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WJCC mainly publishes articles reporting research results and findings obtained in the field of clinical medicine and covering a wide range of topics, including case control studies, retrospective cohort studies, retrospective studies, clinical trials studies, observational studies, prospective studies, randomized controlled trials, randomized clinical trials, systematic reviews, meta-analysis, and case reports.

INDEXING/ABSTRACTING
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CASE REPORT

CYP27A1 mutation in a case of cerebrotendinous xanthomatosis: A case report

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Abstract

BACKGROUND
Cerebrotendinous xanthomatosis (CTX) is a rare autosomal recessive metabolic disease caused by mutations in CYP27A1. It has a low incidence rate, insidious onset, and diverse clinical manifestations. It can be easily misdiagnosed and can go unrecognized by clinicians, leading to delayed treatment and worsened patient outcomes.

CASE SUMMARY
A 38-year-old male was admitted to our hospital with a history of unabating unstable posture and difficulty in walking for more than 30 years. Subsequently based on the patient's medical history, clinical symptoms, magnetic resonance imaging and gene sequencing results, he was finally diagnosed with CTX. Due to the low incidence rate of the disease, clinicians have insufficient knowledge of it, which makes the diagnosis process more tortuous and prolongs the diagnosis time.

CONCLUSION
Prompt diagnosis and treatment of CTX improve patient outcomes.

Key Words: Cerebrotendinous xanthomatosis; CYP27A1; Sterol 27-hydroxylase; c.380G>c.1563dupA; Case report

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Core Tip: Cerebrotendinous xanthomatosis (CTX) is a rare disease for which prompt diagnosis and treatment improve patient outcomes. In addition, unreported new mutation and previously reported mutation were found in this patient. Thus, it provides new data for the further study of the pathogenesis of CTX and enriches the pathogenic mutation spectrum of CYP27A1.

INTRODUCTION

Cerebrotendinous xanthomatosis (CTX) is a rare autosomal recessive lipid deposition disorder characterized by systemic signs and neurological dysfunction[1]. CTX is a treatable genetic metabolic disease, and early diagnosis and treatment can delay the progression of the disease to a considerable extent[2]. We report a case of CTX caused by mutations at two sites in CYP27A1. This case report will help clinicians to better understand CTX and its presentation, leading to early diagnosis and treatment, thereby improving the quality of life of patients.

CASE PRESENTATION

Chief complaints
A 38-year-old male was admitted to our hospital with a history of unabating postural instability and difficulty in walking for more than 30 years.

History of present illness
The patient was first brought for treatment at the age of 5 years. Clinical documentation at that time reported that the patient had exhibited unabating postural instability and difficulty in walking that did not improve with rest. He also exhibited cognitive impairment and irritability.

History of past illness
Since childhood, the patient experienced frequent episodes of chronic diarrhea lasting multiple weeks. At the age of 11 years, the patient underwent bilateral cataract surgery. At the age of 36 years, the patient presented with bilateral masses on the Achilles tendons coupled with thickening of the Achilles tendons.

Personal and family history
The patient had no specific personal and family medical history.

Physical examination
Neurological examination revealed gait ataxia, increased muscle tension in both lower limbs, bilateral hyperreflexia of the Achilles and knee tendons, a bilateral positive Babinski sign, and bilateral positive ankle clonus, indicating that the pyramidal tracts were damaged bilaterally. In addition, the patient also presented with arched feet and egg-sized, hard, painless lumps in both achilles tendons (Figure 1).

Laboratory examinations
Blood lipid level examination revealed a total cholesterol concentration of 4.03 mmol/L (reference range: 2.60–5.20 mmol/L).

Imaging examinations
Magnetic resonance imaging (MRI) (Figure 2A and B) of the brain showed T2-weighted and FLAIR imaging hypertensity in the bilateral cerebellar dentate nuclei. Electroencephalography showed abnormal slow-wave activity, composed of θ and δ waves, bilaterally in the posterior regions. MRI (Figure 2C and D) of the right ankle indicated fusiform swelling and abnormal signals in the Achilles tendons.

Genetic testing
Based on the patient's medical history, clinical manifestations, and imaging analyses, it was unclear if
Figure 1 These two pictures (A, B) showed arched feet and egg-sized, hard, painless lumps in both achilles tendons.

CTX was involved, and gene sequencing was required to confirm the diagnosis. After informing the patient, the patient was eager to identify the underlying cause and had hopes for treatment; therefore he agreed to undergo gene sequencing analyses. Genomic DNA was extracted from the peripheral blood cells of the patient, and first-generation sequencing of the exon coding region of CYP27A1 revealed that the gene had a compound heterozygous mutation of c.380G>A (Figure 3) and c.1563dupA (Figure 4). Further examination demonstrated that the mother and sister of the patient were carriers of the c.1563dupA mutation.

Figure 2 Magnetic resonance imaging. A, B: Magnetic resonance imaging (MRI) of the brain showed T2-weighted and FLAIR imaging hyperintensity in the bilateral cerebellar dentate nuclei; C, D: MRI of the right ankle showed fusiform swelling and abnormal signals in the achilles tendons.

**FINAL DIAGNOSIS**

Based on the patient's medical history, clinical manifestations, auxiliary examinations and gene sequencing results, the diagnosis of CTX was confirmed.
TREATMENT
After the diagnosis of CTX, the patient was prescribed chenodeoxycholic acid (CDCA) at 250 mg three times per day and instructed to adhere to a low-cholesterol diet.

OUTCOME AND FOLLOW-UP
After 1 year of treatment, the patient felt that the symptoms of weakness in both lower limbs had improved slightly, but he did not report any additional changes. The patient reported no adverse reactions to CDCA.

DISCUSSION
The main cause of CTX is sterol 27-hydroxylase deficiency caused by the mutation of CYP27A1 [3]. CYP27A1 encodes sterol 27-hydroxylase and is the only gene known to be associated with CTX [4]. Sterol 27-hydroxylase is involved in the biosynthesis of primary bile acids, including cholic acid and CDCA [5]. Sterol 27-hydroxylase deficiency obstructs the synthesis of primary bile acids, which causes the accumulation of bile acid synthesis pathway intermediates and derivative metabolites such as cholesterol and cholestanol. These substances are easily deposited in various lipophilic tissues, and therefore, they are more common in the brain, lens, and tendons [6]. They can negatively influence the function of cellular calcium channels, destroy the stability of cell membranes, and initiate the apoptosis pathway [7].

Currently, CTX is considered a rare disease, as are only a few hundred reported cases worldwide. However, we believe that this value is likely underestimated owing to the diversity of symptoms and the frequent delay in diagnosis. Consequently, the number of CTX cases is likely to be considerably far
higher than that reported. The primary manifestations of CTX are infant-onset chronic refractory
diarrhea, juvenile-onset bilateral cataracts, tendinous xanthomas, and progressive neurological
dysfunction[8]. Neurophenotypes include ataxia, pyramidal tract signs, cognitive impairment, and
peripheral neuropathy[8]. When the above symptoms are unexplained by common diseases at any age,
the possibility of CTX should be considered, and further examinations should be performed. In the case
of our patient, almost all the aforementioned symptoms occurred, although they varied in age at
presentation. The constellation of symptoms observed in our patient caused us to consider CTX before
other possible diseases.

There are no recognized diagnostic criteria for CTX; thus, clinicians must make their diagnoses based
on medical history, family history, and clinical characteristics, following which the diagnosis must be
confirmed by performing blood biochemistry tests, plasma cholesterol assessment, MRI, and gene
sequencing. Our patient’s gene sequencing results showed two mutation sites in CYP27A1: c.380G>A
located on exon 2 and c.1563dupA located on exon 9. Among these, the mutation site c.380G>A has been
reported previously[9], but the c.1563dupA mutation is novel and yet to be reported. As our patient
possesses a known pathogenic mutation in CYP27A1, we are currently unable to determine whether the
new gene mutation is pathogenic; further research is needed to confirm the same. In addition, the
biochemical diagnosis of CTX is based on the increase in serum cholestanol and urine bile alcohols
levels[4]. The typical imaging findings indicating the prevalence of CTX include T2-weighted and
FLAIR imaging hyperintensity in the dentate nucleus[10]. The current case supports the inclusion of
high signal intensity of the two dentate nuclei on MRI as a typical feature of CTX[11]. The typical
imaging manifestations of CTX are high signal in T2 weighted imaging and FLAIR imaging of dentate
nucleus.

In CTX treatment, there is currently no clear treatment plan, and the condition can be treated sympto-
matically based on the different clinical manifestations. Bile acid supplements, such as CDCA, provide a
source of primary bile acids, which can inhibit the synthesis of bile acids through a negative feedback
mechanism, thereby prevent the accumulation of cholesterol and cholestanol[12]. Consequently, early
oral bile acid supplement treatment is recommended[13]. In addition, cataract extraction is common
performed in these patients, and xanthoma can be surgically removed.
Our patient was prescribed CDCA replacement therapy. At a follow-up visit 6 months post-treatment, the symptoms of the patient had improved slightly. Based on previous studies, we understand that the clinical process of CTX is progressive[14]. Saussy et al[14] compared three cases of CTX and concluded that early and uninterrupted treatment can delay progression of the disease, avert nervous system involvement, and improve the quality of life of patients. Once the neurological symptoms are completely determined, the therapeutic effect will be considerably reduced.

CONCLUSION

Herein, we reported a case where first-generation sequencing of CYP27A1 was performed in a patient with CTX, leading to the detection of an unreported new mutation as well as a previously reported mutation. Consequently, this case provides new data for further examination of the pathogenesis of CTX and enriches the pathogenic mutational spectrum of CYP27A1. In addition, the diagnosis of our patient helped him to receive genetic counseling and guidance regarding fertility. We hope that our case report enables other clinicians to more deeply understand the diagnosis and treatment of CTX, leading to early diagnoses and treatment and improved patient prognoses.

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FOOTNOTES

**Author contributions:** Li ZR acquired patient information and prepared the manuscript; Zhou YL prepared the Figure; Jin Q and Xie YY checked the literature and proposed the idea of publishing this case study; Meng HM reviewed and edited the manuscript; and All authors read and approved the final manuscript.

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