiated stem/progenitor cells during embryogenesis (Artavanis-Tsakonas et al., 1999), may be a mechanism by which differentiated cells revert to a stem/progenitor fate or to convert to a cell type that is sensitive to KrasG12D activity. It has been recently shown that Notch and K-Ras reprogram exocrine cells to mPanINs (De La O et al., 2008). The precise mechanism of, and requirement for, interaction between chronic pancreatitis, Notch, and K-Ras activation for mPanIN formation from different pancreatic cells still needs to be determined.

These data add to the increasing body of evidence pointing to a remarkable plasticity of pancreatic adult differentiated cells (De La O et al., 2008; Habbe et al., 2008; Shen et al., 2000; Zhou et al., 2008) and question the existence of a pancreatic “stem cell” as well as its relevance for PDAC. Our findings suggest that KrasG12D regulates the differentiation status of pancreatic epithelial progenitor/differentiated cells to a certain extent and that in combination with nongenetic stress such as inflammation this is further accelerated. More specifically, a multipotent Pdx1+ insulin− cell may exist in the adult pancreas. This cell type may be the source of the endocrine cells within mPanINs and ductal metaplasia observed in the TM-treated Pdx1CreERTm1;LSL-KrasG12D mice and the endocrine cells arising from the ductal lining after pancreatic duct ligation (Xu et al., 2008). It is possible that this cell type is activated only in certain settings such as injury and neoplasia and may be particularly susceptible to KrasG12D-induced proliferation and differentiation. It might also be the source of the ductal structures located in the islets, if it resides both in the ductal lining as well as the islets or if it migrates to the islets under certain conditions. Alternatively, these structures may arise from a differentiated cell, such as the carbonic anhydrase II− cell that gives rise to endocrine and exocrine cells after birth and injury (Inada et al., 2008), upon KrasG12D activation.

Our study reveals the involvement of different cell types in PDAC initiation and provides a series of models to explore the biology and treatment of PDAC. With the advances in the ability to reprogram adult pancreatic cells in a controlled manner (Zhou et al., 2008), we expect that a better understanding of the reprogramming and transforming events attributed to KrasG12D activation in combination with chronic inflammation will lead to the development of new therapeutic modalities to prevent PanIN formation or to repair the diseased/damaged tissue.

**EXPERIMENTAL PROCEDURES**

**Mouse Strains**

All animal studies and procedures were approved by the Massachusetts Institute of Technology Institutional Animal Care and Use Committee. Research was conducted in compliance with the Animal Welfare Act Regulations and other Federal statutes relating to animals and experiments involving animals and adheres to the principles set forth in the Guide for Care and Use of Laboratory Animals, National Research Council, 1996. The LSL KrasG12D strain (Jackson et al., 2001) was crossed to the following strains: Pdx1CreERTm1 (Gu et al., 2002), proCPA1CreERTm1 (Zhou et al., 2007), RipCreERTm1 (Dor et al., 2004), Trp53flox (Jonkers et al., 2001), and Ink4A/Arflox (Aguirre et al., 2003). Primers used for genotyping by PCR are listed in Table S8 and details of reactions are available upon request. Cre strains were crossed to R26 LSL LacZ mice (Jackson Laboratories) to determine Cre expression patterns.

**Histopathology, Immunohistochemistry, and Immunofluorescence**

Histopathologic analysis of pancreata was carried out by two pathologists (G.C.C. and E.L.S.). mPanIN lesions and PDAC were graded according to consensus criteria for mouse models of pancreatic cancer (Hruban et al., 2006). We carried out IHC and IF analyses according to manufacturers’ recommendations, typically with a modified citric acid unmasking protocol followed by standard detection with 3,3′ diaminobenzidine with a kit from Vector Lab ratories. In some cases, secondary antibodies were conjugated to AlexaFluor 594 (Invitrogen) and nuclei were counterstained with 4,6 diamidino 2 phenyl indole. We used the following primary antibodies: insulin (Zymed), glucagon (Abcam), PYY (RDI), Pdx1 (gift of C.V. Wright), CPA1 (Chemicon), and β galactosidase (AbD Serotec). IF staining was performed as described previously (Jackson et al., 2001). Triple color microscopy and imaging were performed with a Nikon Eclipse E600 and a Spot cooled CCD camera and software. Images were processed with Adobe Photoshop.

**β-Galactosidase Histochemistry**

Pancreata were dissected and fixed in 4% PFA, incubated in 0.5 M Sucrose, and frozen in OCT. β-galactosidase activity was assayed as describe previ ously (Paratore et al., 2002).

**Tamoxifen Treatment**

Mice were treated with TM (Sigma) by intraperitoneal injections (i.p.) of 9 mg/ 40 g body weight in corn oil, one every other day for a total of three injections. For oral delivery, TM was dissolved in 0.5% CMC, 0.4% Tween 80 and 0.9% NaCl solution.

**Caerulein Treatment**

Mice were i.p. injected with 0.1 ml of a 50 µg/ml caerulein (Sigma) in saline solution five times per week.

**Pancreata Size Measurement**

Pancreata areas were determined with Bioquant Image Analysis software in manual measurement mode.

**Molecular Analysis**

For verification of Cre mediated recombination, DNA was prepared from wild type tails and LCM mPanINs of caerulein/TM treated proCPA1CreERTm1;LSL KrasG12D and RipCreERTm1;LSL KrasG12D mice. PCR was performed with primers flanking the Lox Stop Lox cassette (sequences available in Table S8) and on http://web.mit.edu/jacks lab/protocols table.html. Wild type K ras, 2Lox, and 1Lox K rasG12D alleles were detected, yielding 620 bp, 510 bp, and ~1100 bp products, respectively. LCM and DNA isolation were performed with the Veritas Microdissection System and the PicoPure DNA Extraction Kit from Molecular Devices, respectively.

**SUPPLEMENTAL DATA**

Supplemental Data include three figures and eight tables and can be found with this article online at http://www.cell.com/cancer cell/supplemental/S1535 6108(09)00338 9.

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K-Ras-Induced Tumorigenesis in Adult Pancreas

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