



Case Report

A novel mutation in *CYP17A1* gene leads to congenital adrenal hyperplasia: A case report

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Abstract

Background: Congenital adrenal hyperplasia is a rare autosomal recessive disorder where the mutation in P450 family 17 subfamily A member 1 gene (*CYP17A1*) is involved in its etiology. The disorder represents itself with low blood levels of estrogens, androgens, and cortisol that generally couples with hypertension, Hypokalemia, sexual primary amenorrhea, infantilism and in affected individuals.

Case: In this study, the *CYP17A1* gene in a 14-year-old female was examined. The karyotype of the patient was 46, XX, and the analysis of the *CYP17A1* gene by Sanger sequencing revealed a novel homozygous deletion c.1052-1054CCT which led to isolated 17,20-lyase deficiency.

Conclusion: In conclusion, this study report an in-frame deletion which results in isolated 17, 20-lyase deficiency, and the mutation might be used for diagnosis in other patients with distinctive clinical symptoms.

Key words: Congenital adrenal hyperplasia (CAH), *CYP17A1* gene, Ambiguous genitalia.

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1. Introduction

Congenital adrenal hyperplasia (CAH) is a group of rare disorders demonstrated by a failure in one of the five enzymes responsible for cortisol production in autosomal recessive pattern (1). In adrenal glands, Pregnenolone is made from cholesterol which later can be processed to either mineralocorticoids or glucocorticoids or to sex steroids in adrenals and gonads (2).

The p450c17 enzyme is encoded by the *CYP17A1* gene located on 10q24.3 (3) and spans 6.6 kb, which comprises eight exons (4). "This gene transcribes 2.1-kb mRNA molecule, which expresses in both the adrenals and gonads and generates a 57-kDa microsomal cytochrome P450c17 enzyme. The *CYP17A1* enzyme catalyzes both steroid 17-hydroxylase and 17,20-lyase activities" (5). Enzymatic failure of the P450c17 enzyme leads to both glucocorticoids and sex steroids deficiencies. After a reduction in blood glucocorticoids levels due to P450c17 defect, anterior pituitary tries to compensate for the insufficiency of glucocorticoids levels by producing extra Adrenocorticotrophic hormone (ACTH). This results in the extra generation of steroid precursors and elevated ACTH level, which leads to some *Congenital* adrenal hyperplasia and result in hypertension, hypokalemia, and a suppressed renin-angiotensin system (6, 7). Moreover, due to the impairments in *CYP17A1*, the mineralocorticoid precursors (corticosterone and 11-deoxycorticosterone) accumulate, which demonstrate glucocorticoid activity, therefore defect in P450c17 does not associate with adrenal crisis, rather than other CAH variants (8). Mutations in the *CYP17A1* gene are the rarest defects in CAH that yields to steroid 17-hydroxylase and 17,20-lyase deficiencies (9). Several mutations in the *CYP17* gene have been reported that cause either complete or combined 17-hydroxylase/17,20-lyase or isolated 17, 20-lyase enzyme deficiencies (10-14).

The purpose of this study was to investigate the molecular defects in *CYP17A1* gene and its relationship with Congenital adrenal hyperplasia.

2. Case presentation

A 14-year-old female, the first child of consanguineous parents with normal family history was referred to the genetic clinic with high blood pressure, ambiguous genitalia, and lack of pubertal development. The blood sample was taken after receiving written informed consent from her parents. No pubic or axillary hair was seen by physical examination, and she had no clinical symptoms of Turner syndrome with 46, XX karyotype. In the sonographic survey, uterus was infantile. She was hypertensive (150/90 mmHg, 50th percentile for age) with high gonadotropins levels (LH, 19 mU/mL; FSH, 34 mU/mL). Moreover, low peripheral concentrations of sex steroids were seen (Table I).

2.1. Sequencing of *CYP17A1* gene

Genomic DNA was purified from peripheral blood leukocytes (PBL) using QIAGEN Mini Blood kit. All the exons of *CYP17A1* gene were proliferated by PCR (primers listed in Table II), which were designed with Primer3 software (<http://primer3.sourceforge.net>). All the PCR products were sequenced in both directions by sanger sequencing.

2.2. Mutational analysis

As shown in Figure 1, a new in-frame homozygous deletion c.1052-1054CCT in exon 6 was identified that reported for the first time. 17 α -hydroxylase deficiency was first pronounced by Biglieri and colleagues (15), who was phenotypically female and presented with sexual infantilism, primary amenorrhea and hypertension.

Table I. Clinical and hormonal characteristics

Blood pressure (mmHg)	150/90
Karyotype	46,XX
Tanner stage (breast/pubis hair)	B1P1
ACTH (0-60 pg/mL)	76
Cortisol (5-25 µg/dL)	17
LH (Female, 12-18 yr: 0.1-10 mU/mL)	19
FSH (Female, 12-18 yr: 0.3-9 mU/mL)	34
Estradiol (30-120 pg/mL)	15
DHEA (350-4300 ng/mL)	52
Progesterone (0.1-1.3 ng/mL)	4.6

Table II. Primers for *CYP17A1* amplification

Exon	Primers	Sequences (5'-3')	Areas	Product size (bp)
1	1F	CACTGCTGTCTATCTTGCC	1802-2277	476
	1R	CCTTCACATCATCCCACTA		
2	2F	AGGGACCAGAGGTGTAAG	3730-4070	341
	2R	GCAGCAGTAGCCAAGAA		
3	3F	AGGGTGTGATTCATTC	4132-4544	413
	3R	GCAGAGGAGGTAGAGGTG		
4	4F	CGCTTGATGTTTGATTGA	4819-5214	396
	4R	CACCCTGCTCTTGATGATT		
5	5F	ACAGAAGTATGGCAGGAGT	5776-6289	514
	5R	CCAGAGTAGGTTGGAGGT		
6	6F	ACTGGGAAGGGACTGGA	6182-6496	315
	6R	GGCTAGATGTCCTGGGAG		
7	7F	AGTGGGAATGAGGGAGTA	7244-7599	356
	7R	GTCAACAGGTCGGTATAGTT		
8	8F	TCAACCAGGGCAGAACC	7914-8359	446
	8R	GGAAGAATGGCGGAGAA		

F: Forward

R: Reverse

impairment of 17-hydroxylase activity. The alteration in 351Leu which lies in the redox partner-binding domain of P450c17 leads to an impairment in lyase activity of P450c17 (14, 19), in other words, disrupt interactions of redox partner proteins with CYP17A1.

In sum, this case manifested typical feature of 17 α -hydroxylase and 17,20-lyase deficiencies (e.g., hypertension and ambiguous genitalia). This study is the first paper to report an in-frame deletion which results in isolated 17, 20-lyase deficiency, and this mutation might be used for diagnosis in other patients with distinctive clinical symptoms. Identification of 17 α -hydroxylase/17,20 lyase deficiency was confirmed by the particular profile of adrenal steroid levels, and further confirmation by CYP17A mutation analysis.

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Conflict of Interest

The authors declare no conflict of interest.

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