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## Outcomes of Patient Self-Referral for the Diagnosis of Several Rare Inherited Kidney Diseases

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## Abstract

**Purpose**—To evaluate self-referral from the Internet for genetic diagnosis of several rare inherited kidney diseases.

None.

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**Methods**—Retrospective study from 1996–2017 analyzing data from an academic referral center specializing in autosomal dominant tubulo-interstitial kidney disease (ADTKD). Individuals were referred by academic health care providers (HCPs) non-academic HCPs, or directly by patients/ families.

**Results**—Over 21 years, there were 665 referrals, with 176(27%) directly from families, 269(40%) from academic HCPs, and 220(33%) from non-academic HCPs. 42(24%) of direct family referrals had positive genetic testing vs 73(27%) of families from academic HCPs and 55(25%) from non-academic HCPs (P=.72). 99% of direct family contacts were white and resided in zip code locations with a mean median income of \$77,316±34,014 vs. US median income \$49,445.

**Conclusions**—Undiagnosed families with Internet access bypassed their physicians and established direct contact with an academic center specializing in inherited kidney disease to achieve a diagnosis. Twenty-five per cent of all families diagnosed with ADTKD were the result of direct family referral and would otherwise have been un-diagnosed. If patients suspect a rare disorder that is un-diagnosed by their physicians, actively pursuing self-diagnosis using the Internet can be successful. Centers interested in rare disorders should consider improving direct access to families.

#### Keywords

Internet; rare disease; autosomal dominant tubulo-interstitial kidney disease; uromodulin; mucin-1

## Introduction

Poor access to healthcare often refers to individuals in need of care who cannot obtain it for socioeconomic reasons. Individuals with rare disorders also have difficulty accessing proper care, but their obstacles to care are unique. While it is estimated that 10% of individuals suffer from a rare disease [1], each disease is unique, preventing a common pathway to diagnosis. Often, the patient's primary physician is unaware of the correct diagnosis and the presentation of a rare disorder. Other factors preventing a physician's search for a diagnosis include time constraints from a high patient caseload, a simple lack of interest, or a belief that securing the diagnosis will not alter the patient's treatment plan. Specialists likewise may be unable to identify the disorder and do not explore further. If a researcher with expertise is identified, their practice may be very far from the patient, possibly on another continent. The plight of parents of children with undiagnosed, rare disorders has been described in the lay literature [2,3]. Even physician-parents may be unsure how to pursue a diagnosis [4]. While there are anecdotal reports of families finding a diagnosis when their physicians were unable [5], there have been no systematic investigations of individuals with rare disorders and their path to diagnosis.

The Internet has become increasingly important in the self-diagnosis of health conditions [6,7], and focused information on rare disorders may lead to increasing diagnoses of these conditions. The Internet provides the patient with a rare disorder the opportunity to find the rare specialist interested in this disorder.

#### Background

Since 1999, our group has studied three rare genetic conditions that result in autosomal dominant tubulo-interstitial kidney disease (ADTKD) [8]. The cardinal manifestations of these disorders include autosomal dominant inheritance of chronic kidney disease and a bland urinary sediment. Autosomal dominant inheritance results in many affected family members. Patients proceed to end-stage kidney disease requiring a kidney transplant or dialysis between the 3<sup>rd</sup> and 7<sup>th</sup> decades. ADTKD-*MUC1* (OMIM #174000) is due to pathogenic variants in the *MUC1* gene and has no other associated symptoms [9]. ADTKD-*UMOD* (OMIM #162000, #603860) is caused by pathogenic variants in the *UMOD* gene [10] encoding uromodulin and is associated with a high prevalence of gout. ADTKD-*REN* (OMIM #613092) is caused by pathogenic variants in the gene encoding renin [11] and is associated with childhood anemia, gout, and hyperkalemia. These conditions present a diagnostic challenge because clinical findings are nonspecific, the conditions are rare, recently identified, and not well known to nephrologists.

In 1999, there were fewer than ten US families with ADTKD described in the literature. After identification of the genetic cause of ADTKD-*UMOD* [10] in 2002, a concerted attempt was made to identify as many ADTKD families as possible, communicating not only with healthcare providers (HCPs), but also directly with families through Internet webpages [12]. In 2009, pathogenic variants in *REN*[11] and in 2014 pathogenic variants in *MUC1* [9] were identified as causes of ADTKD. Due to complexities in sequencing *MUC1*, the Broad Institute is the only laboratory that provides approved clinical testing for ADTKD-*MUC1* [13]. All samples tested at the Broad Institute are first received and processed at Wake Forest School of Medicine. Thus, this investigation included all individuals who underwent clinical genetic testing for ADTKD-*MUC1*.

The purpose of this analysis was to determine whether providing families with suspected rare disorders direct access to a research team through the Internet would result in genetic diagnoses of ADTKD and to compare direct access to families with indirect access to genetic testing through HCPs. As there was no diagnostic testing for these disorders prior to gene identification and we were the primary clinical academic center in in the US interested in and actively trying to identify families with these disorders, we were also able to characterize the natural history of how rare conditions are diagnosed after they are definitively characterized.

## Methods

#### Recruitment

The following multifaceted approach was used to educate and generate potential referrals: publication of articles [14], reviews [15], and a Kidney Disease: Improving Global Outcomes (KDIGO) consensus report [16], presentations at national meetings, invited lectures at academic centers, publication in other venues, including GeneReviews<sup>®</sup>, Renal and Urology News<sup>®</sup>, and the medical information resource UpToDate<sup>®</sup>. In 2003, 1,000 letters were mailed to US nephrologists requesting referral of any families with a suspected diagnosis of ADTKD. Several times over the last decade, Wake Forest School of Medicine

sent brochures that included articles on ADTKD to US nephrologists. To encourage affected individuals or their families to contact us directly,, a webpage was created [12], encouraging individuals who thought that they or their family might be affected with ADTKD to contact us. Information was also provided through the National Organization of Rare Disorders [17].

#### **Family Evaluation**

*UMOD* and *REN* genetic analysis was performed by commercial laboratories or the laboratory of Stanislav Kmoch, Ph.D [15]. Genetic testing for ADTKD-*MUC1* has previously been described [13].

(Figure 1), (Figure S1), (Figure S2), and (Figure S3) provide flow diagrams of the evaluation process. All referrals were evaluated by the lead author. The study did not include data on bulk shipment of samples specifically for *MUC1* genetic analysis from international centers specializing in renal genetics. For HCP referrals, the HCP would provide patient contact information if the patient was agreeable. In some cases, a clinical diagnosis other than ADTKD was made (either before or after negative genetic testing), and the patients were referred for genetic testing elsewhere. If agreeable, patients were consented and arrangements were made for a local laboratory to obtain a blood sample for genetic analysis at no cost to the patient. If patients did not provide consent or provide blood samples after several reminders by email and phone over one year, they were considered to have declined participation. In all cases, genetic testing was provided free to patients. The characteristics of the person initially referred by HCPs or the first clinically affected family member who agreed to participate are described. Median income according to zip code was obtained from 2010 United States Census Data [18].

Study data were collected and managed using REDCap (Research Electronic Data Capture) electronic data capture tools hosted at Wake Forest School of Medicine [19]. REDCap is a secure, web-based, NIH-sponsored application that supports confidential data capture for research studies.

This study was approved by the institutional review boards of Wake Forest School of Medicine, The First Faculty of Medicine, Charles University, Prague, Czech Republic, and the Broad Institute of MIT and Harvard, Cambridge, MA and adhered to the Declaration of Helsinki.

#### **Statistical analysis**

Statistical analysis was performed with SAS statistical programming (Cary, NC). Chisquared testing was used to compare the three referral groups. For continuous variables, a general linear model was created, with type III sums of squares reported and the direct family referral group selected as the reference group.

## Results

Over 21 years, there were 828 referrals. There were 77 individuals (9%) who already had a genetic diagnosis, 34 individuals (4%) referred directly from the first author's clinical practice, and 52 (6.3%) referrals specifically for pediatric gout. These three groups were

excluded from further analysis. The remaining 665 referrals included 269 (41%) from HCPs at academic institutions, 220 (33%) from HCPs at non-academic institutions, and 176 (27%) direct family referrals, defined as an affected or unaffected member of a family that does not have a known diagnosis of inherited disease contacting us through our website for concerns that they might have ADTKD, without the guidance or assistance from their healthcare providerEight (1%) referrals from family members who were also physicians were included in the direct family referral group.

Of the 489 HCP referrals, 408 (83%) were from nephrologists, 46 (9%) non-nephrologist physicians, 30 (6%) geneticists or genetics counselors, and 5 (1%) HCPs not classified to other groups. There were 37 HCPs who referred 2 families, and 22 HCPs who referred 3 or more families. Referral sources are listed in (Table S1). Multiple avenues led to referrals. Despite increasing availability of electronic media, 252 (52%) referrals were the result of some form of personal contact, of which 97 (20%) were through personal acquaintance with the lead author, 96(20%) were at the recommendation of a colleague, and 59 (12%) resulted from personal contact at meetings or lectures. There were 103 (21%) referrals via the Internet, including UpToDate® and GeneReviews®. There were 370 (76%) referrals from the United States, with 26 (5%) from Canada and 14 (3%) from Australia.

Of the 176 direct family referrals, 116 (66%) of the initial contacts were female, and 108 (61%) were affected personally. Of unaffected individuals referring their family, 27 (24%) were parents and 14 (13%) were spouses, with 46 (41%) unknown. There were 134 (76%) referrals from the US, 7 (4%) from India, and 4 (2%) from Canada, with 2 or fewer referrals from 19 countries. All direct family referrals were through our website, except for one family that learned of our research through the Facebook® page of an affected individual.

A higher number of affected family members could be a catalyst for trying to establish a diagnosis, especially in the case of direct family referrals. Therefore, the number of known, clinically affected individuals in each family was analyzed according to referral group. The number of affected family members was similar between groups (see (Table S2)), with 22% of direct family referrals having greater than 5 affected family members vs. 25% for nonacademic HCPs and 28% for academic HCPs (P=.17).

(Table 1) shows outcomes vs. referral type. Thirty-one percent of direct family referrals were not pursued due to low likelihood of ADTKD vs. 18% for academic HCP and 15% for nonacademic HCP referrals (*P*=.0003). Twenty-seven percent of non-academic HCP and 22% of academic HCP referrals declined participation vs. 17% of direct family referrals (*P*=.04). Twenty-four percent of direct family referrals underwent genetic testing that resulted in a diagnosis of ADTKD-*UMOD/MUC1/REN* vs. 27% of academic HCP and 25% of nonacademic HCP referrals (*P*=.72). Two referrals from non-academic providers led to the identification of other new genetic causes of kidney disease [20,21]. Heterozygous loss-offunction *SEC1A1* missense variants resulted in chronic tubulo-interstitial kidney disease, congenital anemia and pre- and post-natal growth retardation in one family and chronic kidney disease, anemia, and neutropenia in another family [20]. *NDUFAF6* missense variants are responsible for autosomal recessive Fanconi syndrome associated with chronic kidney disease and progressive pulmonary fibrosis [21]. Further testing of other family

members (see (Table S3)) from direct family referrals resulted in the identification of 81 individuals with ADTKD-*UMOD*, 32 individuals with ADTKD-*MUC1*, and 3 with ADTKD-*REN*. Direct family referral resulted in the diagnosis of 116 out of 565 (21%) individuals from all families identified with ADTKD. These families would have remained undiagnosed if family members had not independently sought a diagnosis on the Internet.

Characteristics of the initial contacts from direct family referrals who underwent sample collection (see (Table 2)) included a high proportion of white race, female gender, and higher median income by zip code. Of direct family referrals, 99% were white vs. 93% from academic and 92% of nonacademic referrals (P=.045). Six (5%) academic HCP referrals were African-American families vs. 0 for both non-academic HCPs and direct family referrals. The median income by zip code was substantially higher for direct family referrals (\$77,316±34,014 vs. \$65,301±29,741 academic (P=.04) and \$63,934±24,403 non-academic (P=.03)). Most individuals were referred with very advanced kidney disease, with 93% having an estimated glomerular filtration rate < 45 ml/min/1.73m<sup>2</sup> or on dialysis/ transplanted.

(Figure 2), (Figure 3), (Table S4), (Figure S4), (Figure S5), (Figure S6), and (Figure S7) display the temporal and geographic distribution of referrals. While genetic testing was available for one form of ADTKD as early as 2002, referrals have continued to increase over time. Both patient and academic referrals from Internet sources increased in approximately 2009. Despite increasing Internet resources, the primary source of HCP referrals continues to be some form of personal contact (personal knowledge of the author, referral at the suggestion of a colleague, or lectures), which has resulted in a rising number of referrals each year. Direct family referrals have decreased over the last two years, while HCP referrals have increased.

## Discussion

#### **Comparison with Prior Work**

While prior studies have questioned the effectiveness of Internet self-diagnosis [22, 23], the current investigation showed that the Internet is an important tool for the self-diagnosis of rare disorders. Direct family referrals resulted in the diagnosis of 116 family members with ADTKD, none of whom would have received a diagnosis at that time if family members had not pursued self-diagnosis. Direct family referral via the Internet contributed 29% of the families and 21% of cases of these uncommon diseases, a major resource for clinical characterization and research. Importantly, patients originating from direct family referrals had similar frequencies of positive genetic diagnoses as referrals from HCPs. These observations provide insights about not only the self-diagnosis of rare inherited kidney diseases, but also the increasing empowerment of patients and families with rare disorders [24].

#### **Principle Results**

This article highlights the strengths and weaknesses of direct Internet referral of individuals with rare disorders. Advantages include low cost, low manpower requirements for initial

referral, and the ability to bypass non-diagnosis by physicians. We observed that patients made judicious decisions about initiating contact. Based on clinical characteristics, some conditions may lend themselves better to self-diagnosis. Family members were more likely to diagnose ADTKD-*UMOD* than ADTKD-*MUC1* (see Table 1). ADTKD-*UMOD* is more easily researched on the Internet due to the presence of both gout and inherited kidney disease, while ADTKD-*MUC1* is only associated with inherited kidney disease, making Internet searches nonspecific and more difficult. The presence of many affected family members in autosomal disorders provides more exposure to physicians and/or family members who might be interested to pursue diagnosis. Thus, 36% of our referrals were from families with 4 or more affected individuals. Autosomal recessive disorders, with only one or two affected individuals in each family, may be more difficult to diagnose.

This article also characterized the natural history of diagnosis once the genetic causes of ADTKD were identified. Though genetic testing was available for the first of these disorders in 2002, referrals from HCPs continue to increase over a decade later. Direct family referrals have begun to decrease, possibly due to better recognition by HCPs. Despite the increasing availability of Internet resources, physician education about these rare disorders appears to occur primarily through personal interaction, resulting in increasing referrals over time.

## Limitations

A shortcoming of our study is that we only studied one center that specialized in ADTKD, provided current knowledge on the topic, and provided a path to diagnosis. Patients searching for other conditions may be unable to find accurate information or a path to diagnosis. One report found that only 20% of English searches for health information yielded relevant results [25]. Other studies have noted the poor quality of health information available on the Internet [22,23,26]. Finding accurate information on the Internet is especially problematic for rare diseases, as approximately 50% of these disorders do not have a foundation providing specific patient information.<sup>1</sup> Despite studies showing frequent use of the Internet by parents of children with rare conditions [27–32], one study reviewing 693 websites about rare diseases found that in general the quality of information provided was poor [33]. Our results only describe outcomes that may occur when patients are provided with accurate information, personal contact, and a path to achieving diagnosis.

The potential use of the Internet for self-diagnosis also carries with it the inherent weaknesses of the digital divide. Almost all direct family referrals (99%) providing genetic samples were from white families. The median income of the zip codes from the site of referrals was \$77,316±34,014, significantly higher than the 2010 US median household income of \$49,445 [34]. There were no direct family referrals from African-Americans, and few individuals from non-English speaking countries. There is no evidence that the pathogenic variants causing ADTKD would have a difference in prevalence between races. Factors that affect access to healthcare through the Internet include availability of devices connected to the Internet. Individuals who use the Internet on a daily basis (e.g. for work) are likely to have better Internet search skills. In addition, our webpage information was provided only in English. An Italian study characterizing individuals who used the Internet to access a federation of associations of patients with rare diseases found that users were

African-Americans are more likely to have misgivings about research [35] and therefore may have been less likely to contact us, even though kidney disease is much more prevalent in African-Americans. In a focus group study of African Americans' views on the trustworthiness of physicians, Jacobs et al. reported the importance of interpersonal competence of physicians [36]. Providing a more interpersonal experience on the Internet could include the use of more video material and providing faster methods of direct contact. Screening for genetic disorders in patients in dialysis centers would also provide a better interpersonal experience and enhance access to individuals across the digital divide.

African-Americans were also less likely to be referred by their healthcare providers. The high prevalence of chronic kidney disease in African-Americans may have contributed to decreased consideration of a genetic diagnosis. Decreased access to healthcare providers may have also contributed.

Despite many resources in the literature and on the Internet, 52% of HCP referrals were based on personal acquaintance with the lead author, lectures, or conversations with colleagues who had a personal knowledge of our research. Once nephrologists referred one family with ADTKD, they were likely to refer others. Unfortunately, most individuals from all three referral groups only undertook to establish a diagnosis when kidney disease was quite advanced and patients were near dialysis.

Strengths of this article include that this center was one of very few referral centers involved in the diagnosis of ATKD-*UMOD* and the only center arranging clinical genetic testing for ADTKD-*MUC1*. Weaknesses of this study include the unknown prevalence of this disorder. An Austrian study by Lhotta and colleagues [37] estimated a prevalence of ADTKD-*UMOD* of 1.67 cases per million. Given a US population of 323 million, one would expect approximately 540 cases of ADTKD-*UMOD*, compared with the 283 cases that were identified. The relative contributions of our multifaceted interventions could not be determined. We could not account for how many families reviewed our website information and presented it to their physicians or how many individuals underwent commercial testing. We know that there were 2620 unique page views of our webpages in 2017. Similarly, for the National Organization of Rare Disorders (NORD) ADTKD website, there were 1,518 unique page views in 2017 (personal communication: Marsha Lanes, NORD). Of note, 8.2 million users visited NORD webpages in 2017. We do not know how many families were tested at commercial laboratories for ADTKD-*UMOD*, though we know that all clinical diagnoses for ADTKD-*MUC1* were only made in our laboratory.

## Conclusions

Despite these limitations, this investigation highlights the importance of providing direct knowledge and access to individuals with un-diagnosed rare disorders through the Internet. Such direct access provided many families with a genetic diagnosis that had eluded them for

generations. These families in turn provided a significant proportion of individuals willing to participate in clinical research. We believe that endeavors such as the Broad Institute Rare Genomes Project are likely to further empower patients to find the genetic causes of rare inherited diseases [38]. For aid in the diagnosis of rare inherited kidney diseases, please contact ableyer@wakehealth.edu.

## **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

Disclosures

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Figure 1. Flow diagram of all referrals for evaluation for autosomal dominant tubulointerstitial kidney disease (ADTKD).

Flow diagram of 828 family referrals for ADTKD evaluation.



## Figure 2. Temporal distribution for referral type.

Red represents direct family referrals, blue represents academic healthcare providers (HCP), and aqua represents non-academic HCP.





Red represents direct family referral via the Internet, with all family referrals being generated through Internet searches. Blue represents health care provider (HCP) personal contact, including personal contact between the first author and the provider, colleague referral, and also lectures given by the first author. Green represents HCP referrals via Internet searches. Purple represents HCP referrals via reading of the literature. Aqua represents HCP referrals via mass mailing.

#### Table 1.

## Outcomes according to referral type.

	Direct Family Referrals <sup>a</sup>	Academic HCP <sup>a</sup>	Non-academic HCP <sup>a</sup>	Total <sup>a</sup>	P value
Declined participation or lost to follow- up	29 (16.5)	57 (21.2)	59 (26.8)	145 (21.8)	.04
Genetic diagnosis not pursued due to low likelihood of ADTKD	54 (30.7)	48 (17.8)	33 (15.0)	135 (20.3)	.0003
In progress	7 (4.0)	21 (7.8)	19 (8.6)	47 (7.1)	.16
ADTKD-MUC1	11 (6.3)	36 (13.4)	18 (8.2)	65 (9.8)	.03
ADTKD-UMOD	30 (17.1)	33 (12.3)	35 (15.9)	98 (14.7)	.32
ADTKD-REN	1 (0.6)	4 (1.5)	2 (0.9)	7 (1.1)	.63
Genetic testing negative for ADTKD; pursuing other genes	19 (10.8)	32 (11.9)	28 (12.7)	79 (11.9)	.84
Other clinical diagnosis	25 (14.2)	38 (14.1)	24 (10.9)	87 (13.1)	.45
SEC61A1 pathogenic variant	0	0	1 (0.45)	1 (0.15)	.36
NDUFAF6 pathogenic variant	0	0	1 (0.45)	1 (0.15)	.36
Total	176 (26.5)	269 (40.5)	220 (33.1)	665 (100)	

<sup>*a*</sup>Data shown as number (%).

#### Table 2.

Characteristics of first affected contact in families who underwent sample collection for the study.

	Direct Family Referrals	Academic HCP	Non-academic HCP	P value
Ν	68	120	101	
Gender (% male)	40(58.8)	63(52.5)	52(51.5)	.6
Race (% white)	67(98.5)	111(92.5)	93(92.1)	.045
Age (y)	$47.9 \pm 15.8^{a}$	43.7 ± 17.2	44.7 ± 14.6	
End-stage kidney disease at referral (%)	32(18.2)	47(17.5)	38(17.3)	.97
Estimated glomerular filtration rate (ml/min/1.73m <sup>2</sup> ) <sup>b</sup>	$14.8 \pm 20.1^{C}$	$22.4\pm27.1$	$24.3\pm26.7$	
US referrals (%)	134 (76.1)	191(71.0)	179 (81.4)	.03
Mean median income by zip code (\$)	77,316 $\pm 34,014^{d}$	65,301±29,741	63,934 ± 24,403	

<sup>a</sup>There were no statistical differences between groups.

 $b_{\rm Estimated}$  glomerular filtration rate defined as 0 ml/min/1.73 m<sup>2</sup> for individuals with end-stage kidney disease at the time of referral.

<sup>*C*</sup>The mean estimated glomerular filtration rate was significantly different for direct family referrals vs. nonacademic referrals (P=.03), but not significantly different vs. academic referrals (P=.08).

dMedian income was significantly different for direct family referrals vs. non-academic referrals (P=.03) and vs. academic referrals (P=.04).