

# Adult Inactivation of the Recessive Polycystic Kidney Disease Gene Causes Polycystic Liver Disease

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## Abstract

**Background** A major difference between autosomal recessive polycystic kidney disease (ARPKD) and autosomal dominant polycystic kidney disease (ADPKD) lies in the pattern of inheritance, and the resultant timing and focality of cyst formation. In both diseases, cysts form in the kidney and liver as a consequence of the cellular recessive genotype of the respective disease gene, but this occurs by germline inheritance in ARPKD and somatic second hit mutations to the one normal allele in ADPKD. The fibrocystic liver phenotype in ARPKD is attributed to abnormal ductal plate formation because of the absence of *PKHD1* expression during embryogenesis and organ development. The finding of polycystic liver disease in a subset of adult *PKHD1* heterozygous carriers raises the question of whether somatic second hit mutations in *PKHD1* in adults may also result in bile duct-derived cyst formation.

**Methods** We used an adult-inducible *Pkhd1* mouse model to examine whether *Pkhd1* has a functional role in maintaining bile duct homeostasis after normal liver development.

**Results** Inactivation of *Pkhd1* beginning at 4 weeks of age resulted in a polycystic liver phenotype with minimal fibrosis at 17 weeks. Increased biliary epithelium, which lines these liver cysts, was most pronounced in female mice. We assessed genetic interaction of this phenotype with either reduced or increased copies of *Pkd1*, and found no significant effects on the *Pkhd1* phenotype in the liver or kidney from altered *Pkd1* expression.

**Conclusions** Somatic adult inactivation of *Pkhd1* results in a polycystic liver phenotype. *Pkhd1* is a required gene in adulthood for biliary structural homeostasis independent of *Pkd1*. This suggests that *PKHD1* heterozygous carrier patients can develop liver cysts after somatic mutations in their normal copy of *PKHD1*.

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## Introduction

*PKHD1* is the disease gene for autosomal recessive polycystic kidney disease (ARPKD). ARPKD is a pediatric diagnosis with incidence 1:20,000 live births, characterized by diffusely cystic kidneys progressing to renal failure and congenital hepatic fibrosis (CHF) (1). CHF is a fully penetrant component of the diagnosis characterized by liver fibrosis and portal hypertension, whose consequences require liver transplant in 1%–7% of survivors of early childhood (2,3). CHF pathogenesis is attributed to malformation of the ductal plate during liver development, and ARPKD-related CHF can be distinguished from other forms of liver fibrosis by the occurrence of biliary-derived cysts. Patients with missense mutations tend to have a milder kidney phenotype compared with those with truncating mutations and in some cases are not diagnosed until liver abnormalities are identified in adulthood (4,5). Mouse models for ARPKD recapitulate the human liver phenotype with fibrosis and cysts (6–10). Pancreas phenotypes are variable. ARPKD mice lack a kidney phenotype at least until late in life when mild tubular dilations have been

reported (7,8). The function of fibrocystin is not well understood, and this hinders attempts to identify ARPKD treatments.

The parents of ARPKD children are healthy, and there is no kidney or liver phenotype in heterozygous rodent models (6,9). Monoallelic loss of *PKHD1* has been thought to be inconsequential. Nonetheless, 10% of asymptomatic carrier adults were found to have multiple cysts in their livers or kidneys, and a gene discovery analysis of individuals ascertained on the basis of isolated polycystic liver disease (PCLD) identified *PKHD1* heterozygous carrier genotype as the causative gene mutation (11,12). Genetic studies and modeling of several other gene causes of PCLD have shown that these occur by somatic second hit mutations akin to the mechanisms underlying autosomal dominant polycystic kidney disease (ADPKD) (12–14). These findings allow for two possible genetic models for cyst pathogenesis in *PKHD1* heterozygous mutation carriers. Bile duct cysts may form either because of haploinsufficiency or because of a focal cellular recessive genotype resulting from somatic loss of the single

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normal allele. This second possibility can only be true if the *PKHD1* gene has an ongoing homeostatic role after liver development. We used an adult-inducible mouse knockout model of *Pkhd1* to determine whether postdevelopmental inactivation of *Pkhd1* results in a PCLD phenotype. We found that somatic inactivation of *Pkhd1* is sufficient to cause bile duct cysts in adult livers.

## Materials and Methods

### Mouse Lines

All experiments were conducted in accordance with Yale University Institutional Animal Care and Use Committee guidelines and procedures. The *Pkhd1<sup>flox3-4</sup>* (*Pkhd1<sup>tm1Ggg</sup>* (7)) allele was a generous gift from Dr. Gregory G. Germino (National Institute of Diabetes and Digestive and Kidney Diseases/National Institutes of Health). The *Pkd1<sup>flox</sup>*, *Pkd1<sup>F/H-BAC</sup>*, tamoxifen-inducible *UBC-Cre<sup>ER</sup>*, and doxycycline-inducible *Pax8<sup>rtTA</sup>;TetO-Cre* models have been previously described (13,15,16). The *Pkd1<sup>F/H-BAC</sup>* contains three copies of a bacterial artificial chromosome modified to express dual-epitope-tagged *Pkd1*. All mice are predominantly C57BL6. Littermates were used as controls.

### Pharmacological Induction and Sacrifice

Gene inactivation using both the *UBC-Cre<sup>ER</sup>* and *Pax8<sup>rtTA</sup>;TetO-Cre* models began at postnatal day 28. Tamoxifen 20 mg/ml in sunflower seed oil was administered as a 0.1-ml intraperitoneal injection daily for 5 days. Doxycycline (2 mg/ml in a 5% sucrose solution) was provided in the drinking water for 2 weeks. *UBC-Cre<sup>ER</sup>* mice were sacrificed at age 17 weeks, and *Pax8<sup>rtTA</sup>;TetO-Cre* mice were sacrificed at age 26 weeks (6 months). Organs were perfused with 1× PBS then 4% paraformaldehyde, and stored in 4% paraformaldehyde before sectioning for histology.

### Histology and Immunohistochemistry

Paraffin-embedded tissue sections were stained with hematoxylin and eosin (H&E), Masson Trichrome, Sirius Red, or left unstained. To perform anti-cytokeratin 19 (CK19) immunohistochemistry, unstained slides were treated with xylene and sequential ethanol dilutions to remove paraffin and rehydrate the tissue. Endogenous peroxidase was blocked with 10% hydrogen peroxide in methanol. After water wash, antigen retrieval was achieved with steaming in 1 mM EDTA pH 8 followed by 0.05% Tween 20 in PBS. Samples were then blocked with 1% BSA for 30 minutes before primary antibody application overnight at 4°C. Washes were completed with 0.05% Tween 20. Primary antibody: anti-CK19;TROMA-III (1:100; Developmental Studies Hybridoma Bank, Iowa City, IA); secondary antibody: anti-rat horseradish peroxidase (1:500; Jackson ImmunoResearch, West Grove, PA). Horseradish peroxidase substrate was provided by the 3'-diaminobenzidine peroxidase substrate kit (Vector Laboratories, Burlingame, CA).

### Cystic Parameters

Slides stained with H&E or immunohistochemistry were scanned on a Nikon Eclipse TE2000 microscope using MetaMorph software (Molecular Devices, San Jose, CA) to black and white TIFF files, respectively. For both hepatic cystic

index (HCI) and the percentage of the two-dimensional liver area that stained positive for CK19 (CK19+%) calculations, a denominator was set with the manually outlined perimeter of the liver excluding large internal venous structures. Autothresholding within this space was performed by MetaMorph to determine white (cystic) from dark (noncystic) for HCI. HCI is defined as the percentage of cystic/total parenchymal area. CK19+% is defined as the percentage of CK19+ staining over the total parenchymal area.

### Plots and Statistics

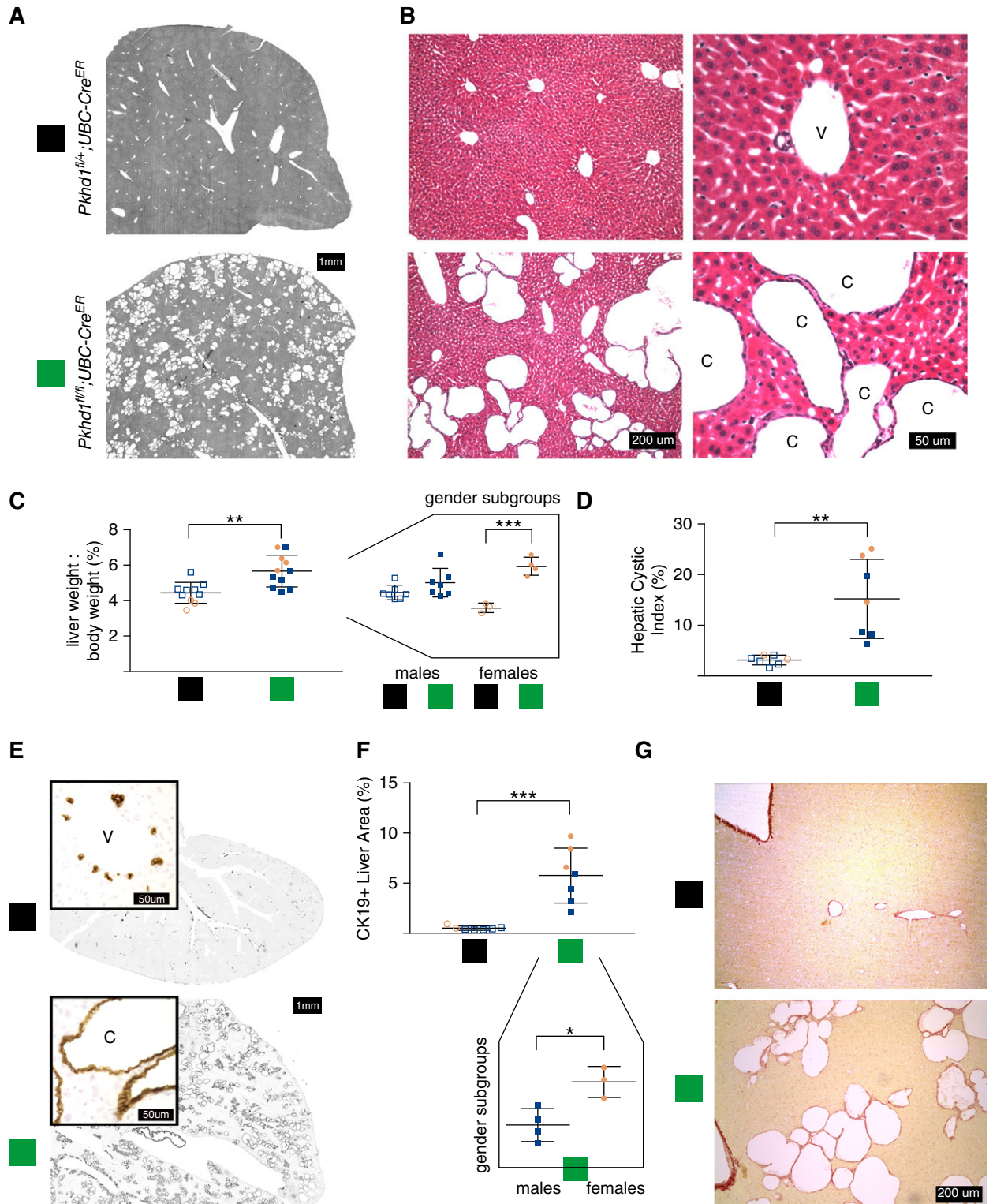
Two-way comparisons of sample means were calculated using the unpaired two-tailed *t* test, a parametric test using Prism version 7 (GraphPad Software, Inc., San Diego, CA). Fisher's exact test was used to make pairwise comparisons of proportions. Sample size calculation was performed using online calculators (<http://powerandsamplesize.com>) for the *Pkd1* dosage experiments to suggest a target of 15 mice of each genotype to have 80% power to detect a difference with 5% risk for type 1 error between a HCI of 12 and 20, with estimated SD of 7.8 from the initial experiment.

## Results

### Postdevelopmental Inactivation of *Pkhd1* Results in Liver Cysts

Mouse liver achieves the mature portal triad morphology of bile ducts, hepatocytes, and vascular structures and an adult liver weight/body weight (LW/BW) ratio in the first 2–4 weeks of postnatal life (17,18). We inactivated the *Pkhd1* gene at age 4 weeks using a tamoxifen-inducible, ubiquitously acting *UBC-Cre<sup>ER</sup>* that expresses in bile ducts and has previously been used to model PCLD with conditional *Pkd1* alleles (16). *Pkhd1<sup>f/f</sup>;UBC-Cre<sup>ER</sup>* mice sacrificed at age 17 weeks showed bile duct cysts throughout the liver (Figure 1). This cystic phenotype was fully penetrant and, in most cases, grossly visible as a stippled texture of the organ upon perfusion (Supplemental Figure 1). Normal ductal plate formation is reported in mice with the germline heterozygous *Pkhd1* genotype (6) and, indeed, tamoxifen-induced littermate mice with a monoallelic *Pkhd1<sup>f/+</sup>;UBC-Cre<sup>ER</sup>* genotype had normal-appearing livers, including portal triads consisting of artery, vein, and bile duct, without cysts (Figure 1, A and B). Histologic analysis of livers from *Pkhd1<sup>f/f</sup>;UBC-Cre<sup>ER</sup>* mice showed clusters of cysts among normal-appearing hepatocytes (Figure 1, A and B). *Pkhd1<sup>f/f</sup>;UBC-Cre<sup>ER</sup>* mice ( $n=11$ ) have an increased LW/BW ratio of  $5.7\% \pm 0.3\%$  compared with that of *Pkhd1<sup>f/+</sup>;UBC-Cre* mice ( $n=10$ ),  $4.4\% \pm 0.2\%$ ,  $P=0.0016$  (Figure 1C), without any effect of genotype on body weight (Supplemental Figure 2A). The female subgroup of *Pkhd1<sup>f/f</sup>;UBC-Cre<sup>ER</sup>* mice ( $n=4$ ) had the most significant difference from the noncystic *Pkhd1<sup>f/+</sup>;UBC-Cre<sup>ER</sup>* genotype female mice ( $n=3$ ),  $P<0.001$  (Figure 1C, right panel).

We further characterized the cystic phenotype by two complementary measures. To quantify the area of the cysts as an HCI, we applied MetaMorph software thresholding to scanned H&E-stained liver section images (Figure 1A) to measure the two-dimensional cystic area as a percentage of the total area of the liver section defined within a perimeter that excluded large vascular structures. Mean HCI was  $15.2 \pm 2.9$  in *Pkhd1<sup>f/f</sup>;UBC-Cre<sup>ER</sup>* mice ( $n=7$ ) versus  $3.14 \pm 0.4$  in



**Figure 1.** | Biallelic loss of *Pkhd1* with *UBC-Cre<sup>ER</sup>* results in diffuse liver cysts. (A) Representative liver sections of the indicated genotype at age 17 weeks. The colored boxes denote genotypes throughout the figure. (B) Hematoxylin and eosin-stained histology of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* versus control monoallelic deletion (*Pkhd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>*) at age 17 weeks. Representative structures marked as: v, vein; c, cyst. (C) Liver weight/body weight ratio of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* versus *Pkhd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* mice. Right panel, sex subgroups visually separated. Blue shapes, male; pink shapes, female. (D) Hepatic cystic index, the percentage of liver section area that is background density (cysts+venous structures) rather than parenchyma on liver lobe sections scanned in greyscale as in (A). (E) Anti-cytokeratin 19 (CK19) immunohistochemistry localizes to the biliary epithelium, which lines all cysts, and not venous structures. (F) Percentage of liver section area positive with CK19 staining. Bottom panel, sex subgroups of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice are visually separated. (G) Sirius Red stain at  $\times 10$  magnification.

*Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup> mice ( $n=7$ ),  $P=0.0015$  (Figure 1D). Immunohistochemical staining of CK19, a marker of biliary epithelium, showed that the liver cysts were uniformly lined with a CK19-positive epithelium, indicating that they were derived from the bile ducts (Figure 1E). In *Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup> mice, the CK19 staining highlighted the normal bile ducts, which represent a tiny minority of the cells in the liver (Figure 1E). Additional cells around the central veins were noted to be CK19-positive as has been defined in wild-type rodent liver CK19 immunostaining (19). To quantify the increased burden of biliary epithelium in our *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> mice, we again applied tissue perimeter outlining and color thresholding, this time to tissue sections stained with CK19. We calculated the CK19+%. Mean CK19+% was  $5.8 \pm 1$  in *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> mice ( $n=7$ ) versus  $0.5 \pm 0.1$  in *Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup> mice ( $n=7$ ),  $P=0.0003$  (Figure 1F).

Both HCI and CK19+% calculations were done by authors blinded to sample genotype on the first seven mice sacrificed for each genotype as only those had histologic sections. Nonzero values in the noncystic *Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup> mice were expected because of the inability of the semiautomatic method to distinguish venous versus cystic structures on two-dimensional histology, and the expected presence of CK19 staining of native bile ducts. HCI and CK19+% were far superior to LW/BW at numerically quantifying the cystic phenotype distinguishing *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> and *Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup> mice. For both HCI and CK19+%, the comparison between genotypes maintained statistical significance ( $P<0.05$ ) even in the sex subgroups with small numbers. CK19+% was the one parameter by which the trend toward increased severity of liver cysts in cystic females versus cystic males met statistical significance ( $P=0.016$ ) despite the small numbers once split by sex (Figure 1F). There was a robust positive correlation ( $r^2=0.47$ ), between the CK19+ area and the HCI values for each mouse showing that these measures were congruent with each other (Supplemental Figure 2B).

As the lesion in recessive polycystic kidney disease is CHF, we also sought to assess whether there was any significant fibrosis in our model. *Pkhd1* null mice develop a significant burden of fibrosis as defined by Sirius Red staining between 3 and 9 months of age (6,20). We stained paraffin sections with Sirius Red stain and Masson's trichrome stain (data not shown) to localize collagen. Sirius Red stained sections show positive staining outlining liver margins, central veins, and cysts (Figure 1G). There was no additional collagen staining beyond these structures on Sirius Red or Masson's trichrome-stained sections, suggesting an absence of significant fibrosis in this model.

#### Analysis of Postdevelopmental *Pkhd1* Loss in the Kidney

Germline *Pkhd1* null mouse kidneys have no phenotype for at least 6 months of life, or not at all (6,8,9). In keeping with this, the kidneys of the *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> mice at 17 weeks of age showed no gross, histologic, or kidney weight/body weight ratio abnormalities. *Pkd1*<sup>+/-</sup> mouse kidneys are noncystic other than a potential small number of discrete cysts if aged over 9 months (21). The combined *Pkhd1*<sup>-/-</sup>; *Pkd1*<sup>+/-</sup> genotype has diffusely cystic kidneys (7,13). In order to evaluate whether *Pkhd1* plays a role after development in the kidney, we made use of this established

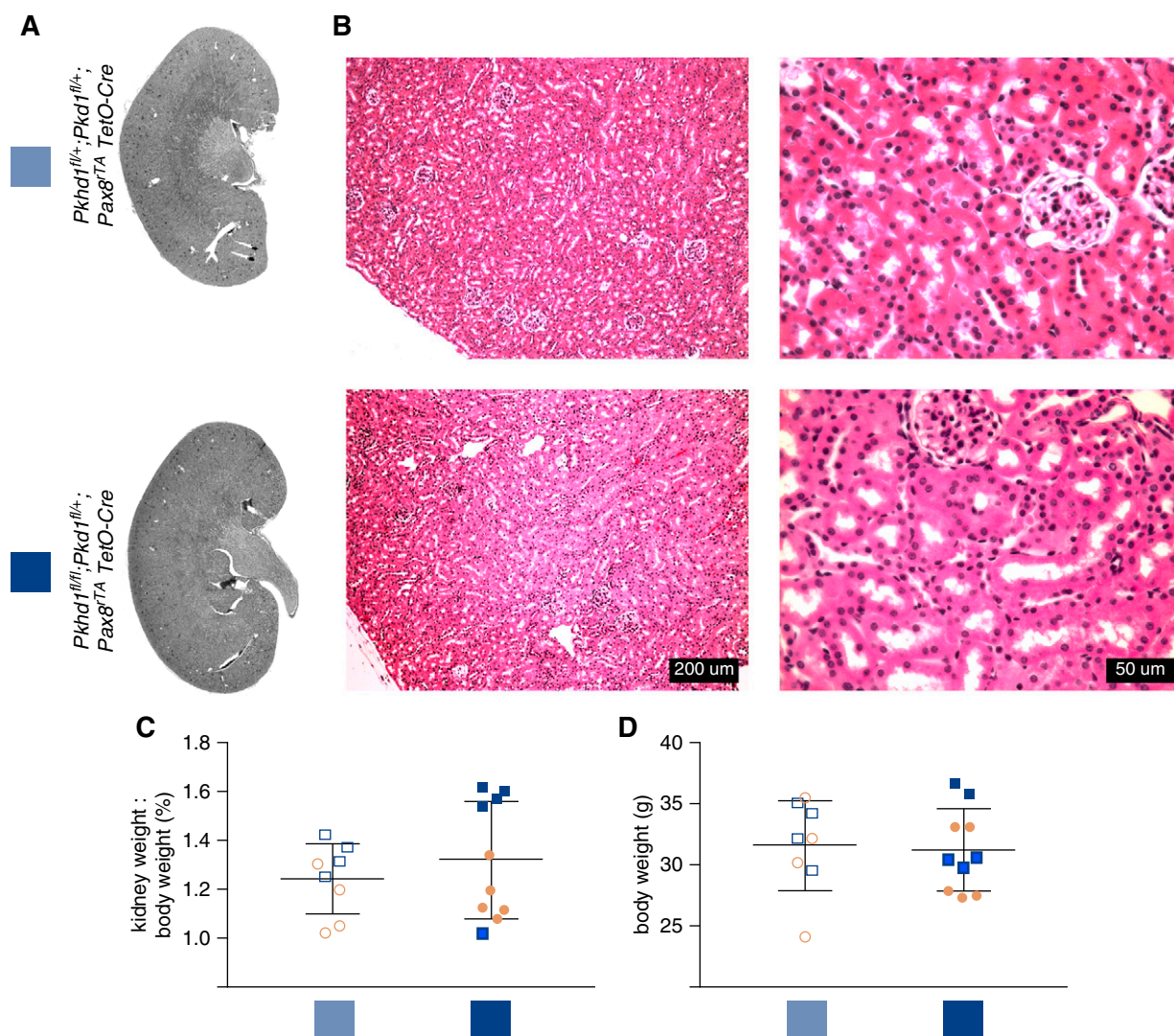
genetic interaction between *Pkhd1* and *Pkd1*. To make the conditional equivalent, we used *Pkhd1*<sup>fl</sup> and *Pkd1*<sup>fl</sup> alleles with the doxycycline-inducible *Pax8*<sup>rtTA</sup>TetO-*Cre* system. This model has Cre activity in the majority of renal tubule segments including the collecting duct. We induced *Pkhd1*<sup>fl/fl</sup>; *Pkd1*<sup>fl/+</sup>; *Pax8*<sup>rtTA</sup>TetO-*Cre* and *Pkhd1*<sup>fl/+</sup>; *Pkd1*<sup>fl/+</sup>; *Pax8*<sup>rtTA</sup>TetO-*Cre* littermates with doxycycline for a 2-week duration from 4 to 6 weeks of age. We sacrificed the mice at age 6 months. Histologic analysis of the perfusion-fixed kidneys showed no apparent abnormalities in the renal tubules, interstitium, or glomeruli (Figure 2, A and B). There was no difference in the kidney weight/body weight ratio (Figure 2C). This ratio was higher in male mice ( $n=9$ ) than female mice ( $n=9$ ),  $P=0.0039$ , regardless of genotype. Scanned kidney sections from this experiment are shown in Supplemental Figure 3.

#### *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre* Model Liver Cyst Phenotype Is Independent of *Pkd1* Dosage

ARPKD mouse liver is more severely affected when *Pkd1* gene dosage is reduced (13). Specifically, the liver phenotype in *Pkhd1*<sup>del4/del4</sup>; *Pkd1*<sup>+/-</sup> is more severe than in *Pkhd1*<sup>del4/del4</sup>, even though the *Pkd1*<sup>+/-</sup> genotype itself has no liver phenotype. Increased *Pkd1* dosage in the form of a three-copy *Pkd1*<sup>F/H</sup>-BAC rescued conditional mouse models of PCLD disease genes *PRKCSH* and *SEC63*, but had no effect on germline ARPKD mouse models (*Pkhd1*<sup>del4/del4</sup>; *Pkd1*<sup>F/H</sup>-BAC) (13). However, given that the *Pkhd1*<sup>del4/del4</sup> mouse liver develops abnormally, this does not exclude the possibility that the postdevelopmental loss of *Pkhd1* modeled in this study could be rescued by increased *Pkd1* dosage. To assess the effect of *Pkd1* dosage on our postdevelopmental *Pkhd1* model, we generated *Pkhd1*<sup>fl/fl</sup>; *Pkd1*<sup>fl/+</sup>; UBC-*Cre*<sup>ER</sup> and *Pkhd1*<sup>fl/fl</sup>; *Pkd1*<sup>F/H</sup>-BAC; UBC-*Cre*<sup>ER</sup> mice to compare their littermates with the cystic *Pkhd1*<sup>fl/fl</sup>; UBC-*Cre*<sup>ER</sup> genotype. We induced these mice with tamoxifen injections at age 4 weeks, identical to our prior experiments, and followed them until 17 weeks of age. There was no appreciable difference in the cystic liver phenotype between these genotypes as assessed grossly by LW/BW or by HCI (Figure 3, A–C, Supplemental Figures 4 and 5). The body weight was also unchanged between these three genotypes (Figure 3D).

#### Bile Duct and Extrahepatic Findings

We performed gross and histologic analysis of the spleen and pancreas, in addition to the liver and kidney, in a subset of *Pkhd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup>, *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup>, *Pkhd1*<sup>fl/fl</sup>; *Pkd1*<sup>fl/+</sup>;UBC-*Cre*<sup>ER</sup>, and *Pkhd1*<sup>fl/fl</sup>; *Pkd1*<sup>F/H</sup>-BAC; UBC-*Cre*<sup>ER</sup> mice. Germline *Pkhd1* null mouse models have an age-dependent penetrance of common bile duct dilation, cholangitis, pancreatic cysts, and splenomegaly in the setting of portal hypertension (7–9). In our postdevelopmental *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> model, we did not find common bile duct dilation or splenomegaly. Spleen histology appeared normal (Figure 3E). Considering liver histology on 67 *Pkhd1*<sup>fl/fl</sup>;UBC-*Cre*<sup>ER</sup> mice with or without concurrent alterations in *Pkd1* dosage, we noted findings consistent with varying degrees of cholangitis in four mice. This ranged from very mild focal pericyclic and intracyclic infiltrates including PMNs ( $n=3$ ) to more diffuse histologic findings



**Figure 2.** | *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;Pax8<sup>rtTA</sup>;TetO-Cre* has no kidney phenotype. (A) Representative kidney sections of the indicated genotype at 6 months of age. (B) Hematoxylin and eosin-stained kidneys from *Pkhd1<sup>fl/fl</sup>; Pkd1<sup>fl/+</sup>;Pax8<sup>rtTA</sup>;TetO-Cre* mice at 6 months of age. (C) Kidney weight/body weight ratio and (D) body weight of *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;Pax8<sup>rtTA</sup>;TetO-Cre* mice. Genotypes indicated by color corresponding to labels in (A). Blue shapes, males; pink shapes, female.

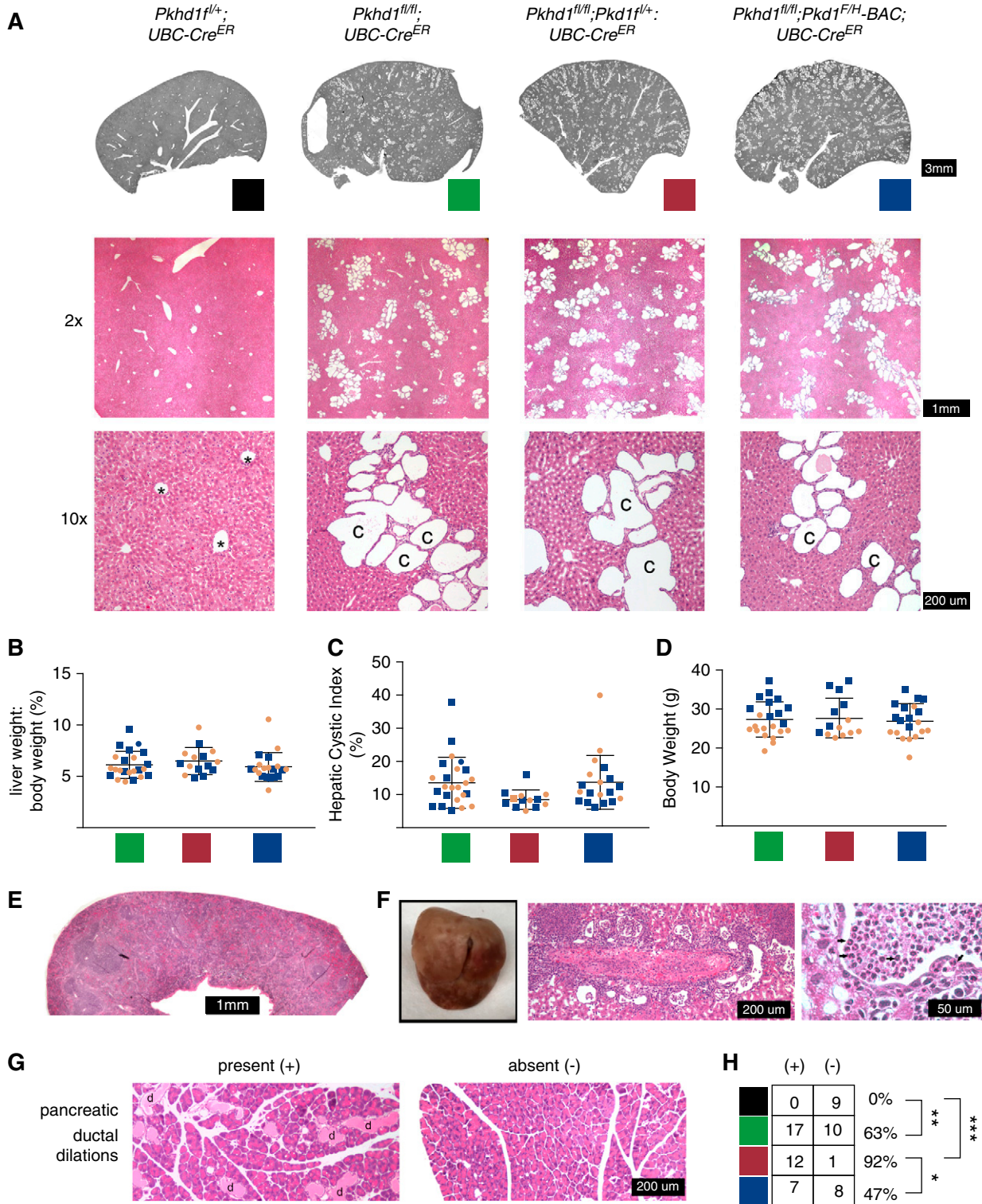
and grossly visible patchy erythema on the liver surface ( $n=1$ , Figure 3F). The four mice in which this was noted each had biallelic *Pkhd1* loss, but included at least one mouse of each sex and *Pkd1* dosage tested.

Histology of the pancreas in *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice demonstrated microscopic ductal dilations containing eosinophilic material, albeit with incomplete penetrance. The gross appearance of the pancreas was unremarkable. Figure 3G shows representative histologic images of mouse pancreatic tissue with or without pancreatic ductal dilations. This phenotype was either notable throughout the sampled tissue or it was absent. This pancreatic ductal dilation was seen in 17 of 27 (63%) of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice, but was never seen in *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* pancreas ( $n=7$ ),  $P=0.007$  (Figure 3H). The pancreatic ductal dilations were seen in a significantly higher percentage of the mice with reduced *Pkd1* compared with those who had extra copies of *Pkd1*: 12 of 13 (92%) *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>; UBC-Cre<sup>ER</sup>*

mice versus 7 of 15 (47%) *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>F/H</sup>-BAC UBC-Cre<sup>ER</sup>* mice,  $P=0.016$ .

## Discussion

This study is the first to evaluate whether *Pkhd1*, and thus its protein product fibrocystin, continue to play a critical role in biliary homeostasis in the developed liver. We show that grown mice that lose *Pkhd1* after normal liver development have a dramatic and fully penetrant cystic liver phenotype. Whereas ductal plate malformation is known to be involved in the pathogenesis of CHF, we now show that it is not required for the *Pkhd1*-related liver cysts themselves. This finding could serve to distinguish the pathogenesis of cysts that are seen in variable degrees in patients with ARPKD from the fibrotic phenotype of CHF. CHF accompanied by cysts is a distinguishing feature of ARPKD unique from other genetic causes of such hepatic fibrosis. Further, this



**Figure 3. | Liver phenotype does not change with alterations of *Pkd1* in *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice.** (A) Liver section scan and hematoxylin and eosin-stained histology for the indicated genotypes. Representative structures marked as: \*, vein; c, cyst. Colors indicated for each genotype are used throughout the figure. (B) Liver weight/body weight ratio. (C) Hepatic cystic index as defined in Figure 1. (D) Body weight. (E) Hematoxylin and eosin-stained histology of spleen at  $\times 2$  magnification. (F) Patchy appearance of liver in one *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* mouse affected by cholangitis. Middle and right panels, hematoxylin and eosin-stained histology shows cystic biliary structures full of polymononuclear lymphocytes (arrows). (G) Pancreatic ductal dilations were seen in a subset of mice with *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* with or without alterations in *Pkd1* genotype. Hematoxylin and eosin-stained histology of pancreas at  $\times 10$  magnification representative of presence or absence of ductal dilations. d, ductal dilation. (H) Quantification of number of mice of each genotype with pancreatic ductal dilations present or absent.

finding demonstrates that the development of the recessive genotype in adulthood, as would be the case in a heterozygous individual who has a somatic mutation to the normal allele, likely results in cysts as is known to occur in PCLD. One limitation to conditional mouse models of dominantly inherited polycystic kidney disease and PCLD is that all biliary epithelial cells lose the disease gene simultaneously, in contrast to human organs in which somatic mutations occur in individual cells. Given this, we cannot yet definitively conclude that a single cell affected with *PKHD1* loss is sufficient to initiate a cyst as is known to be the case in patients with *PKD1* and *PKD2*. Nonetheless, a recent study followed *Pkhd1* heterozygous mice by magnetic resonance imaging and although they showed no abnormalities at 10 months of age, by 1.5 years when somatic mutations could have occurred with time for subsequent cyst growth, the mice had cysts analogous to those found in human heterozygotes (22). This finding in mice, together with the focal nature of some cysts seen in *PKHD1* mutation carriers, supports the idea that recessive loss in individual cells is sufficient to form cysts.

Our discovery of *PKHD1* as an autosomal dominant PCLD disease gene prompted the investigations in this study. Unlike all other established PCLD disease genes, *Pkhd1* does not cause PCLD by affecting polycystin-1 (PC1) maturation (12,23). Biallelic loss of *PrkcsH*, *Sec63*, *Alg8*, *Ganab*, *Sec61b*, *Dnajb11*, or *Alg9* results in significant reduction of the expression of mature PC1, whereas biallelic loss of *Pkhd1* does not (12,13,24–26). Instead, we asked the question of whether *Pkhd1* loss affected the PC1 functional dosage without affecting PC1 protein levels of trafficking. We found that in contrast to mouse models of *PrkcsH* and *Sec63*, in which the presence of extra genomic copies of *Pkd1* in the form of the *Pkd1<sup>F/H</sup>-BAC* rescues the liver cystic phenotype, the extra copies of PC1 had no effect on bile duct cysts because of loss of *Pkhd1*. Similarly, whereas reduced *Pkd1* dosage worsens PCLD because of *PrkcsH* and *Sec63*, it has no effect on *Pkhd1* liver cysts resulting from adult inactivation. This shows that the *Pkhd1*-related liver cystic phenotype is independent of PC1 dosage and thus likely occurs *via* a non-PC1-dependent mechanism.

Adult-inducible inactivation of *Pkhd1* does not recapitulate the genetic interaction between *Pkhd1* and *Pkd1* that occurs with germline *Pkhd1<sup>-/-</sup>;Pkd1<sup>+/-</sup>* mice. The loss of one *Pkd1* allele was not able to bring out a kidney phenotype in our *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;Pax8<sup>rtTA</sup>TetO-Cre* system, and the cystic liver phenotype was no worse in *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* mice than those with two functional copies of *Pkd1*. This suggests that the previously described genetic interaction between the *Pkhd1<sup>-/-</sup>* and *Pkd1<sup>+/-</sup>* genotypes in both the kidney and liver is not involved in adult tissue homeostasis, but rather requires abnormal organogenesis as likely occurs with the germline null *Pkhd1* alleles. Both patients with ADPKD and mice, and ARPKD mice, develop pancreatic cysts with incomplete penetrance (7–9,27,28). Pancreatic ductal dilation in our postdevelopmental *Pkhd1* model occurred with significantly higher frequency in mice with *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;UBC-Cre* versus *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>F/H</sup>-BAC UBC-Cre<sup>ER</sup>* genotypes, even though there was no appreciable difference in severity. This was the only parameter in the *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice that differed with PC1 dosage in adult inactivation of *Pkhd1*.

We describe a novel mouse model of *Pkhd1*/fibrocystin inactivation after liver development. We learn that *Pkhd1* plays a crucial ongoing role in biliary homeostasis. This may suggest that the pathogenesis of ARPKD liver cysts is not intertwined with that of CHF. Further, our findings support the possibility that the polycystic liver phenotype in *PKHD1* heterozygous carriers could occur through somatic mutation, and the lack of *Pkhd1/Pkd1* genetic interaction in this model lends support to a hypothesis that liver cyst formation in ARPKD and ADPKD occur through different mechanistic pathways. Such investigations are relevant to understanding whether future targeted treatments for ADPKD should be expected to be relevant to ARPKD. The robust liver cystic phenotype in our *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* mice makes this a valuable model for future studies to compare the functional pathways of *Pkhd1* and *Pkd1* and their treatment.

#### Disclosures

S. Somlo is a founder, shareholder, and consultant for Goldfinch Bio, and reports personal fees from Otsuka Pharmaceuticals and Goldfinch Bio, outside the submitted work. All remaining authors have nothing to disclose.

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#### Author Contributions

W. Besse, S. Ghosh Roy, C. Roosendaal, and L. Tuccillo were responsible for investigation; W. Besse was responsible for formal analysis, funding acquisition, methodology, visualization, and wrote the original draft; A. Gallagher was responsible for data curation; S. Somlo was responsible for funding acquisition and methodology; W. Besse, A. Gallagher, and S. Somlo conceptualized the study, and reviewed and edited the manuscript.

#### Supplemental Material

This article contains the following supplemental material online at <http://kidney360.asnjournals.org/lookup/suppl/doi:10.34067/KID.K3602020000252/DCSupplemental>.

Supplemental Figure 1. Liver histology of all *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* and *Pkhd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* mice evaluated by this method.

Supplemental Figure 2. Additional information for the *UBC-Cre<sup>ER</sup>* model.

Supplemental Figure 3. Kidney histology of all *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;Pax8<sup>rtTA</sup>TetO-Cre* mice in the study.

Supplemental Figure 4. Males from contemporaneous mouse cohort of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* and *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>F/H</sup>-BAC UBC-Cre<sup>ER</sup>*.

Supplemental Figure 5. Females from contemporaneous mouse cohort of *Pkhd1<sup>fl/fl</sup>;UBC-Cre<sup>ER</sup>* *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>fl/+</sup>;UBC-Cre<sup>ER</sup>* and *Pkhd1<sup>fl/fl</sup>;Pkd1<sup>F/H</sup>-BAC UBC-Cre<sup>ER</sup>*.

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