Menin determines K-RAS proliferative outputs in endocrine cells

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Introduction

K-RAS is a member of the RAS superfamily of membrane-bound GTPases that exist in either a GTP-bound active or GDP-bound inactive state (I). Signaling through G protein–coupled receptors and receptor tyrosine kinases can stimulate the formation of RAS-GTP, which then binds one of several downstream effectors to activate a signaling cascade. The best understood function for K-RAS is that of a mitogen and protooncogene working through the RAF/MAPK pathway to drive the expression of proproliferative genes in the nucleus. Activating mutants such as K-RASG12D, which block GTP hydrolysis and lock K-RAS in the activated state, act as dominant drivers of cellular proliferation and occur frequently in many types of cancer (Figure 1A and ref. 2).

However, K-RAS coordinates multiple, diverse, and sometimes opposing signaling networks (3), suggesting that K-RAS may have context-dependent functions. For example, expression of K-RASG12D stimulates transformation and tumor formation in pancreatic ducts and acini — but not in the pancreatic endocrine cells found in the islets of Langerhans, even after reducing the level of tumor suppressors Cdkn2a and Trp53 (4). In addition, while KRAS is the most frequently mutated gene in pancreatic ductal adenocarcinoma (5), human pancreatic endocrine tumors never carry activating KRAS mutations (6).

In contrast, pancreatic endocrine tumors more commonly inactivate tumor suppressors such as menin (6) or RASSF1A (7–9). RASSF1A, a product of the RASSF1 gene, is an antiproliferative effector of activated K-RAS (10). Menin is the product of the MEN1 gene. Heterozygous null mutations in MEN1 cause multiple endocrine neoplasia type 1 (MEN1), an autosomal dominant cancer syndrome characterized by tumors of the endocrine cells of the pancreatic islets and the parathyroid and pituitary glands (11).

We hypothesized that the signaling networks downstream of K-RAS in pancreatic endocrine cells differed from those in K-RASG12D-sensitive cells such as the pancreatic acinar and duct cells. In testing this hypothesis, we discovered that mice heterozygous for a null mutation in Kras had increased numbers of pancreatic endocrine cells. Focusing on the insulin-producing β cells in the pancreatic islets, we found that their expansion came from 2 sources: increased production from neurogenin 3–expressing endocrine progenitors during embryogenesis and accelerated β cell proliferation during the perinatal period. In contrast, increased K-RAS signaling from constitutively active K-RASG12D suppressed both sources of new β cells, while still activating both the MAPK pathway and the RASSF1A pathway.

Kras heterozygosity also increased the proliferation of endocrine cells in 2 other tissues sensitive to MEN1 mutation: the parathyroid and pituitary glands. In addition, we found that the dominance of the antiproliferative K-RAS effect in β cells depended on the expression of menin. Our data suggest a model in which K-RAS activates both the proproliferative MAPK pathway and the antiproliferative RASSF1A pathway. In the tissues endocrine cell proliferation fluctuates dramatically in response to signals that communicate hormone demand. The genetic alterations that override these controls in endocrine tumors often are not associated with oncogenes common to other tumor types, suggesting that unique pathways govern endocrine proliferation. Within the pancreas, for example, activating mutations of the prototypical oncogene KRAS drive proliferation in all pancreatic ductal adenocarcinomas but are never found in pancreatic endocrine tumors. Therefore, we asked how cellular context impacts K-RAS signaling. We found that K-RAS paradoxically suppressed, rather than promoted, growth in pancreatic endocrine cells. Inhibition of proliferation by K-RAS depended on antiproliferative RAS effector RASSF1A and blockade of the RAS-activated proproliferative RAF/MAPK pathway by tumor suppressor menin. Consistent with this model, a glucagon-like peptide 1 (GLP1) agonist, which stimulates ERK1/2 phosphorylation, did not affect endocrine cell proliferation by itself, but synergistically enhanced proliferation when combined with a menin inhibitor. In contrast, inhibition of MAPK signaling created a synthetic lethal interaction in the setting of menin loss. These insights suggest potential strategies both for regenerating pancreatic β cells for people with diabetes and for targeting menin-sensitive endocrine tumors.

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het; Figure 1B). We discovered that pancreata from E13.5 Kras\textsuperscript{het} embryos contained more endocrine cells, as determined by combined glucagon and insulin staining, than did WT pancreata, while overall pancreatic size remained unchanged (Figure 1, C–E, and Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI69004DS1). By P5, Kras\textsuperscript{het} animals had a relative expansion of both the glucagon-expressing α cell and insulin-expressing β cell populations without any increase in the exocrine pancreas (Figure 1, F–H, and Supplemental Figure 2). Further, Kras\textsuperscript{het} adults had improved glucose tolerance compared with that of WT controls (Supplemental Figure 3), consistent with an increase in functional β cell capacity.

Next, we sought to determine the source of additional endocrine cells in Kras\textsuperscript{het} animals. The fetal pancreas generates endocrine cells by neogenesis, followed by rapid expansion of those cells by proliferation during perinatal and postnatal growth (Figure 2A and ref. 13). Using expression of the transcription factor NEUROG3 to identify the transient precursor cells generated susceptible to MEN1 gene mutation, menin normally prevents the MAPK effector pathway from driving proliferation, while leaving inhibitory effector pathways such as RASSF1A intact. In this model, loss of menin causes proliferation in susceptible cells due to removal of the blockage of MAPK-driven proliferation downstream of K-RAS, while loss of K-RAS signaling increases proliferation by decreasing unopposed RASSF1A activity. Our data explain the absence of activating KRAS mutations and the high frequency of MEN1 and RASSF1A inactivation in pancreatic endocrine tumors. Our study also suggests potential antiproliferative strategies for treating these tumors and proproliferative therapies for diseases that result from a deficiency of endocrine cell types, such as β cells, in both type 1 and type 2 diabetes.

Results
To test K-RAS function in pancreatic endocrine cell growth, we used mice carrying the Kras\textsuperscript{G12D} allele, which is a null allele in the absence of Cre recombinase (12), in place of 1 WT allele (Kras-
function allele that expresses constitutively active K-RASG12D (Figure 3, A–C). In expression field and in glucagon- and insulin-expressing cells (Figure 2, C–E, and Supplemental Figures 5 and 6). Thus, loss of 1 Kras allele led to increases in both neogenesis and proliferation of endocrine cells in the pancreas.

To test the effect of increased K-RAS signaling, we used Cre recombinase to convert KrasG12D from a null allele into a gain-of-function allele that expresses constitutively active K-RASG12D (Figure 1B) in pancreatic progenitor cells (Pdx1-Cre) or in β cells (Ins2-Cre) (Supplemental Figure 7). Pdx1-Cre KrasG12D embryos had no change in pancreatic size, but a marked reduction in the NEUROG3 recombine to convert KrasG12D from a null allele into a gain-of-function allele that expresses constitutively active K-RASG12D (Figure 1B) in pancreatic progenitor cells (Pdx1-Cre) or in β cells (Ins2-Cre) (Supplemental Figure 7). Pdx1-Cre KrasG12D embryos had no change in pancreatic size, but a marked reduction in the NEUROG3 expression field and in glucagon- and insulin-expressing cells (Figure 3, A–C). In Ins2-Cre KrasG12D pups at P5, we observed that β cells proliferated at markedly lower rates than was detected in Ins2-Cre control animals (Figure 3D and Supplemental Figure 8). These data demonstrate that K-RAS signaling directly antagonizes endocrine neogenesis in the embryo and proliferation in neonates.

Through distinct effectors, K-RAS can activate downstream pathways with opposing effects on cellular proliferation (3, 15). K-RAS effector B-RAF, which acquires activating mutations in many tumors, drives proliferation through activation of the MAPK pathway (2), while K-RAS effector RASSF1A blocks cell-cycle progression and is epigenetically silenced in many tumors (10). Pancreatic β cells express mRNA encoding multiple members of the RAS, RAF, and RASSF protein families (16 and Supplemental Figure 9). B-RAF and RASSF1 were broadly detected by immunostaining in the pancreas at P5, with relatively high expression in islets (Figure 3, E and F). To test the ability of K-RAS to activate these pathways in islets, we assayed for the phosphorylation of RASSF1 (Ser131) (p-RASSF1), which is critical for its tumor-suppressive activity (17), and ERK1/2 (Thr202/Tyr204) (p-ERK1/2), which lies downstream of B-RAF in the MAPK pathway (18), and detected an increase in the phosphorylated forms of both proteins broadly in β cells of Ins2-Cre KrasG12D mice as compared with the β cells of Ins2-Cre controls (Figure 3, G–J, and Supplemental Figure 10). We observed no changes in either p-ERK1/2 or p-RASSF1 in whole pancreas extracts from Kras heterozygous inactivating germline mutations of the MEN1 tumor-suppressor gene that encodes menin cause multiple endocrine neoplasia type 1 (MEN1), an autosomal dominant cancer syndrome characterized by tumors in a subset of endocrine cell types, especially pancreatic islet, parathyroid, and pituitary (11). Similar to Kras heterozygous inactivating germline mutations of the MEN1 tumor-suppressor gene that encodes menin cause multiple endocrine neoplasia type 1 (MEN1), an autosomal dominant cancer syndrome characterized by tumors in a subset of endocrine cell types, especially pancreatic islet, parathyroid, and pituitary (11). Similar to Kras heterozygous inactivating germline mutations of the MEN1 tumor-suppressor gene that encodes menin cause multiple endocrine neoplasia type 1 (MEN1), an autosomal dominant cancer syndrome characterized by tumors in a subset of endocrine cell types, especially pancreatic islet, parathyroid, and pituitary (11).
selectively increases islet cell growth and β cell proliferation as well as tumor formation in the MEN1 tissues, without affecting the exocrine pancreas or other cell types (22–24). We therefore examined whether other MEN1 tissues were affected in Kras-het mutants. We found that parathyroid glands were enlarged in Kras-het neonates relative to those in WT littermates (Figure 4, A and B, and Supplemental Figure 13), with increased proliferation of parathyroid chief cells, but not follicular cells, in the adjacent thyroid (Figure 4, C, D, and G, and Supplemental Figure 14). In the pituitary, we found that size and shape were unchanged (data not shown), but prolactin-expressing lactotrophs showed increased proliferation in Kras-het neonates relative to those in their WT littermates.
(Figure 4, E–G, and Supplemental Figure 14). Thus, reduced *Kras* gene dosage increased the growth of endocrine cells in the pancreas, parathyroid, and pituitary. To test the interaction between K-RAS and menin, we used a conditional *Men1*-null allele (25). We found that reducing *Men1* gene dosage increased proliferation and switched K-RAS to an activator of proliferation in β cells. Reducing *Men1* K-RAS and menin, we used a conditional –null allele (25).

It has been proposed that menin blocks growth by preventing p-ERK1/2 from driving cell-cycle progression (26). Consistent with this hypothesis, transgenic expression of menin in MIN6 insulinoma cells (Supplemental Figure 15) rendered the proliferation of these cells insensitive to the highly specific MEK1/2 inhibitor PD0325901 (20), which blocked ERK1/2 phosphorylation (Supplemental Figure 15). Conversely, we found that PD0325901 rescued the hyperproliferation of β cells with reduced *Men1* gene dosage (Figure 5, A–G). ERK1/2 phosphorylation was unaffected in *Ins2-Cre Men1*β−/− islets (Supplemental Figure 11). Remarkably, complete loss of *Men1* resulted in a synthetic lethal interaction (27) with PD0325901 (Figure 5F) or GSK1120212, another highly specific MEK1/2 inhibitor (data not shown) (28). Also, while FTI-GGTI elevates proliferation of adult WT and *Ins2-Cre* β cells, this effect was reversed in *Men1* mutant β cells (Figure 5G). Together, these data demonstrate that menin functions to block MAPK-driven proliferation in endocrine β cells.

Our data suggest a model in which menin determines K-RAS effects on endocrine cell proliferation by blocking proliferation driven by the MAPK pathway, while leaving K-RAS inhibitory pathways, such as RASSF1A, intact (Figure 5I). In this model, reduced menin gene dosage increases endocrine cell growth due to a reduction in blockage of MAPK-driven proliferation downstream of K-RAS, while reducing K-RAS gene dosage in endocrine cells with normal menin activity increases growth by decreasing the activity of the antiproliferative RAS effector RASSF1A. This model predicts that reduced menin activity might permit activators of the MAPK pathway to increase β cell proliferation. As predicted, the glucagon-like peptide 1 (GLP1) agonist exendin-4, which stimulates ERK1/2 phosphorylation in β cells (29), did not affect human β cell proliferation when added alone, but synergistically enhanced proliferation when combined with MI-2, an inhibitor of the interaction between menin and the histone methyltransferase MLL (ref. 30 and Figure 5, H and I).

**Discussion**

Despite its dominant role in stimulating growth in the exocrine pancreas (5), our data reveal that K-RAS acts to suppress growth in the pancreatic endocrine cells of the islets of Langerhans. This dichotomy further demonstrates how the same signaling pathway can generate opposing outputs in different cells (1–3, 31, 32) and provides an explanation for the divergent role of *KRAS* in the transformation of these 2 major pancreatic cell types (4–7, 33). The antiproliferative activity of K-RAS required the RAS effector RASSF1A, suggesting that RASSF1A silencing in pancreatic endocrine tumors (7, 9, 34–36) represents a release from the physiological growth inhibition imposed by this pathway. RASSF1A can inhibit cell proliferation by controlling the transcription of cell-cycle genes, the stability of cell-cycle proteins, or the stability of
microtubules (37–43). RASSF1A may also cooperate with other RASSF members to effect its action downstream of K-RAS (44).

In addition to the pancreas, endocrine tumors that arise from the parathyroids or pituitary are devoid of activating KRAS mutations (45–48) and frequently experience RASSF1A gene silencing (49, 50). Together with the endocrine pancreas, these tissues comprise the most common sites of tumorigenesis in people who inherit a single inactivating mutation in the menin-expressing MEN1 gene (11, 51). Given the wide expression of K-RAS, RASSF1A, and menin, what mediates the cell specificity of tumor formation in people carrying MEN1 gene mutations? We hypothesize that discrete cell types use distinct regulatory networks to specify and restrict the biological actions of K-RAS. In MEN1-sensitive endocrine cells, K-RAS activates opposing growth pathways, but antiproliferative pathways dominate due to the activity of menin, which prevents the MAPK pathway from driving growth, while leaving the inhibitory RASSF1A pathway intact. In this model, loss of menin increases proliferation due to removal of the blockage of MAPK-driven proliferation downstream of K-RAS, while loss of K-RAS signaling increases proliferation by decreasing unopposed RASSF1A activity (Figure 5J). The importance of the RAF/MEK/ERK pathway in driving the inappropriate growth and survival of endocrine cells with reduced menin activity offers a rationale for using inhibitors of this pathway for antiproliferative and cytotoxic therapy for tumors resulting from germline or somatic MEN1 mutations.

How menin blocks MAPK-driven proliferation in endocrine cells remains unclear. Menin localizes to the nucleus and chromatin-modifying complexes including MLL and can regulate gene expression (52, 53), but it has been shown to block cell proliferation by multiple mechanisms, including modulating both...
transcriptional regulators and cell signaling pathway components through direct protein-protein interactions (52–54). Identifying the unique targets of the MAPK pathway in menin WT and mutant endocrine cells may provide insight into the unique sensitivity of endocrine cells to loss of menin.

In addition to providing new insights into endocrine growth and tumorigenesis, the model in Figure 5J also suggests potential therapeutic targets for diabetes. Diabetes treatment has the opposite objective of antineoplastic therapies, that of enhancing proliferation to regenerate β cell mass, which is a major and currently unmet goal in the treatment of both type 1 and type 2 diabetes. Therefore, appropriate combinations of inhibition and activation of specific K-RAS pathway components may represent a rational strategy for inducing β cell proliferation and restoration for people with diabetes.

In conclusion, our study provides fresh insight into the unique pathways that regulate endocrine cell growth and presents what we believe to be a novel mechanism for the cell type-specific control of K-RAS function.

Methods

Animals. Mice were housed on a 12-hour light/12-hour dark cycle in controlled-climate rooms (21.5–22.5°C). The Kras^{G12D} (12), Men1^{−/−} (25), Rasffla^{−/−} (21), R26^Ins2-Cre^{lox/lox} (55), Pdx1-Cre (56), and Ins2-Cre (57) transgenic mouse lines were maintained and genotyped as previously described. Noon of the day on which a vaginal plug was detected was considered E0.5.

Glucose tolerance tests. A glucose tolerance test (GTT) was performed on 10-week-old male mice that had been placed on a high-fat diet (Research Diets) for 6 weeks beginning at 4 weeks of age. All mice were fasted overnight and placed in fresh cages to ensure that no food was available on the cage floor. The following morning, each mouse was intraperitoneally injected with glucose dissolved in water (2 g/kg body weight). Glucose measurements were taken from tail blood using a Freedom Lite glucose meter (Abbott Laboratories).

Pancreatic histology. Mice were euthanized by carbon dioxide asphyxiation, and embryos were harvested by dissection using a dissecting stereoscope. Tissues were then fixed in 4% PFA at 4°C overnight and placed in 100% methanol. Next, tissues were treated with 5% hydrogen peroxide for 5 hours and then washed with methanol 3 times and rehydrated into PBS containing 0.1% Tween-20 (Sigma-Aldrich). Tissues were then incubated with primary antibodies against neurogenin 3 (German laboratory; ref. 14), insulin (Dako), glucagon (Dako), and E-cadherin (Invitrogen), which had been diluted in PBTX with 5% donkey serum overnight at 4°C, followed by 3 washes in cold PBTX. For neurogenin 3, insulin, and glucagon staining, tissues were then incubated with fluorescent secondary antibodies (Invitrogen) that had been diluted in PBTX with 5% donkey serum overnight at 4°C, followed by 3 washes in cold PBTX. For E-cadherin, tissues were instead incubated with a biotinylated rabbit anti-rat antibody (Vector Laboratories) overnight at 4°C, followed by 3 washes in cold PBTX. Tissues were then incubated with a streptavidin-conjugated horseradish peroxidase from the VECTASTAIN Elite ABC kit (Vector Laboratories) for 2 hours at room temperature, followed by 3 washes in cold PBTX. Tissues were then incubated with fluorescent tyramide (PerkinElmer) for 30 minutes at room temperature, followed by 3 washes in cold PBTX. For imaging, stained tissues were placed in 100% methanol and cleared using a 1:2 mixture of benzyl alcohol/benzyl benzoate and placed into a custom-made glass chamber. A z series through the intact embryonic pancreas was collected on a Zeiss LSM 510 confocal microscope using a ×10 objective. The area of fluorescent staining throughout the sample was quantified using ImageJ software. Measurements were normalized using the average value from control samples.

Gene expression studies. Gene expression from adult β cells (Supplemental Figure 9) was derived from reanalysis of the data from ref. 16.

Islet culture and proliferation assessment. Mouse and human pancreatic islets were isolated by collagenase digestion and were then hand-picked as previously described (58). Mouse islets were obtained from P5 neonates and 12-week-old male mice. Human islets were obtained by the UCSF Diabetes Research Center Islet Core Laboratory from deceased donors (donor 1: 22-year-old male; donor 2: 23-year-old female; donor 3: 28-year-old male; donor 4: 42-year-old male; donor 5: 48-year-old female; donor 6: 55-year-old male). For proliferation studies, islets were cultured for 5 days in RPMI-1640 media and treated with FTI-GGTI inhibitors (10 μM and 20 μM; respectively) (Sigma-Aldrich); 10 μM PD0325901 (Sigma-Aldrich); 10 μM GSK1120212 (Selleck Chemicals);
90 nM exendin-4 (Sigma-Aldrich); 1 μM MI-2 (XcessBio); or DMSO (Sigma-Aldrich). Medium with these additives was freshly made and replaced daily. On day 5 of treatment, islets were incubated with 10 μM Edu (Invitrogen) (mouse islets for 16 hours; human islets for 48 hours) and then fixed in 4% PFA for 2 hours. Fixed islets were washed 3 times with PBS and then incubated with primary antibodies raised against NKX6.1 (Sigma-Aldrich) in PBTX with 5% donkey serum overnight at 4°C, followed by 3 washes in cold PBTX. Islets were then incubated with fluorescent secondary antibodies (Invitrogen) that had been diluted in PBTX with 5% donkey serum at 1:300 overnight, followed by 3 washes in cold PBS. Islets were then stained for DNA using either Topro3, DAPI, or Hoechst 33342, at a dilution of 1:10,000 in all cases. After staining, replicating cells were labeled with the Click-iT Edu Alexa Fluor Imaging Kit (Invitrogen). Islets were placed in VECTASHIELD and imaged using a Leica SP5 confocal laser scanning microscope. ImageJ software was used to count NKX6.1/Edu-double-positive nuclei. The relative percentage of proliferating β cells was calculated by dividing the number of nuclei containing Edu and NKX6.1 by the total number of NKX6.1+ nuclei and then normalized using control values. For immunoblots, islets were cultured overnight (PD0325901) or overnight and then for 30 minutes with fresh inhibitors (FTI-GGTI), before harvesting.

MIN6 insulinoma cell culture. MIN6 insulinoma cells (59) were maintained in DMEM supplemented with 15% FBS, 100 units/ml penicillin, 100 μg/ml streptomycin, and 71.5 μM β-mercaptoethanol. A MIN6 subclone with low menin expression was used for this study. For proliferation studies, MIN6 insulinoma cells were plated on Permanox 8-well chamber slides (Electron Microscopy Sciences) on day 0, transfected on day 1, treated on days 2 and 3, and then fixed in 4% PFA for 1 hour on day 4. Transient transfection of MIN6 insulinoma cells was performed using Lipofectamine 2000 reagent (Invitrogen).

Immunoblotting. Immunoblotting was performed using standard procedures. Briefly, cells or islets were lysed in buffer containing 50 mM HEPES (pH 7.5), 150 mM NaCl, 1% Triton X-100, and Halt Protease Inhibitor Cocktail (Pierce Biotechnology). Lysates containing 40 μg of protein, as determined by Bradford Reagent (Bio-Rad), were resolved by SDS-PAGE and transferred to nitrocellulose or PVDF membranes (Bio-Rad). Membranes were blocked with 5% nonfat milk and/or 5% fraction-V BSA (Sigma-Aldrich) in Tris-buffered saline containing 0.1% Tween-20 (TBST). Membranes were incubated with primary antibodies overnight at 4°C. The following primary antibodies were used: p-ERK1/2 (Sigma-Aldrich and Cell Signaling Technology) and ERK1/2 (Abcam and as described in ref. 60). RASSF1 (Sigma-Aldrich) and p-RASSF1A (Ser131) (17) antibodies were preabsorbed using pancreatic tissue powder derived from neonatal or adult Rassf1a+/− mice. Membranes were washed with TBST and incubated with the following secondary antibodies: donkey anti-rabbit IRDye 680RD (LI-COR Biosciences); donkey anti-mouse IRDye 800CW (LI-COR Biosciences); donkey anti-rabbit HRP (Cell Signaling Technology); and donkey anti-mouse HRP (Cell Signaling Technology). For multiple detection, membranes were stripped using stripping buffer (Thermo Fisher Scientific). Membranes were imaged using either the LI-COR Odyssey Infrared imaging system or x-ray film and a flat-bed scanner.

Generation of menin iresGFP plasmid. The plasmid pMenin ires-GFP was generated by inserting coding in tandem sequences for human menin (Addgene; ref. 61), an internal ribosomal entry site (ires), and GFP into the pCAGG mammalian expression vector (62).

Statistics. All data points were compared using a 2-tailed Student’s t test, and a P value of less than 0.05 was considered statistically significant.

Study approval. All procedures were approved by the IACUC of the UCSF and were conducted in accordance with regulations of the UCSF, which gave ethical approval for these studies.

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