Functional megalin is expressed in renal cysts in a mouse model of adult polycystic kidney disease

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ABSTRACT

Background. Autosomal dominant polycystic kidney disease (ADPKD) is characterized by the progressive growth of cysts and a decline of renal function. The clinical feasibility of the number of potential disease-modifying drugs is limited by systemic adverse effects. We hypothesize that megalin, a multiligand endocytic receptor expressed in the proximal tubule, may be used to facilitate drug uptake into cysts, thereby allowing for greater efficacy and fewer side effects.

Methods. The cyst expression of various tubular markers, including megalin and aquaporin 2 (AQP2), was analysed by immunohistochemistry (IHC) of kidney sections from the ADPKD mouse model (PKD1 RC/RC) at different post-natal ages. The endocytic function of megalin in cysts was examined by IHC of kidney tissue from mice injected with the megalin ligand aprotinin.

Results. Cyst lining epithelial cells expressing megalin were observed at all ages; however, the proportion decreased with age. Concomitantly, an increasing proportion of cysts revealed expression of AQP2, partial expression of megalin and/or AQP2 or no expression of the examined markers. Endocytic uptake of aprotinin was evident in megalin-positive cysts, but only in those that remained connected to the renal tubular system.

Conclusions. Megalin-expressing cysts were observed at all ages, but the proportion decreased with age, possibly due to a switch in tubular origin, a merging of cysts of different tubular origin and/or a change in the expression pattern of cyst lining cells. Megalin expressed in cysts was functional, suggesting that megalin-mediated endocytosis is a potential mechanism for drug targeting in ADPKD if initiated early in the disease.
INTRODUCTION

Autosomal dominant polycystic kidney disease (ADPKD) is the most common inherited kidney disorder, with a reported prevalence of 1/1000–1/2000 in the general population [1, 2]. ADPKD is genetically heterogeneous and is mainly associated with pathogenic variants in two genes, PKD1 and PKD2, which encode the membrane proteins polycystin 1 (PC1) and polycystin 2 (PC2), respectively [3, 4]. The disease is characterized by progressive growth of kidney cysts, causing a decline in renal function and leading to end-stage renal disease (ESRD) [5] in ~50% of patients by the sixth decade of their life [6, 7]. While present-day interventions may slow disease progression, there is currently no available treatment that efficiently blocks the progression of cyst formation and growth, in part because the dosing is limited by systemic adverse effects resulting from targeting of other organs [8]. Multiple studies in rodent models of ADPKD have shown that in the early stage of disease, cyst formation mainly initiates from the proximal tubules (PTs), after which a tubular switch in cyst origin is observed [9–11]. In the kidney, megalin is selectively expressed within the brush border located at the apical membrane of PT cells, where it serves as a multiligand receptor essential to reabsorption of a large number of filtered substances, including drugs [12]. Megalin has previously been explored as a kidney-specific drug-targeting receptor in which the conjugate was efficiently endocytosed and accumulated exclusively in the PT cells [13]. Based on these findings, we hypothesized that megalin may be used to facilitate drug uptake in megalin-expressing cyst lining epithelial cells (referred to as cystic cells hereafter), thereby increasing efficacy and reducing adverse effects. To substantiate the applicability of this approach, we examined the temporal expression and function of megalin in cystic kidney tissue from a PKD1RC/RC mouse model of ADPKD.

MATERIALS AND METHODS

Ethics statement

Animals used in all experiments were maintained and subjected to procedures according to the guidelines and approval of the Danish Animal Experiment Inspectorate (2019-15-0201-01631).

Animal model

The ADPKD mouse model PKD1RC/RC has previously been published [9]. The mice were housed in a 12-h day/night cycle, a temperature of 21 °C and bred homozygous in the animal facility at the Department of Biomedicine, Aarhus University. Ten mice (five of each sex) were anaesthetized by isoflurane inhalation at the post-natal (P) age 10, 20, 30, 90 and 180 days and, when possible, kidneys were perfusion fixed through the aorta (or otherwise immersion fixed) with 2% paraformaldehyde in 0.1 M sodium cacodylate buffer, pH 7.4. Kidneys were subsequently dehydrated and embedded in paraffin by standard methods. Four micrometer sections were cut from the centre of the kidney and placed on SuperfrostPlus object glass slides (Thermo Fisher Scientific, Waltham, MA, USA). The sections were stained for immunohistochemistry as described previously [16] using IP and IF labelling of segment-specific markers: megalin (PT), aquaporin 1 (AQP1; PT, thin descending limb of Henle’s loop, vasa recta), sodium–chloride cotransporter (NCC; early distal convoluted tubule), calbindin (late distal convoluted tubule) and aquaporin 2 (AQP2; connecting tubule and collecting duct, collectively referred to as CD). Images were acquired using a motorized, upright widefield slide scanner (Olympus BX61VSU, Olympus Life Science, equipped with an ORCA-Flash 4.0 camera (Hamamatsu Photonics, Hamamatsu City, Japan). Whole slide scans were obtained with a ×40 objective. Image processing and measurements were performed using cellSens Entry version 2.3 software (Olympus Life Science) and basic image acquisition was achieved using OlyVIA version 2.9 software (Olympus Life Science).

IP staining

Primary antibodies Included sheep anti-megalin (1:20 000; gift from Pierre Verrout, France), rabbit anti-AQP1 (1:2000; Merck, Darmstadt, Germany), rabbit anti-NCC (1:1000; gift from Mark A. Knepper, USA), mouse anti-calbindin (1:5000; gift from Fitzgerald, UK) and rabbit anti-AQP2 (1:2000; Merck).

Secondary antibodies included polyclonal rabbit anti-sheep immunoglobulin G (IgG; Dako, Glostrup, Denmark), polyclonal goat anti-rabbit IgG (Dako) and polyclonal goat anti-mouse IgG (Dako). All were diluted to a concentration of 1:200. Horseradish peroxidase–conjugated antibodies were activated by incubation with 3,3-diaminobenzidine (PanReac AppliChem ITW Reagents, Chicago, IL, USA) for 10 min. The sections were counterstained with Mayer’s haematoxylin (Ampliqon, Odense, Denmark) for 2 min and dehydrated before mounting of coverslips with Eukitt (PanReac AppliChem ITW Reagents).
IF staining

Primary antibodies included sheep anti-megalin (1:10,000; gift from Pierre Verroust, France), rabbit anti-AQP2 (1:1000; Merck) and rabbit anti-aprotinin (1:2000; MyBioSource, San Diego, CA, USA).

Secondary antibodies included Alexa Fluor546 donkey anti-sheep IgG (Invitrogen, Carlsbad, CA, USA) and Alexa Fluor488 donkey anti-rabbit IgG (Invitrogen). All were diluted to a concentration of 1:300. Sections were mounted with coverslips using Fluoroshield mounting medium (Sigma-Aldrich).

Serial sections

From each mouse injected with aprotinin, 200 serial sections were cut from the centre of one kidney. Every second slice was IP labelled with anti-aprotinin (1:2000; MyBioSource) to identify any possible connection between aprotinin-positive cysts and the tubular system and every 10th slide was IF labelled with anti-megalin and anti-AQP2 to establish the tubular expression pattern of the cysts. Cysts were traced through the consecutive sections using OlyVIA version 2.9 software (Olympus Life Science).

Statistical analysis

All quantitative data are presented as mean ± SD. Two-way analysis of variance (ANOVA) was used to compare mean cyst count between the groups (sex and age) and Student’s t-test was used to assess the differences in mean cyst count between sex within each age group. P-values < 0.05 were considered statistically significant.

RESULTS

The number of renal cysts in PKD1RC/RC mice increases with age

Renal cysts were observed at all ages and were often located in clusters rather than scattered (Supplementary data, Figure S1). The total number of cysts increased from 22.3 ± 4.54 at P10 to 91.1 ± 15.72 at P180 (Figure 1; Supplementary data, Table S2), showing a statistically significant, positive correlation between the number of cysts and age (P = 0.0000001, two-way ANOVA). A trend towards more cysts in male mice was observed (Figure 1), but this was not statistically significant (P = 0.8), supporting an age-dependent increase in the number of cysts independent of sex. Five mice at age P90 were excluded due to an extreme outlying phenotype (excessive total kidney volume and number of cysts) when compared with the rest of the animals at the same age (Supplementary data, Table S3). These mice were born following embryo transfer and the phenotype was most likely related to this procedure. Data obtained from these animals is therefore not part of the presented results in the following sections.

PKD1RC/RC mice show a switch in tubular marker expression of cysts with age

As cysts develop from different and specific tubular segments, we wanted to describe tubular marker expression in cystic cells, focusing on identifying cysts expressing megalin. As expected, all tubular markers (megalin, AQP1, NCC, calbindin and AQP2) stained the tubule segments in which they are normally expressed (Supplementary data, Figure S2). Expression of all markers, except for NCC, was observed in cysts at all ages. Megalin and AQP1 are co-expressed in the apical membranes of the PT and a similar co-expression was observed in cysts with very few exceptions (data not shown). Also, calbindin and AQP2 are both expressed in connecting tubules, hence most calbindin-positive cysts also expressed AQP2. Kidney sections from all mice were stained with all tubular markers, but in the subsequent data analysis we focused on megalin and AQP2, as they were the most commonly expressed of the analysed membrane proteins in cystic cells and no colocalization of these two markers was observed.

Megalin was expressed in the brush border and endocytic apparatus in cystic cells at all ages (Figure 2A and B). At P10, 56% of the cysts were megalin positive, which decreased to 45% at P20, 25% at P30, 21% at P90 and 16% at P180 (Figure 3). Concomitantly, an increase with age in cysts expressing AQP2 was observed (Figure 3). This switch in tubular marker expression was also noticed when looking at cyst occupancy, with megalin-positive cysts constituting the largest area of the kidney in younger mice and AQP2-positive cyst occupying the largest area in older mice, when comparing the two markers (Figure 4). Interestingly, the fraction of cysts that did not stain for any of the examined tubular markers increased with age (14% at P10 versus 44% at P180) (Figures 3 and 5C). In addition, some of the larger cysts at P20, P30, P90 and P180 seemed to partially lose their expression of megalin, with some parts of the cystic cells being megalin positive, while others appeared negative (Figure 2A [P90 red *] and Figure 5A). Similar results were observed in cysts expressing AQP2 at P90 and P180 (Figure 5B). Some cystic cells at all ages revealed an altered morphology, with flattened and elongated cells, when compared with the surrounding tubules. The change in cell morphology was independent of megalin and AQP2 expression but was associated with the loss of expression of both markers (Supplementary data, Figure S3).

Megalin mediates endocytosis of aprotinin in cystic cells

To establish if megalin expressed in cystic cells is capable of mediating endocytosis, we injected PKD1RC/RC mice at P25 and
P90 with aprotinin and visualized potential uptake using anti-aprotinin antibody. Aprotinin co-localized with megalin in PTs and cysts. Some megalin-positive cysts with no apparent aprotinin uptake were also observed, demonstrating that the expression of megalin is not sufficient for the uptake into cystic cells (Figure 6A). Diffuse, intracellular positive staining was also observed in some AQP2-positive segments, which may reflect a non-specific mechanism of aprotinin uptake. Megalin-positive cystic cells and PTs revealed granular staining for aprotinin consistent with endocytosis, suggesting that megalin in cystic cells is capable of mediating endocytosis (Figure 6B and C).

Aprotinin uptake into cystic cells depends on a tubule–cyst connection

To determine if a connection between cysts and the tubular system was required for aprotinin uptake, 200 serial sections were prepared from the PKD1RC/RC mouse injected with aprotinin (Figure 7). All aprotinin-positive cysts were at some point attached to an adjacent tubule segment and aprotinin was absent in megalin-positive cysts with no tubular connection. Hence a persistent connection to the tubular system is necessary for the efficient uptake of injected and filtered aprotinin into megalin-positive cyst lining cells. No significant difference in the proportion of megalin- and aprotinin-positive cysts was observed when comparing mice at P25 (~62% (33–100%)) with mice at P90 (~67% (50–85%)) (P = 0.79) (Supplementary data, Table S4). During the tracing of a mouse at P25, we observed a possible fusion of two separately defined cysts, both attached to the tubular system (Supplementary data, Figure S4A). One part of the cyst showed evidence of aprotinin uptake, whereas the other part was aprotinin negative, suggesting that the cysts may originate from two different tubule segments. Based on morphology and location of a nearby GL, the tubule segment attached to the cyst part positive for aprotinin was most likely a PT segment, whereas the segment without aprotinin uptake could not be identified (Supplementary data, Figure S4B). Sections prior to the tubule–cyst connection were not available and it was therefore not possible to further characterize the tubular marker expression pattern of this tubule segment. Additional immunolabelling showed that the ‘fused’ cyst was megalin positive, at least in some parts (Supplementary data, Figure S4C).

DISCUSSION

In this hypothesis-generating study, we demonstrate that megalin is expressed in cystic cells and megalin is functional and capable of endocytosing ligands freely filtered in the glomeruli if an intact tubule–cyst connection is present. The PKD1RC/RC mouse model used is characterized by progressive formation of renal cysts over time. The cystic cells selectively express markers specific to various segments of the renal tubular system consistent with previous observations in other ADPKD models [10, 11]. Our observations suggest that
cysts originate from both PT and CD segments and a switch in cyst origin occurs, with more cysts being of CD origin with age.

The fractions of cysts with partial or absent expression of any of the examined tubular markers increase with age as disease progresses. This may reflect dedifferentiation of cystic cells characterized by a loss of apical–basal polarity [17], which may also explain the changed morphology of cystic cells, which become flatter and more elongated [18]. Consistent with this, similar results have been observed in other ADPKD mouse models [19]. Thus partial expression of megalin or AQP2 may reflect dedifferentiation that eventually results in loss of expression. Interestingly, the number of cysts originating from PT remains quite stable over time, suggesting dedifferentiation is mainly affecting cysts originating from other tubules. As an alternative to dedifferentiation, the fusion of cysts originating from different tubular segments may lead to partial expression of particular markers. We identified one case of what appears to be two fused cysts clearly attached to different tubular segments. We did not identify any cysts expressing both megalin and AQP2, which may have been expected if cyst fusion was common. Thus it remains unclear if this observation represents a fusion of two cysts or some other process.

Injected aprotinin was internalized by PT cells and megalin-expressing cystic cells, suggesting megalin-mediated endocytosis; however, an intact tubule–cyst connection was required, consistent with the notion that such ligands must be filtered in the GL prior to endocytosis [20]. It is generally accepted that cysts develop from the tubular system and eventually pinch off as they become enlarged [21]. In the PKD1<sup>RC/RC</sup> mouse model, we observed cysts both at P25 and P90 with a diameter >300 µm that remained connected to the tubular system. The ratio between cysts positive for both megalin and aprotinin (suggesting an intact tubule–cyst connection) and cysts positive for megalin only (suggesting detachment of cysts from the tubular system) is similar at P25 and P90, implying that the fraction of megalin-positive cysts connected to the tubular system does not decrease with time, at least not until after P90. In humans, the diameter of cysts before detaching has been estimated to 2–3 mm [22], and it has been reported that 20–30% of examined cysts at ESRD remain attached to the tubular system [23].

Megalin-mediated endocytosis has previously been explored experimentally in mice for kidney-specific drug delivery using peptide-based ligands, showing peptides accumulating almost exclusively within the kidneys [13, 24], although megalin is also expressed in other absorptive epithelia [12]. While the above findings may suggest a similar capability of megalin to facilitate uptake of ADPKD-specific agents in renal cysts, our findings...
have some important limitations that must be considered. First, manual counting of cysts provides a source of error, as it is a subjective assessment depending on the definition of cysts. We believe that the relatively large number of animals included in the study provides some assurance against random variation and that the relative changes in cyst number and tubular marker expression observed are reliable. Second, all observations are cross-sectional in time, rendering it impossible to make any definite conclusions about the time-space dynamics of cyst formation in the kidney, including cyst origin. However, the use of segment-specific marker expression to characterize cyst origin has been applied in other studies with similar findings [9–11]. Third, the results in rodent models of ADPKD are not always directly transferable to humans [8]. Even though the PKD1RC/RC mouse model used in this study is based on a PKD1 allele (p.R3277C) identified in humans with ADPKD and the model would compare to a consanguineous family with adult-onset ADPKD [9], some differences do exist. In humans, significant gender variability [25, 26], as well as inter- and intrafamilial variability are observed [27], while PKD1RC/RC mice of the same age reveal histological phenotypes with very little variation, most likely related to the very homogeneous background genotypes and similar environmental conditions typical to the animal experimental setting. Despite such potential differences, a number of observations in this study are similar to findings in humans, including the increase in cyst number [23, 28] as well as the increase in AQP2-positive or marker-negative cysts with increasing age [23]. In addition, the original article states that reduced renal function was observed in mice from 9 months of age [9], which suggests the observations in this study compare to a time before glomerular filtration rate starts to decline in humans. However, the capability of megalin to facilitate uptake of ADPKD-specific agents in renal cysts was only examined in four mice, hence the results need to be verified in other species, including humans.

In conclusion, in a mouse model of ADPKD, megalin is expressed in renal cysts at ages from P10 to P180, although the proportion of positive cysts decreases with age. The megalin ligand aprotinin is internalized in megalin-positive cystic cells if an open tubule–cyst connection can be preserved. Thus potential treatment approaches using megalin to facilitate uptake of ADPKD-modifying agents probably should be applied in the early stages of the disease and should exert an ADPKD-specific effect that does not affect other megalin-expressing cells and tissues [16].

![Figure 6: Aprotinin is endocytosed into megalin-positive cysts. Representative images showing the expression of megalin and the localization of injected aprotinin by IF and IP labelling in kidney sections from a PKD1RC/RC mouse at P25. (A) A megalin- (red) and aprotinin-positive (green) cyst (top row), as well as a megalin-positive, but aprotinin-negative cyst (bottom row). Scale bar: 100 µm and 50 µm, respectively. (B) Megalin- and AQP2-positive cysts (IF, left) compared with aprotinin staining by IP (right) of an adjacent section. Scale bar: 50 µm. (C) Magnification of (B). Scale bar: 20 µm. In the megalin-positive cysts as well as neighbouring PTs, aprotinin labelling is clearly granular and vesicular, reflecting endocytosis, whereas aprotinin labelling of the AQP2-positive cysts is diffuse, possibly reflecting non-specific uptake. Asterisk: megalin-positive cyst; arrow: proximal tubule.](#)
SUPPLEMENTARY DATA
Supplementary data are available at ckj online.

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CONFLICT OF INTEREST STATEMENT
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DATA AVAILABILITY STATEMENT
The data underlying this article are available in the article and in its supplementary material.

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