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Defects in Steroidogenic Enzymes. Discrepancies Between Clinical Steroid Research and Molecular Biology Results*

M. Zachmann

Department of Pediatrics, University of Zurich, Kinderspital, CH-8032 Zurich, Switzerland

Molecular biology has clarified the understanding of steroidogenic enzyme genetics. Nevertheless, there are discrepancies between fundamental and clinical experience. (1) Why do patients with "pure" 17α-hydroxylase or 17,20-desmolase deficiency exist, when one cytochrome regulates both steps? A case of interest is discussed, who had "pure" 17,20-desmolase deficiency until adolescence, but additional 17α -hydroxylase deficiency thereafter. (2) In 11β -hydroxylase deficiency, it was puzzling to find 18-hydroxylated compounds, and, in isolated hypoaldosteronism, normal cortisol, since 11\beta- and 18-hydroxylation were thought to be regulated together. This has now been explained by differences in the fasciculata and glomerulosa. The occurrence of 11β-hydroxylase deficiency of 17-hydroxylated steroids only, however, remains enigmatic. (3) 3β -Hydroxysteroid dehydrogenase deficiency does not only seem to exist in classic (mutations of type II gene), but also in late-onset cases. In them, no molecular basis could be found. (4) Also, in cholesterol side-chain cleavage, there is an inequity: while evidently one cytochrome regulates 20- and 22-hydroxylation, pregnenolone is formed when 20xOH-cholesterol, but not when cholesterol, is added to adrenal tissue of deficient patients. Other factors (promoters, fusion proteins, adrenodoxin, cAMP-dependent expression of genes, and/or proteases), or hormonal replacement in patients may be responsible for these discrepancies.

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INTRODUCTION

Understanding of the genetics and regulation of steroidogenic enzymes and of their disorders has made much progress as a consequence of new developments in molecular biology. Nevertheless, there are some discrepancies between newly acquired fundamental knowledge and older clinical experience, which can at present not be completely understood.

In the following, selected examples are discussed to illustrate some apparent differences between clinical observation and molecular findings.

17α-HYDROXYLASE AND 17,20-DESMOLASE (OR 17,20-LYASE) DEFICIENCIES

Classic deficiency of 17α -hydroxylase originally described in adults [1] blocks the biosynthesis of cortisol,

androgens and estrogens. Among steroids not hydroxylated in position 17, DOC and its 18-hydroxylated metabolites may cause hypertension. Deficiency of 17,20-desmolase [2] allows efficient 17-hydroxylation, but prevents the side-chain cleavage of 17α OH-pregnenolone and/or 17α OH-progesterone. Androgens and estrogens are lacking, whereas cortisol and aldosterone are normal. Male pseudohermaphroditism (XY-individuals) or absent pubertal development (XX-individuals) are the only symptoms [2–8].

In spite of these two clearly separate clinical entities, biochemical studies with 17-(bromoacetoxy) progesterone already performed years ago suggested that a cytochrome P450 from testicular microsomes consisted of a single protein with both enzymatic activities [9]. Estradiol-17 β was also found to inhibit both activities [10]. Later, it was shown that most mature cytochromes are derived from a precursor with a higher mass [11]. The enzyme was active with $\Delta 4$ and $\Delta 5$ substrates as a 17α -hydroxylase and with the corresponding 17α -hydroxysteroids as a 17,20-desmolase. It was, however, observed that the activities were not

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^{*}Dedicated to Professor Andrea Prader on the occasion of his 75th birthday.

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evenly distributed, but that e.g. the 17,20-desmolase activity was largely suppressed in the microsomes.

Kinetic comparison of 17α -hydroxylase and 17,20-desmolase activities have been carried out in human adrenal microsomes [12]. The median maximum velocity of the 17α -hydroxylase was significantly greater than that of the 17,20-desmolase. Most steroids with the exception of cortisol competitively inhibited both enzyme activities. The inhibition constant was, however, always higher for the 17α -hydroxylase than for the 17,20-desmolase activity. It thus became evident that both, steroid 17α -hydroxylase and 17,20-desmolase activities reside within the same polypeptide chain, but that there are still some differences of reaction. Various genetic causes have thereafter been discovered as molecular bases for the defects [13–16].

The questions clinicians ask in this context are: why do clinical cases with "pure" 17α -hydroxylase [1] or "pure" 17,20-desmolase deficiency [2] exist separately within separate families, when a single cytochrome regulates both steps? Why do combined cases not always occur [17, 18]? What are the factors that lead to the clinical and biochemical picture of 17α -hydroxylase deficiency in one, and to that of 17,20-desmolase in another patient? Does one of the two defects remain unchanged during the whole life of a patient, or is there a developmental aspect to it?

A patient of interest in this context, whose data have been presented [19], has to be briefly discussed. This XY-individual with female external genitalia and compound heterozygous mutations [16] was followed from adolescence to adulthood. While up to the age of 16 years her 17α -hydroxylating capacity was efficient, and steroids not hydroxylated in position 17 were low, drastic steroid changes were observed thereafter. Simultaneously, the previously normal blood pressure increased. While conventional antihypertensives had no effect, dexamethasone normalized the pressure. The only iatrogenic difference after age 17 years was that estrogen and gestagen replacement at low doses had been started. The changes of integrated values of urinary steroids are shown in Table 1. The 17α-hydroxylated compounds (cortisol metabolites) decreased drastically, while compounds not hydroxylated in position 17 increased. The high quantities of compounds not hydroxylated in position 17, however, remained readily dexamethasone suppressible.

Table 1. Urinary steroids ($\mu mol/d$) in a patient with 17,20-desmolase/17 α -hydroxylase deficiency from adolescence to adulthood

Age (yr)	Cortisol metabolites	Pregnanediol	THB + allo-THB
16.9	13.1	6.1	1.9
20.7	5.0	7.7	25.5
25.2	< 0.1	26.5	76.8

From this observation we conclude that, in the same individual, 17α-hydroxylase activity was normal throughout childhood and adolescence, but decreased in adulthood under estrogen supplementation. 17,20desmolase activity, by contrast, was already completely absent in the fetus, leading to complete absence of masculinization. This case thus demonstrates that previously pure 17,20-desmolase deficiency can with time be complicated by additional 17α-hydroxylase deficiency. The factors leading to this change are unknown. A modifying effect of estrogen replacement on the cytochrome P450 C17α-activity is, however, a possibility. Also insulin inhibits adrenal 17α-hydroxylase activity in man [20], but there is no evidence in our patient that there might have been newly acquired insulin hypersecretion. It is thus conceivable that, for unknown reasons, fetuses, infants, and children with a deficiency of cytochrome P450 C17a present the signs of "pure" 17,20-desmolase deficiency, while adults show the signs of the combined defect. This might be the reason why we as pediatric endocrinologists have originally described patients with 17,20-desmolase deficiency [2], while 17α -hydroxylase deficiency (which in retrospect rather has to be regarded as "combined" deficiency) was first described in adults [1].

118- AND 18-HYDROXYLASE DEFICIENCIES

Steroid 11β -hydroxylase deficiency is, after 21-hydroxylase deficiency, the second most common cause of congenital adrenal hyperplasia. The conversion of 11-deoxycortisol (compound S) to cortisol, and that of DOC to corticosterone is blocked. In addition to androgen excess similar to that in 21-hydroxylase deficiency, hypertension may develop because of the hypersecretion of DOC, and salt-wasting is usually absent. 18-Hydroxylase deficiency has been postulated to be a cause of isolated hypoaldosteronism, where the formations of cortisol and corticosterone are normal.

The enzyme is encoded by the CYP11B1 gene on chromosome 8q22. Two mutations in CYP11B1 have previously been reported in patients with the defect (Arg-448 \rightarrow His and a 2 bp insertion in codon 394). In Jews of Morocco and in corticosterone methyl-oxidase (CMO) II deficiency in Jews of Iran, steroid 11β -hydroxylase has been found to be encoded by two homologous genes, CYP11B1 and CYP11B2. CYP11B1 encodes a specific cytochrome necessary for cortisol biosynthesis, with predominantly 11β -hydroxylase and moderate 18-hydroxylase activity, whereas CYP11B2 encodes another isozyme necessary for the biosynthesis of aldosterone with 11β -hydroxylase, 18-hydroxylase and 18-oxidase activities [21].

CYP11B2 has been shown to be the same gene as that for CMO type I and II [22]. Molecular genetic analysis of CYP11B2 of three patients affected with CMO I deficiency has revealed that deletion of 5 nucleotides occurs exclusively in exon 1, resulting in a frameshift

to form a stop codon in the same exon. Thus, P450 C18 is not produced, causing a complete lack of aldosterone biosynthesis in the patients. Additional, previously uncharacterized mutations causing 11β -hydroxylase deficiency have been reported by others [23, 24]. Some are point mutations (nonsense and missense), and one is a single base pair deletion causing a frameshift.

It has originally been puzzling for clinicians to find 18-hydroxylated compounds in patients with congenital adrenal hyperplasia due to 11β -hydroxylase deficiency, and conversely in isolated hypoaldosteronism to find normal cortisol formation, since 11β - and 18hydroxylation were thought to be regulated together [25]. This problem has been partly solved by the observation of different regulations in the zona fasciculata and glomerulosa of the adrenal cortex: it is now known that, in the rat, there are two distinct species of cytochrome $P450 \text{ C}11\beta/18$, aldosterone synthase and 11β -hydroxylase. Whereas aldosterone synthase is located exclusively in the zona glomerulosa, the distribution and site of production of 11β -hydroxylase is not entirely clear [26, 27]. Four forms of rat CYP11B genes were isolated and characterized [28], which were and -B4. CYP11B1,-B2,-B3, Southern analyses indicated that the members of the rat CYP11B gene subfamily were confined to these four genes. Among them, CYP11B1 and -B2 encoded steroid 11β -hydroxylase and aldosterone synthase, respectively, while CYP11B3 was a gene homologous to CYP11B1 without a known expression product. Müller and Oertle [29, 30] identified two isozymes of cytochrome $P450 \text{ C}11\beta$, CYP11B1 and CYP11B2 in the rat adrenal cortex. CYP11B1 is found in all zones and catalyzes a single hydroxylation of DOC in 11β - or 18-position. CYP11B2 is produced exclusively in the zona glomerulosa and catalyzes all three reactions involved in the conversion of DOC to aldosterone. In vivo and in vitro, the expression of the genes encoding CYP11B1 and CYP11B2 is regulated by two separate control systems which appear to operate both independently and interdependently. Nomura et al. [31] isolated three genomic clones of rat CYP11B. Two of them corresponded to 11β -hydroxylase gene (CYP11B1) and aldosterone synthase gene (CYP11B2). The third was a novel gene resembling CYP11B1 and CYP11B2, and was named CYP11B3.

When patients with congenital adrenal hyperplasia due to 11β -hydroxylase deficiency are studied, there are, however, still observations to be made, which are not completely clarified by the known genetic aspects [32]: some patients have a block of 11β -hydroxylation in the 17α -hydroxylated pathway only [33–35], while the 11β -hydroxylation of DOC is functioning normally. The respective steroid secretion rates of such a patient are shown in Table 2. In others, no 18-hydroxylated compounds can be found, while in most classic cases, 18OH-DOC is present or increased. Especially, the clinical fact that 11β -hydroxylase defi-

Table 2. Steroid secretion rates in a patient with congenital adrenal hyperplasia due to 11β-hy-droxylase deficiency of 17α-hydroxylated steroids only (from [34])

Compound	Secretion rate— patient (mg/m²/day)	Secretion rate— normal mean (mg/m²/day)
S	82.6	0.31
DOC	0.13	0.05
F	3.9	8.7
В	2.4	2.3

ciency may be present in the 17α -hydroxylated pathway only remains completely unexplained by the molecular findings.

3β-HYDROXYSTEROID DEHYDROGENASE DEFICIENCY

In this rare defect, steroid biosynthesis is blocked at the stage of the $\Delta 5$ -compounds, and $\Delta 4$ -compounds cannot be formed in the adrenals, but only by nonspecific reductases in the liver. DHEA is the only increased androgen. Because its action is weak, no or only mild virilization occurs. 46XY-patients are incompletely masculinized, while the genitalia in females show slight hypertrophy of the clitoris or are normal. In most cases, there is salt-wasting, but cases without it have been observed [36, 37]. Since the defect is also located in the gonads, no or insufficient pubertal development occurs [38].

The molecular basis of classic 3β -hydroxysteroid dehydrogenase deficiency has consistently been found to be a mutation of the type II gene [39–44].

Clinically, the syndrome does not only seem to exist in the mentioned classic form, but also in mild or late-onset cases, be it in children with premature adrenarche or advanced bone maturation [45, 46], or in hirsute women [47–50]. In them, unlike in the different types of 21-hydroxylase deficiency, a molecular basis could, however, so far not be found. The question from the clinicians therefore is: do such genetically determined mild cases really exist and if yes, how frequent are they? Most likely, they are much less frequent than has been suspected, and the diagnosis of mild or late-onset 3β -hydroxysteroid dehydrogenase deficiency has probably been made too often solely on the basis of increased DHEA or other 45-steroids. Since many different molecular causes have been found in late-onset 21-hydroxylase deficiency, and none in "mild" 3β -hydroxysteroid dehydrogenase deficiency, it seems that the latter has less variable causes, which could be limited to classic cases.

CHOLESTEROL SIDE-CHAIN CLEAVAGE

The incapability to cleave the side chain of cholesterol causes the syndrome of lipoid adrenal hyperplasia

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(synonyms: Prader syndrome [51], deficiency of cytochrome *P*450scc). With the exception of some inactive cholesterol metabolites, no steroids can be produced [52]. Newborns of both chromosomal sexes have female external genitalia, adrenal insufficiency, and salt-wasting.

So far, no molecular basis of this syndrome could be found. Southern blotting patterns of the P450scc gene were normal, and PCR of all exons showed normal sequences on multiple amplifications and sequencing reactions, indicating normal P450scc genes [53]. Also, Northern blots of testicular RNA from a 6-month-old patient and from a control fetus showed normal P450scc mRNA, indicating a normal P450scc promoter. Reprobing with cloned human cDNAs for adrenodoxin reductase and adrenodoxin showed that these electron transport cofactors used by P450scc were also normal, as was probing with cDNAs for all three known factors involved in cholesterol transport to the mitochondria-sterol carrier protein 2, endozepine, and steroidogenesis activator peptide. These results suggest that the lesion in lipoid CAH is not in the P450scc system or in any step upstream from P450scc. In other studies, similar results were obtained [54-57].

Also concerning the clinical lack of this enzyme, there is a biochemical observation, which is difficult to understand: while it has been sufficiently demonstrated that one single cytochrome $P450\mathrm{scc}$ regulates the 20α -and 22-hydroxylation of cholesterol simultaneously, it has been shown in earlier, but still valid and careful studies, that pregnenolone could be formed when $20\alpha\mathrm{OH}$ -cholesterol, but not when cholesterol, was incubated with adrenal tissue of patients with lipoid adrenal hyperplasia (Table 3). 20α - and 22-hydroxylation thus for unknown reasons seem to be regulated partly independently [58].

In conclusion, the progress in molecular studies, although rapid and very impressive, has so far not explained all clinical and biochemical observations in the field of the congenital defects of steroidogenic enzymes. Other factors such as promoters and the influence of fusion proteins of type II *P*450 enzymes [59], adrenodoxin as an electron donor [60], cAMP-dependent expression of genes encoding the steroidogenic enzymes [61], and/or proteases [62], which are possibly also modulated by developmental factors and/or treatment of the patients with cortisol, mineralocorticoids and sex hormones, may be responsible for the discrepancies.

Table 3. Cholesterol-cleaving activity (°0) of enzyme preparation of adrenal tissue of a patient with lipoid adrenal hyperplasia and of normal tissue (from [58])

Substrate	Patient	Normal $(n = 3)$
[4- ¹⁴ C]cholesterol	< 0.01	2
[7α-3H]20α-hydroxycholesterol	2.6	3-5

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