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Inhibition of glucosylceramide accumulation results in effective blockade of polycystic kidney disease in mouse models

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Polycystic kidney disease (PKD) represents a family of genetic disorders characterized by renal cystic growth and progression to kidney failure¹. No treatment is currently available for people with PKD, although possible therapeutic interventions are emerging^{2, 3, 4, 5, 6, 7, 8}. Despite genetic and clinical heterogeneity, PKDs have in common defects of cystic epithelia, including increased proliferation, apoptosis and activation of growth regulatory pathways¹. Sphingolipids and glycosphingolipids are emerging as major regulators of these cellular processes⁹. We sought to evaluate the therapeutic potential for glycosphingolipid modulation as a new approach to treat PKD. Here we demonstrate that kidney glucosylceramide (GlcCer) and ganglioside GM3 levels are higher in human and mouse PKD tissue as compared to normal tissue, regardless of the causative mutation. Blockade of GlcCer accumulation with the GlcCer synthase inhibitor Genz-123346 effectively inhibits cystogenesis in mouse models orthologous to human autosomal dominant PKD (Pkd1 conditional knockout mice) and nephronophthisis (*ick* and *pcy* mice). Molecular analysis *in* vitro and in vivo indicates that Genz-123346 acts through inhibition of the two key pathways dysregulated in PKD: Akt protein kinase-mammalian target of rapamycin signaling and cell cycle machinery. Taken together, our data suggest that inhibition of GlcCer synthesis represents a new and effective treatment option for PKD.

PKD is transmitted as an autosomal dominant (AD) or autosomal recessive (AR) trait. Mutations in polycystin-1 and polycystin-2 are responsible for ADPKD, the most common form of PKD^{1, 10, 11}. Recessive forms of PKD include ARPKD, a childhood disease, and nephronophthisis, the most frequent genetic cause of end-stage renal disease in the first three decades of life^{11, 12}. Although multiple forms of PKD have differing clinical manifestations, common mechanisms promote cystogenesis at the cellular and molecular levels¹. The products of the various genes mutated in PKD are expressed in primary cilia or centrosomes¹². Multiple molecular mechanisms contribute to PKD, including aberrant ciliacell cycle signaling, intracellular calcium dysregulation, Wnt pathways, cAMP-activated proliferation and the Akt–mammalian target of rapamycin (mTOR) pathway^{13, 14, 15, 16}. Enhanced understanding of these mechanisms and development of animal models orthologous to human ADPKD, ARPKD and nephronophthisis has led to the discovery of new potential therapies^{1, 17}. At present, there is no mechanism-based treatment available for PKD. Sphingolipids and glycosphingolipids regulate many cellular processes, including proliferation, apoptosis and modulation of cell signaling pathways^{1819, 20, 21}. It is becoming increasingly recognized that glycosphingolipids have key roles in the progression of a number of diseases, including diabetes and cancer¹⁸. Glycosphingolipids are key components of membrane rafts, modulating cell surface receptors, including the epidermal growth factor, insulin and insulin-like growth factor-1 receptors^{22, 23, 24, 25}. Alterations of glycosphingolipid metabolism, with elevated GlcCer and lactosylceramide (LacCer) abundance, have been documented in human ADPKD and the *cpk* mouse model, suggesting a role in cystogenesis^{19, 26}. LacCer can act as a growth factor on kidney epithelial cells, thereby directly contributing to cystogenesis¹⁹. Recently, the gangliosides GM3 and GM1 were detected in primary cilium of epithelial cells²⁷. Therefore, changes in glycosphingolipid metabolism in cystic epithelial cells may have a major role in driving cyst growth.

We hypothesized that inhibiting glycosphingolipid synthesis and lowering the abundance of GlcCer and its derivatives with specific GlcCer inhibitors might effectively treat PKD. We used Genz-123346, an orally available inhibitor of GlcCer synthase that shares some structural features with previously developed compounds and blocks the conversion of ceramide to GlcCer^{28, 29, 30}. A consequence of inhibiting GlcCer is decreased expression of downstream lipids, including GM3, which has been linked to a positive effect on glycemic control in rodent models of diabetes³⁰. A similar compound very effectively reduced GlcCer accumulation in preclinical models of Gaucher's disease and was well tolerated in phase 1 and 2 clinical trials^{31, 32}.

To determine whether abnormal glycosphingolipid metabolism is a common feature of PKD, we compared GlcCer and GM3 levels in normal and cystic kidneys of human ADPKD, its orthologous *Pkd1* conditional knockout mouse model and *jck* and *pcy* mice orthologous to human nephronophthisis^{33, 34, 35} (Fig. 1). We observed significantly increased GlcCer and GM3, but not ceramide, abundance in all cystic samples analyzed compared to normal controls (Fig. 1b). These data reveal that altered glycosphingolipid metabolism is a hallmark of human and mouse PKD and may be mechanistically involved in dysregulation of cell cycle progression and proliferation. If so, lowering GlcCer abundance by specifically targeting GlcCer and GM3 from cultured rat kidney epithelial cells with either Genz-123346 or GlcCer synthase–specific siRNA delayed cell cycle progression (Supplementary Fig. 1).

We reasoned that reducing GlcCer levels by treatment with the GlcCer synthase inhibitor Genz-123346 may block cell cycle and proliferation and attenuate cystogenesis *in vivo*. We gave *jck* mice 0.1% or 0.2% Genz-123346 in their feed from 4 to 9 weeks of age. Genz-123346 treatment resulted in a dose-dependent reduction of renal GlcCer and GM3 levels (Fig. 2a) that translated into effective inhibition of cystic disease (Fig. 2b,c and Supplementary Table 1). Because PKD develops over the life of an individual, safety is a major consideration. Genz-123346 was generally well tolerated in mice, although we noted a slight reduction in body weight gain at the highest dose (Supplementary Table 1). Notably, efficacy was achieved at lower doses as well, with no associated body weight loss (Supplementary Table 1). A similar compound has recently proven to be well tolerated in people with Gaucher's disease in phase 1 and 2 clinical trials³².

We have previously shown that mechanisms of cystogenesis in *jck* mice have multiple similarities to human ADPKD, including activation of mitogenic signaling pathways, dysregulated cell cycle and increased apoptosis^{33, 36}. To determine primary molecular targets affected by glycosphingolipid modulation in response to Genz-123346 *in vivo*, we

acutely treated *jck* mice with established disease (7 weeks of age) with Genz-123346 for only 5 d (Fig. 3a-d). Such short-term treatment proved sufficient to reduce kidney GlcCer levels without considerably affecting cystic growth (data not shown). Therefore, we could assess primary targets responsible for the treatment effects. Western blot analysis of treated kidneys showed a direct effect of Genz-123346 on the cell cycle machinery, as evidenced by reduced cyclin D expression and reduced Rb phosphorylation, suggesting G1/S cell cycle arrest (Fig. 3b). Reduced proliferating cell nuclear antigen (PCNA) levels confirmed the inhibitory effect of Genz-123346 on proliferation (Fig. 3b). This mechanistic effect of GlcCer synthase inhibition might be responsible for its therapeutic efficacy in PKD, as we have previously demonstrated that direct blockade of the cell cycle with the CDK inhibitor roscovitine results in a robust arrest of PKD in preclinical models². A large body of evidence suggests that glycosphingolipids play a key part in mediating cell proliferation⁹. Exogenous LacCer can increase kidney proximal tubular cell proliferation¹⁹. Also, increasing GlcCer levels in Madin-Darby canine kidney cells with a β -glucosidase inhibitor promotes proliferation, whereas inhibition of GlcCer synthase activity with 1-phenyl-2decanoylamino-3-morpholino-1-propanol hydrochloride (PDMP) decreases proliferation³⁷. PDMP treatment of NIH3T3 cells also results in effective inhibition of proliferation through blockade of the cell cycle 20 .

We also observed a direct effect of Genz-123346 on the Akt-mTOR signaling pathway, with reduced phosphorylation of Akt and ribosomal protein S6 (Fig. 3d and Supplementary Fig. 2). Furthermore, mTOR has also been validated as a target for PKD through multiple preclinical trials^{3, 7, 8}. In contrast, apoptosis or mitogen-activated protein kinase kinase (MEK)-extracellular signal–regulated kinase (ERK) signaling were not directly affected by Genz-123346 (Fig. 3c,d).

Because primary targeting of the Akt-mTOR and cell cycle pathways by Genz-123346 may indirectly affect other pathways of cystogenesis upon chronic treatment, we analyzed the long-term effects of GlcCer synthase inhibition in *jck* mice treated with Genz-123346 from 4 to 9 weeks (Fig. 3e–h). In addition to the inhibition of the cell cycle and Akt-mTOR pathways, chronic administration of Genz-123346 indirectly inhibited apoptosis and MEK-ERK signaling in *jck* kidneys (Fig. 3g,h). Of note, Genz-123346 had no effect on any molecular pathway analyzed in wild-type treated kidneys, suggesting its effect is limited to diseased kidneys (Fig. 3f–h).

To determine whether cyst-lining cells are responsible for elevated glycosphingolipid levels in diseased kidneys, we derived immortalized epithelial cells from wild-type and *jck* kidneys. Cultured cystic cells showed high expression of GlcCer synthase mRNA (Supplementary Fig. 3a) and elevated GlcCer and GM3 levels (Supplementary Fig. 3b) compared to wild-type cells. Notably, Akt-mTOR signaling was activated in cultured cystic cells and was attenuated in response to Genz-123346 treatment (Supplementary Fig. 3c). The *in vitro* data support the idea that altered glycosphingolipid metabolism modulates cell signaling and proliferation cell autonomously in kidney epithelial cells. Although our data support a link between aberrant glycosphingolipid synthesis and cystogenesis, the exact mechanisms are largely unknown and need to be investigated.

To further evaluate whether GlcCer synthase inhibition can effectively treat a slowly progressive, adult form of PKD characterized by cyst formation and fibrosis, we tested efficacy in *pcy* mice with 0.2% Genz-123346 in feed between 4 and 15 weeks of age. Genz-123346 lowered kidney GlcCer and GM3 abundance (Fig. 4a) and effectively inhibited cystogenesis and fibrogenesis in *pcy* mice (Fig. 4b–d and Supplementary Table 2). Therefore, inhibition of GlcCer synthase activity retards PKD progression in two different models of nephronophthisis.

To strengthen the argument for glycosphingolipid modulation as a new therapeutic strategy for ADPKD, we tested Genz-123346 efficacy in an orthologous mouse model with a conditionally inactivated *Pkd1* gene. Such models have only recently been produced, and confirmatory trials of drugs shown to be effective in nonorthologous models are emerging^{4, 8}. We generated mice with a germline null allele for *Pkd1* (*Pkd1*^{tm1Gzbd}), a conditional knockout allele with *lox* sites flanking exons 21–23 (*Pkd1*^{tm1Gztn}) and a tamoxifen-regulatable *Cre* gene^{36, 38}. Cystogenesis was induced by injecting tamoxifen into nursing females at postnatal day 5 (P5) and progressed with renal functional decline over 4–5 weeks. Genz-123346 treatment between day 7 and day 33 significantly lowered kidney GlcCer and GM3 amounts (Fig. 4e) and inhibited cystogenesis, as evidenced by reduced kidney to body weight ratio, cystic volume and blood urea nitrogen (BUN) (Fig. 4f,g and Supplementary Table 3).

The data presented here show that glycosphingolipid metabolism is altered in *jck* and *pcy* mouse models of nephronophthisis and the *Pkd1* conditional knockout mouse model of ADPKD. Inhibition of GlcCer synthase alters glycosphingolipid metabolism and effectively blocks disease progression in mouse PKD. Mechanism-of-action studies suggest that GlcCer synthase inhibition results in effective cell cycle arrest and inhibition of the Akt-mTOR pathway, ultimately leading to decreased apoptosis and mitogenic signaling. Together, these results demonstrate that modulation of glycosphingolipid metabolism is a new and effective approach for the treatment of PKD.

Methods

Mouse handling and treatment

Mice were handled in accordance with Genzyme Institutional Animal Care and Use Committee guidelines. Genz-123346 is a specific GlcCer synthase inhibitor that does not inhibit other enzymes such as nonlysosomal glucocerebrosidase, acid β-glucosidase, digestive saccharases and debranching enzyme at concentrations that effectively inhibit GlcCer synthase activity³¹. We administered Genz-123346 ad libitum to ick mice by mixing in powdered 5053 diet (Pharmaserv) at 0.225% or 0.1125% (wt/wt) from 26 to 64 d of age or 50 to 55 d of age, as indicated in the Results and Figures 2 and 3 legends. Because *ick* mice show sex dimorphism in disease progression, we analyzed males and females separately³³. Fibrosis is an insignificant component of PKD in *jck* mice and was therefore not tested in this model. We performed *ick* genotyping as previously described³³. We maintained pcy mice on a CD1 genetic background by intercrossing homozygous mice³⁴. We administered Genz-123346 ad libitum to pcy mice from 4 to 15 weeks of age by mixing in powdered 5053 diet at 0.225% (wt/wt). Pkd1 conditional knockout mouse generation is described in the Supplementary Methods (*Pkd1*^{tm1Gztn} allele). We bred females homozygous for the Pkd1 conditional knockout allele to males homozygous for a tamoxifeninducible *Cre* allele³⁸ and heterozygous for a *Pkd1* germline mutation³⁶ (*Pkd1*^{tm1Gzbd} allele) to generate mice heterozygous for the Cre allele, heterozygous for the Pkd1^{tm1Gztn} conditional allele and either heterozygous for the *Pkd1*^{tm1Gzbd} germline allele (mutant) or carrying a wild-type *Pkd1* allele (wild-type controls). We induced Cre recombinase activity by injecting the nursing females with tamoxifen (250 mg per kg body weight in sunflower oil) on P5 to deliver it to the pups in the milk. We treated pups with Genz-123346 at 25 mg per kg body weight per dose twice daily from P7 to P20 and then administered 0.15% Genz-123346 in feed from P21 to P33. We killed the mice by CO₂ asphyxiation before organ harvest. BUN was determined with a VetAce analyzer (Alfa Wasserman).

Histological analysis

We quantified cystic volume as described previously². To quantify fibrosis, we digitized Mallory's trichrome–stained kidney sections with an ACIS II system (Chromavision) and used Metamorph software (Molecular Devices) to quantify the percentage of fibrotic area to total tissue area.

Glycosphingolipid analysis

We obtained kidney samples from people with ADPKD undergoing nephrectomy for endstage renal disease from Bioserve Biotechnologies and Cooperative Histology Tissue Network (CHTN). We obtained normal human kidney samples from Bioserve Biotechnologies. Informed consent was obtained by Bioserve Biotechnologies and the Cooperative Histology Tissue Network prior to sample collection, following approval by the relevant Institutional Review Boards. Kidney samples were homogenized at 100 mg ml¹ in distilled water with a Mini Beadbeater (Biospec Products) following the manufacturer's protocol. We extracted sphingolipids with a modified Folch method³⁹. We dried the supernatant under a stream of nitrogen, reconstituted in a methanol-chloroform-water mixture, which we then diluted fivefold with 0.2% formic acid (vol/vol) and 5 mM ammonium formate in 1:1 methanol-acetonitrile. Sphingolipids were seperated with an Agilent 1100 HPLC system (Agilent) equipped with a Waters Xbridge Phenyl 3.0×100 mm 3.5-µm column (Waters) and analyzed the eluent by electrospray ionization mass spectrometry with an API-4000 mass spectrometer (Applied Biosystems). Measurements took place in positive ion mode. All multiple reaction monitoring transitions included m/z264.2 as the product ions. We normalized the results to total phosphate level, determined with a previously developed method⁴⁰. The sphingolipid extract was digested in 15% nitric acid in a microwave oven and then analyzed by inductively coupled plasma atomic emission spectroscopy (Varian Instruments). We verified the results of relative quantification by performing absolute quantification of GlcCer levels in kidney tissues using HPLC as described previously 31 (data not shown).

Western blot analysis

We homogenized samples on ice in RIPA buffer (Boston BioProducts) containing 1 mM dithiothreitol, 5 mM EDTA, 2 mM NaF, 1 mM Na₃VO₄ (all supplied by Sigma-Aldrich), Pefabloc SC and Complete protease inhibitor cocktail (both from Roche Applied Science). We determined protein concentrations by BCA protein assay (Pierce). We loaded equal amounts of protein on 4-12% NuPage Bis-Tris gels following the manufacturer's protocols (Invitrogen). We performed electrophoretic transfer onto polyvinylidene difluoride membranes (Millipore) in a semi-dry apparatus according to the manufacturer's instructions (Genomic Solutions). We blocked membranes with 5% nonfat milk in Tris-buffered saline (TBS) containing 0.1% Tween-20 and incubated with primary antibodies overnight at 4 °C. Primary antibodies were detected with horseradish peroxidase-labeled secondary antibodies at 1 in10,000 dilution (antibody to rabbit or mouse IgG: Promega; antibody to rat IgG: eBioscience). Immunoreactive proteins were elucidated by enhanced chemiluminescence (GE Healthcare). Primary antibodies to the following proteins were used: Bcl-x_L, Akt, ApaF-1, cleaved poly-(ADP-ribose) polymerase, cyclin D3 (all from BD Biosciences), phospho-Akt (Ser473), IGF-IRB, S6 ribosomal protein, phospho-S6 ribosomal protein (Ser235/236), cyclin D1, phospho-Cyclin D1 (Thr286), phospho-Rb (Ser780), total ERK, phospho-ERK (Thr202/Tyr204) (all from Cell Signaling Technologies), cyclin D2 (Biosource International), caspase-2, caspase-3 proform, MEK1/2, epidermal growth factor receptor (all from Millipore), PCNA and GAPDH (US Biological).

Statistical analysis

Data are expressed as means \pm s.e.m. Comparisons were made by two-tailed *t* tests with GraphPad Prism software (GraphPad Software, Inc.); significance was accepted at the 0.05 level of probability (*P* < 0.05).

Additional methods

Detailed methodology is described in the Supplementary Methods.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1. Altered glycosphingolipid metabolism in human and mouse PKD kidneys

(a) Schematic representation of glycosphingolipid metabolism. The *de novo* synthesis of sphingolipids is a multistep process that results in the production of ceramide from serine and palmitoyl-CoA. Ceramide can be deacylated by ceramidase to generate sphingosine, which can be phosphorylated by sphingosine kinase to generate sphingosine-1-phosphate. Alternatively, ceramide can be converted to or generated from sphingomyelin, a membrane lipid, or glycosylated by UDP-glucose:ceramide glucosyltransferase (GlcCer synthase) to generate GlcCer. GlcCer can be further modified by the addition of galactose to LacCer, which can be modified by GM3 synthase to GM3. GM3 can subsequently be modified to generate other gangliosides. Genz-123346 is a direct inhibitor of GlcCer synthase. (b) Kidney GM3, GlcCer and ceramide levels measured by liquid chromatography–mass spectrometry analysis in human ADPKD, 24-day-old *Pkd1* conditional knockout (cKO) mice (deleted on P1 and P2), 9-week-old *jck* mice and 30-week-old *pcy* mice relative to normal controls. AU = arbitrary units. Data are expressed as means \pm s.e.m. *n* = 10 humans with PKD and 3 normal humans; *n* = 3 for all mouse samples. **P* < 0.05 compared to control.

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Figure 2. Blockade of GlcCer synthase activity with Genz-123346 lowers renal GlcCer abundance and effectively inhibits PKD in *jck* mice

(a) Kidney GlcCer and GM3 amounts in male mice treated with 0.1% or 0.2% Genz-123346. *P < 0.05 compared to vehicle controls. n = 3 per group. (b) Dose-dependent effect of Genz-123346 on PKD progression in *jck* males. Shown are quantitative analyses of kidney to body weight ratio (Kidney / BW), cystic volume and BUN in *jck* males. *P < 0.05 compared to vehicle controls. (c) Representative kidney sections from *jck* mice treated with 0.2% Genz-123346 and vehicle controls. Data are expressed as means \pm s.e.m. The number of mice in each group is shown in Supplementary Table 1.

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Figure 3. Molecular pathways of cystogenesis affected by inhibition of GlcCer synthase *in vivo* (**a**–**d**) The effect of acute inhibition of GlcCer synthase. Representative immunoblots of protein extracts from 55-day-old male *jck* kidneys after 5 d of treatment with vehicle or 0.2% Genz-123346. (**a**) Schematic of the treatment regimen. (**b**) Immunoblots of cell cycle markers in kidneys of vehicle-treated (V) and Genz-123346–treated (T) mice. (**c**) The effect of Genz-123346 on apoptosis. Immunoblots of apoptotic markers expressed in kidneys of vehicle-treated and Genz-123346–treated mice. (**d**) Western blot analysis of Akt-mTOR and MAP kinase pathways. (**e–h**) The effect of long-term inhibition of GlcCer synthase activity on cystogenesis. Western blot analyses of whole kidney lysates from wild type (WT) and *jck*

males treated with either vehicle or 0.2% Genz-123346 from 26 to 64 d of age. (e) Schematic of the treatment regimen. (f) Cell cycle marker analysis. (g) Apoptotic marker analysis. (h) Analysis of cell signaling pathway activity. GAPDH, glyceraldehyde-3phosphate dehydrogenase; S6, ribosomal protein S6; P-S6, phosphorylated ribosomal protein S6; Cyc, cyclin; Bcl-x_L, Bcl-2–like protein-1; ApaF-1, apoptotic peptidase activating factor-1; IGF-1R, insulin-like growth factor-1 receptor.



Figure 4. Genz-123346 effectively inhibits PKD in *pcy* mice and in *Pkd1* conditional knockout mice

(**a**–**d**) The effect of Genz-123346 on cystogenesis in *pcy* mice. (**a**) Kidney GlcCer and GM3 levels after Genz-123346 treatment from 4 to 15 weeks of age. n = 3 per group. (**b**) Quantitative analysis of kidney/body weight (BW) ratio, cystic volume and fibrosis in treated *pcy* mice. (**c**) Representative kidney sections from treated *pcy* mice and vehicle controls at 15 weeks. (**d**) Mallory's trichrome–stained kidney sections showing reduced fibrosis in Genz-123346 treated *pcy* mice. (**e**–**g**) The effect of Genz-123346 on cystogenesis in *Pkd1* conditional knockout mice. (**e**) Kidney GlcCer and GM3 abundance in conditional knockout mice treated from 7 to 33 d of age. n = 3 per group. (**f**) Kidney/body weight ratio, cyst volume and BUN in treated *Pkd1* conditional knockout mice. (**e**) conditional knockout mice. (**e**) Representative kidney sections from vehicle-treated and Genz-123346–treated *Pkd1* conditional knockout mice. **P* < 0.05 compared to vehicle controls. Data are expressed as means ± s.e.m. The number of mice per group for each experiment is indicated in Supplementary Tables 2 and 3.