Repair after nephron ablation reveals limitations of neonatal neonephrogenesis

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Introduction

The number of nephrons in the human kidney, once established, does not increase during life but rather declines with age through attrition or disease. The “nephron endowment” is determined by 36 weeks of gestation (1), although new nephron formation after birth in humans has been suggested (2). Low birth weight (from any cause, including preterm birth) is strongly correlated with reduced nephron endowment (3, 4) and results in hyperfiltration from birth (5), leading to an increase risk of essential hypertension (6–9) and chronic kidney disease later in life (10, 11); these infants are thus placed at long-term risk for disease. Currently, there is no known method to increase nephron number in newborn or adult humans. Establishing an experimental system that induced neonephrogenesis postpartum would be an important first step in the development of mechanism-based therapeutics to augment nephron number in patients.

In the newborn mouse, nephron formation (nephrogenesis) has been postulated to continue until P3 (taking P1 as the day of birth), with the presence of Six2+ nephron progenitor cells and early nephron precursor structures, including renal vesicles, being seen as circumstantial evidence for de novo nephrogenesis (12). Fifty-one percent of mature nephrons are formed between P0 and P4, as shown in 3D modeling studies (13). Maturation of the nephrons continues after Six2+ cells are no longer present, and nephron growth accounts for the increase in kidney size. While de novo nephrogenesis is a possible contributor to kidney growth, an alternative explanation would be that the cap mesenchyme contains already fate-determined nephron progenitors that are set to complete nephrogenesis with the number of nephrons predetermined at birth by a multitude of different factors, including nutrition, oxygen tension, and genetics (14). This distinction is important because it has consequences for studies in regenerative medicine when evaluating the nephron-forming potential in the newborn mouse. Studies in the 1970s utilizing different nephron-counting procedures were discordant in demonstrating nephrogenic potential after birth with or without stimulation through contralateral nephrectomy (15). While the mouse model offers a postnatal window to investigate and potentially manipulate the final stages of nephrogenesis, formal lineage-tracing studies unequivocally demonstrating the extent of nephrogenesis after birth have never been conducted. The signals responsible for the cessation of nephrogenesis are unknown. It has been reported that BMP7...
pathway inhibition with LDN-193189, which blocks SMAD1/5, results in an extension of the time after birth that CITED1+/SIX2+ nephron progenitors are present and an increase in nephron number (16). It is unclear whether this increase occurs with additional ureteric bud branching or a further increase in the number of nephrons per ureteric bud tip. Furthermore, it remains unclear whether the kidney has the ability to increase neonephrogenesis after birth when stimulated, for instance, following the reduction of kidney mass by cryoinjury. An evaluation of the endogenous ability of the kidney to modulate neonephrogenesis would offer potential paths for pharmacological interventions to prolong nephrogenesis in the premature birth neonate and potentially provide insight into how to replace nephrons in the adult.

In newborn animals, regenerative mechanisms have been reported for some organs. For example, newborn hearts in mice have been reported to regenerate cardiomyocytes and heal without fibrosis after resection of the tip, an ability that is lost in the first week after birth (17). Others, however, have challenged this observation (18). It has been proposed that organ formation and growth are regulated according to the size and functional demand of the newborn (19). Recent work suggests that nephron progenitors can be provided with signals that extend their self-renewal, raising the possibility of extending the neonephrogenesis in vivo (16). We hypothesized that reducing nephron mass during active nephrogenesis would alter this putative regulation, resulting in compensatory new nephron formation, a response that would be only possible during the early neonatal period when nephron progenitors were present.

To test this hypothesis, a model of cryoablation in newborn mice was developed to create a reproducible and localized injury to the kidney that reduces total kidney nephron mass. Acute reduction in nephron mass within 24 hours of birth resulted in an adaptive hypertrophic response and healing, with small amounts of fibrosis detected at 1 week (P7) and 3 weeks (P21). If contralateral nephrectomy was performed at the time of injury, the healing appeared more pronounced, with no signs of scarring by P21. By contrast, when cryoinjury-induced nephron reduction was performed at 1 week (P7), more extensive scarring was observed at P21. Genetic lineage-analysis of the nephron progenitor cell population and marker analysis by immunofluorescence analysis was done to assess the regenerative response of the newborn kidney. Whether the injury occurred within 24 hours of birth or at 7 days of age we detected no enhanced production of new nephrons, as measured by glomerular number, which remained reduced. Thus, the reduction of nephron number in the neonatal period when nephron progenitors are present does not stimulate de novo creation of additional glomeruli or other nephron segments. Rather, the newborn mouse responds to injury by compensatory hypertrophy of remaining nephrons.

Results
Cryoinjury to reduce nephron mass in newborn mice. In order to examine the regenerative potential of the newborn kidney, we developed a surgical approach using cryoinjury and nephrectomy. Newborn mice (we designate the day of birth as “P1,” which includes the first 24 hours of life) were subjected to cryoinjury using a liquid nitrogen–cooled probe placed on the kidney surface for 3 seconds. This procedure led to a highly reproducible area of coagulative necrosis in the kidney (Figure 1, A and B). Twenty-four hours after injury, a necrotic area was documented involving approximately 15%–20% the size of the cortex and penetrating to the medullary area (Figure 1A). To assess the extent of injury beyond the necrotic area, we performed immunofluorescence analysis to detect KIM-1 (Havcr1), a marker of tubular injury (20), and LTL, a marker of proximal tubules (21). This revealed a penumbra of containing KIM-1+/LTL+ proximal tubular cells, indicating viable but injured tubules around the necrotic area (Figure 1, C and D).

Kidney architecture is restored after injury. In the mouse, it is generally accepted that the induction of new nephrons, or active neonephrogenesis, continues to occur during the first 3 days after birth while the nephron progenitors are still present (12, 13, 16, 22). The timeline of our experiments, as indicated in Figure 1E, was designed to evaluate whether there was potential for “de novo” induction of new nephrons in response to injury during this period. Mice were subjected to either newborn (P1) or late (P7) cryoinjury as a stimulus for repair processes that might include new nephron formation. Cryoinjury was performed on P1, when Six2+ cells were present in the cap mesenchyme (Figure 1F). There was some variability in the exact timing of Six2 downregulation, likely due to the fact mothers delivered naturally during the night, resulting in a small range of delivery times. We examined the injury response through the first week (P3–P7) and at weaning age (P21). At P5, a time point 2 days after the Six2+ nephron progenitor cells have been exhausted during normal development (12, 13) and beyond the expected time of Six2 positivity with normal variation, we observed a persistence of only a few dispersed Six2+ cells in a subset of cryoinjured mice (2 of 6) (Figure 1G).
Coagulative necrosis was apparent 1-day after injury. This was followed by reconstitution of the kidney architecture from the edges of the injury. This was discernible by day 3 after injury, with only a small residual indentation remaining on the kidney surface at the injury site at 1 week of age (P7) (Figure 2A). The weights of the injured left kidneys (normalized to body weight) were initially not significantly different from the weights of the contralateral control kidneys (Figure 2B) but started to lag behind the control kidneys 3 days after injury and remained decreased compared with contralateral controls on P8 and P21 (Figure 2, B and C), despite gross anatomical restoration of tissue architecture. Uninjured contralateral control kidneys did not undergo significant hypertrophy, as reflected by kidney-weight-to-body-weight ratios that were similar to sham-operated kidneys. The lack of detectable hypertrophy in the contralateral kidney was also seen when animals were cryoinjured on P7 (Figure 2C).

We hypothesized that the continued presence of an uninjured contralateral kidney might inhibit the regenerative response. To investigate this, we performed unilateral nephrectomy with or without cryoinjury. Unilateral nephrectomy in newborn animals without cryoinjury caused compensatory hypertrophy in the kidney, as indicated by an increased kidney-weight-to-body-weight ratio (compare sham with sham/nephrectomy in Figure 2C). Unilateral nephrectomy at the time of cryoinjury also resulted in compensatory hypertrophy in the injured kidneys, as indicated by a kidney-weight-to-body-weight ratio that did not significantly differ from that of uninjured kidneys after contralateral nephrectomy (Figure 2, C and D). The total nephron number 1 week after newborn cryoinjury was determined by glomerular counting using the acid maceration method (23, 24). This method is based on glomerular counting of a sample of whole kidney tissue to minimize bias. An example of a sample used for counting is shown in Supplemental Figure 1 (supplemental material available online with this article; doi:10.1172/jci.insight.88848DS1). The injured kidneys had a significantly decreased number of total glomeruli compared with contralateral control kidneys (Figure 2E). Sham animals had glomerular numbers comparable to those of the uninjured contralateral control kidneys, without a significant difference between the right and left kidney. In addition, contralateral
Figure 2. Cryoinjured kidneys exhibit reduced kidney mass and glomerular number. (A) Paraffin sections of kidney parenchyma after cryoinjury, with contralateral kidney intact, show a progressive size reduction in the injured necrotic area (asterisks) compared with the surrounding kidney parenchyma. On P7, there was resolution of the injury, with only a small indentation remaining at the site of the injury. Original magnification, ×3. (B) Kidney weight progressively declined in the first 7 days after injury compared with that in uninjured control kidneys and sham controls. n = 104 animals, n > 6 per time.
nephrectomy did not affect glomerular number in injured kidneys, as there was an equivalent decrease in glomerular number despite an increased kidney weight. Overall, the absence of an increase in glomerular counts indicated compensatory hypertrophy rather than new nephron formation as the mechanism for restoration to normal kidney weight (Figure 2D).

To determine the long-term implications of the cryoinjury and subsequent whole kidney development after injury, animals were followed into adulthood and euthanized at 12 weeks. The kidney-weight-to-body-weight ratio increased with age in both injured and sham controls. The differences in kidney-weight-to-body-weight ratios observed at 1 week after injury were still evident at 12 weeks after left kidney injury (Figure 2F). Additionally, we examined kidneys subjected to resection of a kidney lower pole as a method to reduce nephron mass at birth. At 5 weeks after resection, kidney weight of the resected kidneys remained reduced compared with contralateral controls (Figure 2G), indicating a complete lack of a compensatory response to this type of injury also.

An increase in the kidney-weight-to-body-weight ratio after injury could be caused by a hypertrophic response solely, increased epithelial cell proliferation, or de novo nephron formation in response to injury. Having ruled out the last possibility, we examined the proliferative response by staining for Ki67 to determine if accelerated proliferation accounts for restitution of kidney architecture after cryoinjury (Figure 3A). The number of proliferating cells in the area of cortex or outer medulla surrounding the injury was not significantly different between injured kidneys and uninjured contralateral controls or sham controls on P3, P4, P6 and P8, which is consistent with a hypertrophic, rather than hyperplastic, response to the injury (Figure 3B). This hypertrophic response is greater if the contralateral kidney is removed.

Developmentally regulated genes are not altered in response to cryoinjury. Expression patterns of master regulator genes expressed during kidney development were assessed by immunofluorescence and in situ hybridization to identify cells engaged in nephrogenesis. The transcription factor Pax2 is expressed during nephrogenesis and regulates ureteric bud branching and outgrowth (25). The pattern of expression of Pax2 in injured and contralateral control kidneys from P3 to P6 was not affected by cryoinjury. Pax2 expression was seen in cap mesenchyme cells in the subcortical area as well as in collecting duct cells extending into the medulla (Figure 4A). Lhx1 plays an essential role in nephrogenesis and is expressed in the intermediate mesoderm and in the nephric mesenchyme (26). After cryoinjury, Lhx1 transcripts are equally expressed in injured and contralateral control kidneys in the cap mesenchyme until P6, with no expression at P7 or later in injured kidneys, contralateral kidneys, and sham control kidneys (Figure 4B).

Persistent Six2+ cells at the injury site do not form additional new nephrons. To determine the fate of Six2+ nephron progenitors after cryoinjury, we used the tamoxifen-dependent Cre line, Six2CRE (27), to lineage trace these cells. Six2CRE/+ heterozygous mice were crossed to the Cre-responsive R26RtdTomato/tdTomato homozygous reporter mice, resulting in Six2CRE/+; R26RtdTomato/+ double heterozygous animals and R26RtdTomato/+ littermate controls (Figure 5A). Tamoxifen dosing was used to label Six2+ cells prior to or during injury, permanently labeling Six2+ descendant daughter cells from that time forward with tdTomato expression. Glomeruli that were tdTomato+ were therefore formed after tamoxifen-induced labeling of the Six2+ nephron progenitors. We measured the percentage of glomeruli formed after tamoxifen induction by staining all glomeruli by immunofluorescence (Figure 5C) and determining the number of tdTomato/podocalyxin double-positive glomeruli. When control pregnant mice were administered tamoxifen at E17.5, approximately 50% of glomeruli were labeled with tdTomato when examined on P7, indicating that the contribution of Six2+ progenitor cells at day 17.5 is due to ongoing nephrogenesis (Figure 5B). R26RtdTomato/+ littermate controls given tamoxifen did not show any labeling. Thus, about half of nephrons are formed after E17.5 from the Six2+ cap mesenchyme in our system, approximating the
shown), there were similar amounts of Six2-derived tdTomato+ single cells in the outer cortex of cryoinjured subject to cryoinjury on P1, with tamoxifen injected at the end of nephrogenesis (P3–P4, n = 3–5 animals per group). In box-and-whisker plots, whiskers depict the minimum and maximum values, boxes depict 25th to 75th percentiles, and lines depict the median. Ordinary 1-way ANOVA, Tukey’s multiple comparison test.

We next determined the fate of the rare dispersed Six2+ cells found in the area adjacent to the injury by labeling Six2+ cells 2 days after injury. If the dispersed Six2+ cells entered into a nephrogenic program, we would predict fully labeled nephrons to form. However, in animals subjected to cryoinjury on P1, with tamoxifen injected at the end of nephrogenesis (P3–P4, n = 4, data not shown), there were similar amounts of Six2-derived tdTomato+ single cells in the outer cortex of cryoinjured and contralateral kidneys. (Figure 5, D and E, right). No increase in the number of completely labeled tdTom+ nephrons was present, indicating absence of de novo nephrogenesis. tdTom+ cells were scattered as single cells within glomeruli and tubules of nephrons, indicating that Six2+ cells were already lineage committed and not truly multipotent anymore, giving rise only to parts of nephrons without stem cell self-renewal.

Increased development of fibrotic lesions after injury in 1-week-old (P7) mice. To characterize the development of fibrosis following cryoinjury, we examined kidneys at 3 weeks and 12 weeks after injury induced at either P1 or P7, with or without contralateral nephrectomy. When cryoinjury was performed on P1 newborns without contralateral nephrectomy, the injured kidneys recovered either without noticeable surface abnormalities or had, in about 50% of the cases, a small residual macroscopic indentation at the site of cryoinjury 3 weeks after surgery (Figure 6A). Microscopically, Sirius red–stained wedges were observed at the injury site, with an indentation on the cortical surface. When contralateral nephrectomy was performed at the time of injury, fibrosis was not detected when injury was induced in P1 newborns (Figure 6A, P1 cryoinjury/nx), indicating that contralateral nephrectomy prevented the development of fibrosis. In contrast, when injury was induced on P7, macroscopic injury persisted to week 3, even with contralateral nephrectomy (compare Figure 6A, P1 cryoinjury/nx with P7 cryoinjury/nx). In addition, Sirius red staining and α-SMA staining at 3 weeks were increased in the P7 injured kidneys in the presence of contralateral nephrectomy, indicating that contralateral nephrectomy did not prevent fibrosis when injury occurred at the P7 time point. When P1 and P7 injured animals were examined at a longer time point (12 weeks after injury), a region of cortical thinning was noted in all kidneys and a persistent trichrome+ area remained in the cortex at the injury site in approximately one-third of the P7 injured animals (Figure 6B).

Discussion

Nephrogenesis necessitates the interplay of different progenitor cell populations in a coordinated fashion, with reciprocal induction and maturation into various differentiated cell types. It is estimated that there are 26 highly specialized cell types populating a nephron, making nephrogenesis one of the most detailed and specialized processes in organogenesis (29). Factors that govern the final number of nephrons at birth are unknown, but in humans low nephron endowment correlates with low birth weight, female gender, short adult stature, and prematurity (30). While in vitro approaches for the differentiation of stem cells to kidney structures are showing promise (31–34), facilitating nephrogenesis in vivo would be a highly desirable and efficient tool in treating disease. The de novo generation of nephrons to substitute for the low nephron
number associated with low birth weight or nephrons lost to acute or chronic injury is a priority of regenerative medicine in nephrology and would open new avenues for therapy for millions of people.

In contrast to the human, in which de novo nephrogenesis is complete by 36 weeks of gestation, it has been known that there are nephron progenitor cells and immature nephron forms present in the rodent kidney at the normal day of parturition. New nephrons are continuously being generated during kidney development, with the induction of nephron progenitor cells in the cap mesenchyme through interactions with the ureteric bud and the surrounding cortical interstitium. Recent work suggests this niche depends on growth factor signaling pathways (16). This raised the possibility that we might be able to prolong nephrogenesis after parturition by invoking an intrinsic repair response to acute reduction of nephron endowment and learn how we might do so in humans from this response.

The kidney has significant capacity for repair after injury, yet lost nephrons cannot be replaced, leading to decreased function and, together with progressive scarring, end-stage renal disease (35). In rodent models, examples of adaptive regeneration that have been reported to be present exclusively in neonates include replacement of heart tissue (17) and regeneration of fingertips (36) and cochlear cells (37). Importantly, these capabilities are lost with maturation.

**Figure 4. Expression of developmentally regulated genes is unaltered in response to cryoinjury.** (A) Immunofluorescence staining for Pax2 and (B) in situ hybridization for Lhx1 were used to determine expression patterns around the injury area compared with contralateral control kidneys. (A) Pax2 expression is not different around injured areas compared with controls (red, Pax2; green, DBA; white, LTL). Original magnification, ×20. (B) Lhx1 is not differentially expressed after injury, as determined by in situ hybridization. Original magnification, ×3. Asterisks indicate site of injury. DBA, Dolichos Biflorus Agglutinin; LTL, Lotus Tetragonolobus Agglutinin.
Figure 5. Lineage tracing to determine nephron formation. (A) Breeding strategy of transgenic animals. Six2 heterozygous (Six2^{GCE/+}) mice were crossed to tdTomato reporter (R26R^{tdTomato/tdTomato}) mice. Injury was induced on the day of birth (P1) followed by tamoxifen injection to induce recombination in Six2^{+} cells. Kidneys were harvested at P7. tdTomato is expressed permanently in all cells that were Six2^{+} at the time of tamoxifen injection and their daughter cells. (B) Negative controls injected with tamoxifen at E17.5 showed no recombination, as expected (left). Positive controls injected at E17.5 had approximately 50% of nephrons labeled positive with tdTomato (middle). Animals injected with tamoxifen at P1 had 46% of glomeruli labeled by tdTomato (right). Original magnification, ×4. (C) Podocalyxin staining in combination with tdTomato fluorescence was used to determine the number of tdTomato-labeled glomeruli as surrogate number for labeled nephrons. An overview of a kidney with the red channel on the left side (tdTomato) and the far-red channel (podocalyxin) on the right side (left). Higher-magnification images showing two dual-labeled (tdTomato and podocalyxin) glomeruli (arrowheads) as well as several non-tdTomato-labeled glomeruli. Original magnification left panel, ×4; right panel, ×20. (D) The percentage of tdTomato-labeled glomeruli was not significantly different in animals subjected to cryoinjury (left) compared with uninjured control animals (right). Original magnification, ×4. (E) When tamoxifen was injected at P3 in animals subjected to cryoinjury, we found only single dispersed tdTomato^{+} cells in the outer cortex. Original magnification, ×10. (F) Determination of tdTomato-labeled glomeruli was done using podocalyxin staining (green). Double-labeled glomeruli (podocalyxin and tdTomato) appear yellow (left). The percentage of tdTomato-labeled glomeruli was not significantly different after injury compared with control animals or contralateral control kidneys (right panel). Original magnification, ×4. n = 6 animals per group. In box-and-whisker plots, whiskers depict the minimum and maximum values, boxes depict 25th to 75th percentiles, and lines depict the median. Ordinary 1-way ANOVA, Tukey’s multiple comparison test.
We devised an injury model to study regeneration in newborn animals using cryoablation to reduce functional kidney tissue and induce reversible injury adjacent to the necrotic cryoinjured area. Newborn mice subjected to cryoinjury recover and heal the injury site, yet we did not find evidence for the generation of new nephrons; rather, we found hypertrophy of existing nephrons to replace lost tissue. While there was some destruction of ureteric bud tips at the injury site, adjacent buds were intact and presumably retained the capacity to induce new nephron formation. We found no difference in proliferative activity adjacent to, or away from, the injury site, and the weight of the injured organ was lower compared with control animals if the contralateral kidney remained in place. When the contralateral kidney was removed at the time of injury, the hypertrophic response of the injured kidney was enhanced such that the weight was greater

Figure 6. Development of fibrotic lesions after injury. (A) At 3 weeks of age, animals with cryoinjury at P1 showed staining for fibrotic lesions as determined by Sirius red and α-SMA, whereas P7 injured animals exhibited more residual indentation and fibrotic lesions at the injury site. Injury with contralateral nephrectomy resulted in healing without fibrosis if injury was carried out on P1 whereas despite nephrectomy fibrosis was present 3 weeks after injury on P7. Original magnification, ×3. (B) Long-term (12 week) follow-up of kidneys subjected to cryoinjury shows indentation and minimal fibrosis at the site of injury in P1-injured animals, with a similar picture in animals with cryoinjury subjected to contralateral nephrectomy. P7-injured animals with cryoinjury had more indentation with fibrotic tissue, as determined by trichrome staining, and this increase was mitigated if there had been contralateral nephrectomy. n = 7 animals per group. Scale bar: 1.0 mm. SMA, smooth muscle actin.
than sham-treated control kidneys at 21 days when normalized by the body weight of the animal. With contralateral nephrectomy, there was no detectable fibrosis 21 days after injury despite the fact that a large part of the kidney had undergone cryoablative necrosis. Though our model results in injury that is localized and creates space for expansion of new nephrons, we did not observe evidence of de novo nephrogenesis.

In approximately one-third of the injured mice we detected the persistence of single and small clusters of Six2+ cells around the injury area for 2 days longer than in uninjured kidneys. This persistence, however, consisted of single and isolated cells that are unlikely of significance, as they have been separated from an organized niche, and no new nephrons were formed from these dispersed cells. While Six2 activity is required for maintaining the mesenchymal progenitor population in an undifferentiated state (38), Six2 persistence alone is unlikely to be sufficient for ongoing nephrogenesis. For example, expansion of nephrogenic Six2+ progenitors by Lin28 overexpression is not sufficient to induce nephrogenesis (39). The lack of formation of new nephrons raises the question of whether the “nephrogenic zone,” described to be present at birth in rodents, is capable of creating new nephrons or rather represents a region in which already fate-determined nephrons are maturing. Our data suggest that, in the neonatal mice, Six2+ cells at birth contribute to approximately 45% of glomeruli, but the ability to expand this population and produce additional glomeruli is lost and is not enhanced after injury.

Given the long-term adverse outcomes associated with low nephron number at birth, with increased incidence of hypertension and chronic kidney disease (30), it is important to understand the determinants of nephrogenesis with an eye to developing interventions that can enhance nephron number. New nephrons can be generated in response to partial nephrectomy in adult fish (40) but, to date, attempts to do so in adult mammals have been unsuccessful. Importantly, one study using in utero partial renal ablation in an ovine model showed increased nephron numbers in the contralateral kidney after the removal of one kidney (41). The stage of nephrectomy targeted in this study was well before the completion of kidney formation and thus underlines the importance of timing in inducing additional nephron formation. Our work targeted late-stage nephrogenesis in mice. It is not clear why new nephron formation does not continue after P3 or cannot be induced at P1–P3 in mice with an intact “nephrogenic zone.” Gene expression patterns regulating interactions between early nephrons and the ureteric epithelium are altered by P2 (12, 13), and there is strong expression of TMEM100, a proximal tubule marker, in the cap mesenchyme by P1 (42). Rumballe et al. (13) have proposed an accelerated differentiation at P1. These observations, together with our own, support a model in which the cap mesenchyme has already committed to nephron formation by P1.

In conclusion, our data demonstrate that, despite a persistence of a “nephrogenic zone” at birth, newborn mouse kidneys are unable to generate compensatory new nephrons in response to acute kidney mass reduction. In the rodent, the capability for nephrogenesis postnatally in response to nephron reduction appears to be limited to the nephrons that are already at a committed stage of development by the time of birth.

Methods

Cryoinjury surgery. Outbred CD1 wild-type mice were obtained pregnant from the vendor (Charles River), and mice were left to litter naturally. The morning a new litter was observed was considered the first postnatal day and was labeled “P1,” with subsequent days counted as P2 and up. Neonatal pups within 24 hours of birth (P1), a time at which the nephrogenic zone and Six2+ progenitors are present in the renal cortex, were subjected to cryoinjury, as described as follows. Mice were placed inside a latex glove to keep dry and anesthetized by immersion into ice water for 5 minutes, which causes reversible apnea and asystole, thereby preventing excessive blood loss during surgery. The complete procedure was performed on an ice block to prevent rewarming during surgery. Lateral flank incisions were performed to expose the kidneys.

For cryoinjury, the kidney was gently pulled out of the retroperitoneal space with a cotton swap. A cooling probe, consisting of the tip of a blunt steel framing nail placed into a syringe outlet, cooled by liquid nitrogen placed inside the syringe, was placed for 3 seconds on the kidney surface at the middle of the kidney curvature. The cooling probe produced a consistent and reproducible area of injury that was initially white in appearance while cold but quickly returned to the color of the remaining kidney.

For nephrectomy, the kidney was exposed after a flank incision and gently pulled out, and 6.0 silk suture was placed around the kidney. After closure of the suture around the kidney, the parenchyma was cut with scissors to remove the whole kidney parenchyma, leaving the closed suture in place to prevent bleeding.

Following cryoinjury and/or nephrectomy, the flank incisions were sutured with two 6.0 silk closure sutures.
sutures in one layer, and the mice were warmed slowly under a warming lamp. The return of blood circulation was visible by a change in skin color to pink and movement of the animals. Sham operated mice underwent the same procedure, except the kidneys were exposed without cryoinjury. The entire procedure lasted 10–15 minutes depending on the extend of the procedure (sham versus cryoinjury versus cryoinjury plus nephrectomy). After refinement of the technique and minimization of bleeding, more than 90% of animals survived the surgical procedure immediately after anesthesia. Overall survival was lower because of maternal cannibalization of about 10% of pups after placing pups back with the mother. Mortality was similar when carried out in neonates or P7 animals.

The same anesthesia protocol was employed for kidney pole resection. After placement of a suture around the lower pole, microsurgery scissors were applied to the pole of the kidney to remove 10%–15% of the kidney. Kidneys were harvested 5 weeks after resection. The data presented in Figure 2G are from 10 animals pooled from 3 separate experimental dates.

**Lineage-tracing experiments.** Male Six2<sup>GCE/-</sup> mice (Six<sup>2<sub>mEGFP/cre/ERT2+Amc</sub></sup>) crossed with R26R<sup>tDTomato/tDFto- mate</sup> (Gt(ROSA)26Sgr<sup>tm1Wtsi</sup>/GraCAGtdTomato/Rvtr) female mice were used for lineage-tracing experiments. Transgenic mice were obtained from Jackson laboratory and bred and mated in the local animal facility. Induction of Cre recombinase activity and reporter gene expression was achieved by administering a single dose of 0.2 mg (~0.05 mg/g BW) estrogen analog tamoxifen (Sigma-Aldrich) dissolved in corn oil subcutaneously to neonatal mice either immediately after surgery on P1 or, in a subset of animals, at P4 to lineage-trace Six2<sup>+</sup> cells. Mice with the Six2<sup>GCE</sup> allele express a GFP::CreERT2 fusion protein in the Six2 expression domain (27). This protein remains sequestered outside the nucleus until tamoxifen is administered. Upon tamoxifen treatment, the GFP::CreERT2 protein translocates to the nucleus, acting on the loxp sites flanking a STOP sequence upstream of the tdtomato gene insert, resulting in permanent red (tDTomato RFP) labeling of all cells that express Six2 at the time of tamoxifen administration.

For frozen sections, tissues were fixed in 4% paraformaldehyde in PBS for 1 hour at 4°C, dehydrated using 30% sucrose overnight and snap-frozen in OCT. 20-μm-thick sections were prepared using a cryo-tome for immunofluorescence and in situ hybridizations.

**Tissue preparation.** For H&E and trichrome staining, paraffin embedding and sectioning (5 μm) was performed as follows. Kidneys were fixed overnight in 10% formalin at room temperature, moved to 70% ethanol, and stored up to 1 week before paraffin embedding. H&E and Masson's trichrome staining were performed according to standard procedures.

**Immunofluorescence.** Frozen sections (20 μm thickness) were warmed, dried, and washed in PBS/0.1%-Triton (PBT) and blocked for 1 hour with 5% donkey serum. The primary antibody was applied at 4°C overnight at the tested dilution (in PBT and 1% donkey serum). Following 3 washes with PBT, the secondary antibody was applied together, at times, with either Lotus Tetragonolobus Agglutinin (LTL) (Vector labs, B-1325) or Dolichos Biflorus Agglutinin (DBA) (Vector labs, FL-1031) for 2 hours, followed by a brief rinse with PBT and nuclear stain with Hoechst or DAPI for 10 minutes. Mounting was done with Vectashield (Vector labs). Primary antibodies used were as follows: rabbit polyclonal anti-Six2 (Pro-teintech, 11562-1-AP) and Pax2 (Abcam, ab37129).

In situ hybridization. Frozen sections (20 μm thickness) of mouse kidneys were hybridized with digoxigenin-labeled antisense riboprobes generated using gene-specific PCR. The complete protocol for riboprobe generation and section in situ hybridization is available on the GUDMAP web site (http://gudmap.org/) listed under “Protocols” (43, 44). In situ hybridization was performed after thawing and drying sections stored at −80°C. Sections were fixed in 4% PFA for 10 minutes and washed in PBS 3 times. Incubation with 10 μg/ml of proteinase-K was followed by 3 washes in PBS. After proteinase K treatment, samples were fixed again and prehybridized at 70°C for 1 hour before being hybridized with 500 ng/ml digoxigenin-labeled riboprobes at 70°C overnight. Samples were washed with hot solutions at 65°C and treated with 2 μg/ml RNase A for 15 minutes. After additional stringent hot washes, samples were washed with 1×MBST (0.1 M maleic acid, 0.15 M NaCl, 0.1% Tween-20 [pH 7.5]), blocked in blocking solution (20% heat-inactivated sheep serum in MBST plus 2% Boehringer Mannheim Blocking Reagent), and incubated subsequently with anti-digoxigenin antibody conjugated with alkaline phosphatase (Roche, 1:4,000) overnight at 4°C. Extensive washing in 1×MBST was followed by incubation in NTMT (100 mM NaCl, 100 mM Tris-HCl [pH 9.5], 50 mM MgCl<sub>2</sub>, 0.1% Tween-20, 2 mM levamisole). Hybridization signal was visualized by adding BM purple (Roche). Color reactions were observed at different time points and terminated when signals were strong or a background developed.
After development of the colorimetric assay, samples were post-fixed and stored in 80% glycerol/PBS. Images were then captured with a Nikon digital camera (Nikon) attached to a Nikon SMZ1500 stereoscope (Nikon).

**Quantitative analysis of proliferation.** Frozen sections were stained with anti-Ki67 (Abcam) and LTL. Ki67+ cells were counted in ×40 sections taken from 3 different randomly chosen areas in the mid-cortex, both close to and remote from injury site.

**Glomerular counting.** The acid maceration method was used to determine glomerular number. Acid maceration to quantitate glomerular number is a commonly used and readily accessible method for estimating glomerular number, and the advantages are the ease of performance, low cost, and rapidity. Sources of error are potential counting errors and low precision associated with hemocytometer use or nonuniform suspension. More precise methods, such as the common gold standard, the unbiased physical disector/fracti-onator method, are extremely labor and cost intensive, because a whole kidney is sectioned exhaustively. We believe that acid maceration is reasonably accurate in estimating the glomerular number in a whole kidney, and it is commonly used in the literature as a standard method to estimate nephron number (23).

Briefly, harvested kidneys were decapsulated and chopped into small pieces with a razor blade and then suspended in 1 ml of 6 N HCl at 37°C for 1 hour, followed by trituration by micropipette. Five milliliters of distilled water were added to a total volume of 6 ml, which was incubated overnight at 4°C. Pictures of 10-μl sample volumes on a Neubauer chamber with grid (without using a coverslip) were taken on a Nikon SMZ1500 stereoscope. For each sample, a total of 10 pictures were taken for counting, so that a volume of 100 μl was counted. Manual counting was done blinded to the sample using ImageJ (NIH) (45).

**Statistics.** Statistical analyses were performed using GraphPad Prism 5 software. Significance was tested using the 2-tailed unpaired Student’s t test or 1-way ANOVA with Dunnett or Tukey analysis for multiple comparisons when appropriate. A P value of less than 0.05 was considered statistically significant. All values are expressed as mean ± SD.

**Study approval.** All protocols were approved by the Institutional Animal Care and Use Committee of the Harvard Medical Area Standing Committee on Animals, Harvard Medical School.

**Author contributions**

JVB, FT, BSF, and RI conceived the study. JVB, FT, RI, BSF, MTV, and MG provided study methodology. FT, RI, BSF, and MTV performed experiments. JVB, MTV, and MG provided resources. JVB acquired study funding. FT and MTV wrote the original draft of the manuscript. FT, BSF, MTV, and JVB reviewed and edited the manuscript.

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