The mTOR Pathway Is Activated in Human Autosomal-Recessive Polycystic Kidney Disease

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Key Words
Cystic kidney disease · Autosomal-recessive polycystic kidney disease · mTOR

Abstract
Background: An inappropriate activation of the mTOR pathway was demonstrated in the autosomal dominant (AD) form of polycystic kidney disease (PKD). To date it is unclear whether the mTOR pathway is activated in autosomal-recessive (AR) PKD, a cystic disease which occurs in childhood. The purpose of the present study was to evaluate the mTOR pathway in AR PKD.

Methods: We evaluated the expression of mTOR pathway molecules in paraffin-embedded liver and kidney samples from patients with AR PKD and control specimens from animals as well as humans. Monoclonal antibodies, the phosphorylated proteins pmTOR, pS6-ribosomal-protein (pS6K), p4E-BP1, pelf4G, and phospho-tuberin/TSC2 were used.

Results: mTOR was strongly expressed in renal cyst-lining cells and bile ducts from AR PKD specimen. S6K immunostaining was strong in smaller tubules and weak both in larger renal cysts and in the bile duct epithelium. In controls, mTOR and S6K were expressed in distal tubule segments. 4E-BP1-immunostaining was restricted to noncystic tubules in AR PKD. eIF4G-immunostaining was observed in bile duct epithelium in AR PKD, but not in control tissue. Tuberin/TSC2 immunostaining was negative in all specimens.

Conclusion: Our data suggest that the mTOR pathway may be activated in AR PKD, and mTOR molecules may represent a potential target to slow down cyst development in this disease.

Introduction
Polycystic kidney diseases (PKD) include various disorders characterized by cystic kidneys and variable multiorgan pathology [1]. Autosomal-recessive (AR) PKD is a form of childhood PKD which is characterized by rapidly progressive cyst development and congenital hepatic fibrosis (CHF). CHF is characterized by bridging hepatic fibrosis and proliferation of irregularly shaped small in-
trahepatic bile ducts. The latter can be considered the hepatic equivalent to renal cyst development.

Advances in the understanding of the molecular basis for PKD have led to promising new therapeutic approaches to slow down progression of cyst development [2]. One such strategy includes inhibition of the mammalian target of rapamycin (mTOR), a conserved Ser/Thr kinase that regulates various cellular processes including growth and metabolism [3, 4]. Recently, an activation of the mTOR pathway has been described in cystic tubules in mouse models and in the human autosomal dominant (AD) form of PKD [5].

mTOR is a member of the phosphoinositide kinase-related kinase (PIKK) family that phosphorylates proteins on serine or threonine residues [6]. Target molecules of mTOR include ribosomal S6 kinase and eukaryote initiation factor 4E binding protein 1, key translation regulators for protein synthesis. mTOR activation is regulated by a variety of signaling pathways including small GTPases, Rheb, akt, Raptor, and the TSC1-TSC2 complex [3]. Recently, Wahl et al. [7] demonstrated an enhanced phosphorylation of akt, an upstream regulator of mTOR, in Han:SPRD rats.

To date, several experimental studies in rodent models of PKD have revealed that inhibition of the mTOR pathway results in reduction of kidney size, prevents the loss of kidney function, and lowers cyst volume [5, 8–10]. Furthermore, retrospective observations from patients with PKD and end-stage renal disease who underwent kidney transplantation have shown that treatment with sirolimus, an mTOR inhibitor, reduced kidney volumes by 25% [5]. Currently, several clinical trials have started to investigate the effect of mTOR inhibitors in patients with AD PKD [2].

Fischer et al. [11] have reported an activation of the mTOR pathway in kidneys from patients with AR PKD. In the present study, we investigated the mTOR pathway in kidney and liver specimens from patients with AR PKD. Furthermore, we evaluated the physiological renal and hepatic expression of these molecules in animals as well as humans. Shillingford et al. [5] reported an inappropriate activation of the mTOR pathway in the autosomal dominant form of PKD using antibodies against phosphorylated, active forms of mTOR (pmTOR) and its downstream effector S6 kinase (pS6K). In addition, mTOR-inhibition has been shown to slow down cyst formation in several rodent models of PKD [12]. In our study, we have focused our analysis on phosphorylated forms of the mTOR-related molecules. We analyzed expression of the upstream mTOR-regulator tuberin, and the mTOR effectors S6K, eIF4G, and 4E-BP1.

Materials and Methods

Reagents, Antibodies, and Animals

Rabbit-monomonal antibodies against the following phosphorylated, activated forms of kinases of the mTOR pathway were purchased from Cell Signaling Technology, Frankfurt am Main, Germany: pmTOR (Ser2448), pS6K (Ser23/24), p4E-BP1 (Thr37/46), pElF4G (Ser1108), and p-tuberin/TSC2 (Thr1462). All other chemicals and reagents were supplied by Sigma. BALB/c mice and Sprague-Dawley rats (Charles River, Sulzfeld, Germany) were used for experiments.

Specimen

Human liver and kidney specimens from patients with AR PKD were obtained from subjects who underwent either combined kidney – liver transplantation or received grafts consecutively. Human control kidneys were obtained from patients who underwent complete unilateral nephrectomy or hepatectomy due to carcinoma. Sections were taken from regions morphologically free from any signs of cancer. Rodent kidneys were fixed by perfusion with 4% paraformaldehyde in PBS, and subsequently embedded in paraffin [13]. The study was approved by the local Ethics Committee of the University Hospital Essen.

Immunostaining

Immunostaining was carried out on 4-μm-thick paraffin-wax sections. Antigen retrieval was carried out with 0.01 M citrate buffer at pH 6.0 for 25 min in a hot water bath (95 °C). Primary antibodies were incubated for 60 min and antibodies were demonstrated with a commercially available antibody anti-rabbit IgG detection kit (ZytoChem-Plus HRP Polymer-Kit, Zytomed Systems, Berlin, Germany). Sections were counterstained with Mayer’s hemalum solution (Merck, Darmstadt, Germany). Sections were coverslipped and examined with a Zeiss Axio Imager equipped with an AxioCam MRC Camera (Carl Zeiss Jena GmbH, Jena, Germany).

Patients

Table 1 displays the characteristics of the cohort. Mutation screening was done for the 66 exons encoding the 4074 aa polyductin protein (GenBank NM_138694) by denaturing high-performance liquid chromatography (DHPLC) on a Wave Fragment Analysis System (Transgenomic, Crewe, UK), as recently described in detail [14].

Results

Patients were between 4 and 14 years old when they received a liver transplant (table 1). In 2 patients, two mutations in the PKHD1 gene were detected; in another 2 patients only one mutation in the PKHD1 gene was detected, and in another 2 patients no mutation in the PKHD1 gene was found. All detected missense mutations affect evolutionarily conserved residues of the polyductin/fibrocystin protein and were absent among 200 ethnically matched controls. Specimens from all 6
patients were analyzed; the presented specimens are denoted in Table 1, the specimens studied are shown in Table 1.

Expression of pmTOR
Antibodies against phosphorylated molecules were used to evaluate the activation of the mTOR pathway.

Figure 1 shows the distribution of pmTOR in human kidneys with AR PKD and human and murine control kidneys. In AR PKD kidneys, pmTOR was strongly expressed in cyst lining cells whereas there was no expression or very low levels of expression detectable in non-cystic tubules with the exception of a few tubule cells. The subcellular staining pattern was characterized by an intracellular granular distribution with varying intensity between tubule cells. In control kidneys, there was a clearly positive signal in distal tubules and collecting ducts, whereas proximal tubuli and glomeruli were negative. This staining pattern was observed in human (Fig. 1e, f), murine (Fig. 1g, h) and rat kidneys (data not shown).

Figure 2 shows the expression of pmTOR in human liver specimen from patients with and without AR PKD. Three liver explants from patients with AR PKD are shown to illustrate the variety of the bile duct morphology in AR PKD ranging from few massively enlarged and elongated bile ducts (Fig. 2a, b) to small and irregularly shaped bile ducts (Fig. 2e, f). pmTOR immunostaining was strongly positive in all bile ducts in patients with AR PKD. Bile duct cells showed a strong ubiquitous intracytoplasmic staining pattern. In contrast, bile ducts from human control livers showed only a weak immunostaining for pmTOR restricted to the apical cell border (Fig. 2g, h).

Distribution of Phosphorylated Ribosomal Protein S6 (pS6K)
Figure 3 shows the distribution of pS6K in human kidneys with AR PKD and control kidneys from humans and mice.

The immunostaining showed a heterogeneous distribution with strong positive staining in smaller tubules compared to the pmTOR staining in AR PKD kidneys. Larger cysts exhibited only a weak positive signal for pS6K, in some cystic tubules the signal was even completely absent. Positive staining appeared to involve all cells from one affected tubule. In control kidneys, pS6K immunostaining was restricted to the thin ascending part of the loop of Henle. In addition, there was a weaker staining signal of distal tubules in human (Fig. 3e, f) and murine kidneys (Fig. 3g, h; data not shown for rats).

Figure 4 shows the expression of pS6K in human liver specimen from patients with and without AR PKD. Samples from three AR PKD livers are shown in Figure 4a–f. pS6K immunostaining was most intense in parenchymal hepatic cells. Bile ducts also exhibited a positive but weaker signal for pS6K labeling. In contrast, bile ducts from controls were completely negative for pS6K (Fig. 4g, h).

Distribution of p4E-BP1 and peIF4G
Figure 5 shows the distribution of p4E-BP1 and peIF4G in AR PKD kidneys, livers and controls. p4E-

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**Table 1. Characteristics of study cohort**

<table>
<thead>
<tr>
<th>Patient No.</th>
<th>Sex</th>
<th>Age at transplantation</th>
<th>Age at diagnosis, years</th>
<th>Age at ESRF</th>
<th>PKHD1 mutations</th>
<th>Figure</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>M</td>
<td>combined kidney-liver transplantation at 12 years</td>
<td>1</td>
<td>7</td>
<td>c.4457C&gt;T (p.P1486L) (M) c.9464A&gt;G (p.Y3155C) (P)</td>
<td>2c, d</td>
</tr>
<tr>
<td>2</td>
<td>M</td>
<td>combined kidney-liver transplantation at 7 years</td>
<td>2</td>
<td>7</td>
<td>c.107C&gt;T (p.T36M) (M) c.10219C&gt;T (p.Q3407X) (P)</td>
<td>1a, 1b, 3c, 3d</td>
</tr>
<tr>
<td>3</td>
<td>F</td>
<td>combined kidney-liver transplantation at 4 years</td>
<td>1</td>
<td>4</td>
<td>no mutation detected</td>
<td>3a, 3b</td>
</tr>
<tr>
<td>4</td>
<td>M</td>
<td>kidney transplantation with 5 years combined kidney-liver transplantation at 6 years</td>
<td>1</td>
<td>4</td>
<td>c.5895dupA (p.L1966fs)</td>
<td>1c, 1d, 2a, 2b, 4e, 4f</td>
</tr>
<tr>
<td>5</td>
<td>M</td>
<td>combined kidney-liver transplantation at 13 years</td>
<td>7</td>
<td>13</td>
<td>no mutation detected</td>
<td>2e, f</td>
</tr>
<tr>
<td>6</td>
<td>F</td>
<td>combined kidney-liver transplantation at 14 years</td>
<td>8</td>
<td>12</td>
<td>c.9767C&gt;T (p.P3256L)</td>
<td>4c, 4d</td>
</tr>
</tbody>
</table>

P = Paternally inherited allele; M = maternally inherited allele; ESRF = end-stage renal failure.
BP1 immunostaining was restricted to noncystic tubules in AR PKD, but there was no positive signal in cyst lining tubule cells. In the control kidney, few (distal convoluted) tubules were slightly positive (fig. 5c, d). In the liver, no p4E-BP1 labeling could be detected, neither in AR PKD (fig. 5e, f) nor in controls (data not shown).

AR PKD kidneys and control kidneys were negative for peIF4G immunostaining (data not shown). In contrast, in specimens from AR PKD livers, peIF4G immunostaining was observed in the bile duct epithelium (fig. 5g, h), whereas peIF4G immunostaining in control liver was negative. p-Tuberin/TSC2 immunostaining was negative in all kidney and liver specimens.

Fig. 1. Renal expression of activated mTOR. The left panel shows lower magnifications at 10×, the right higher magnifications at 40× of the same specimen. a–d pmTOR expression in 2 different human AR PKD kidneys. Strong labeling is observed in epithelial cyst lining cells whereas non-cystic tubules (proximal tubule, PT) were negative. e, f pmTOR expression in a human control kidney. g, h pmTOR expression in a murine kidney. In both, pmTOR expression was clearly detectable in distal tubules (DT), the convoluted segments of distal tubules (DCT), and collecting ducts (CD), whereas proximal tubules (PT) and glomeruli (GL) were negative. The same immunostaining pattern was observed in murine kidney sections (g, h).
To preclude an impact of age on the expression profile of the analyzed molecules, experiments were performed using specimens from juvenile human controls (8–13 years) and murine pups. There was no difference in the immunostaining pattern between young and adult controls.

### Discussion

In the present study, we evaluated whether the mTOR pathway is activated in human AR PKD. We found that molecules of the mTOR pathway were strongly expressed in cyst lining renal epithelial cells and the bile duct epithelium.
These data suggest that mTOR pathway activity may be related to the cystic changes found in AR PKD. It is possible that mTOR pathway may participate in the cyst development in renal tubules and bile ducts in patients with AR PKD and in the bpk mouse, a murine model of early-onset AR PKD, mTOR inhibition significantly improved the cystic phenotype [5]. In the current study, we provide additional evidence for an activation of the mTOR pathway in human AR PKD. mTOR inhibitors are already used in children after solid organ transplantation, and may also represent a suitable therapeutic target to slow down cyst progression in human AR PKD.

We observed different labeling patterns for pmTOR and pS6K immunostaining in AR PKD kidneys. pS6K is
not only regulated by mTOR sensing amino acid sufficiency but is also located in the signaling pathway of growth factors including PDK1 and PKB, two protein kinases that are client proteins of Hsp90 [15]. It may be possible that such complex control mechanisms may account for the different staining pattern observed for pmTOR and pS6K in AR PKD kidneys.

When we analyzed the physiological expression of mTOR molecules in the human kidney, we found strong expression of the phosphorylated molecules pmTOR and pS6K in distal tubule segments. While Shillingford et al. [5] did not show immunostaining of these molecules in the normal human kidney, another group reported renal expression of mTOR-related molecules in
the normal kidney. Consistent with our findings, Ken- 

erson et al. [16] report renal expression of mTOR-related 

molecules (e.g. pS6K) in distal tubules. To further eluci-

date the renal expression of mTOR molecules, we im-

munostained murine and rat kidneys for these mole-

cules, and found consistent results. The physiological 

significance of expression of these molecules remains 

currently unclear. However, their constitutive expres-

sion in distal tubules and collecting ducts may contrib-

ute to the susceptibility for cyst development, which 

commences in the distal parts of the nephron segment 

in autosomal recessive forms of human and experimen-

tal PKD in rodents [1, 17–19].

Fig. 5. Renal and hepatic expression of 
p4E-BP1 and peIF4G. a, c, e, g Lower mag-
nifications at 10×. b, d, f, h Higher magni-
fications at 40× of the same specimens. 
p4E-BP1 expression in a human AR PKD 

kidney (a, b), a human control kidney (c, 
d), and in a human AR PKD liver (e, f). 
g, h peIF4G expression in an AR PKD (hu-
man) liver.
Liver specimen from AR PKD patients showed a marked granular intense staining for pmTOR molecules in all bile duct cells suggesting that this pathway is activated in the entire bile duct epithelium. This is in line with the observation in the present study that bile ducts in the same AR PKD specimen showed very similar dimensions and morphology. Our observation that all bile ducts are affected in AR PKD is consistent with the clinical course of this biliary disease which almost obligatory manifests as patients age [20].

However, while bile duct morphology showed only little variability in the same specimen, there was a great variability of bile duct morphology between specimens (fig. 2, 4). This is in line with the wide array of clinical phenotypes and disease progression in AR PKD which result from various factors including genetic heterogeneity, modifying and environmental factors [20].

In noncystic tubules in AR PKD kidneys, pmTOR staining was weakly positive in some tubule cells (fig. 1b), suggesting that activation of the mTOR pathway in AR PKD kidneys may successively initiate. Furthermore, pS6K staining in the AR PKD kidney was observed in smaller noncystic tubules. This is consistent with the data provided by Shillingford et al. [5] showing also pS6K expression in smaller tubules. Similarly, p4E-BP1 staining was positive in noncystic tubules in AR PKD kidneys, but no expression was detected in the cystic epithelium. Indeed, after activation and recruitment of mTOR to the translation preinitiation complex (where it phosphorylates S6K and 4E-BP1), 4E-BP1 and S6K dissociate from this multi-molecule complex [21]. It is possible that this accounts for the differential detection of pmTOR, pS6K, and p4E-BP1 in AR PKD kidneys.

We did not find a positive signal for p-tuberin/TSC2 in AR PKD or control kidney or liver specimens. However, activation of these molecules inhibits mTOR activity, and no information concerning the involvement of these upstream pathway molecules could be obtained.

In conclusion, our data indicate that the mTOR pathway is also activated in AR PKD, and that mTOR molecules may thus represent a potential target to slow down cyst development in AR PKD.

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References


