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High density lipoprotein as a source of cholesterol for adrenal steroidogenesis: a study in individuals with low plasma HDL-C

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Abstract Few studies have addressed the delivery of lipoprotein-derived cholesterol to the adrenals for steroid production in humans. While there is evidence against a role for low-density lipoprotein (LDL), it is unresolved whether high density lipoprotein (HDL) contributes to adrenal steroidogenesis. To study this, steroid hormone profiles in urine were assessed in male subjects suffering from functional mutations in ATP binding cassette transporter A1 (ABCA1) (n = 24), lecithin:cholesterol acyltransferase (LCAT) (n = 40), as well as in 11 subjects with low HDL cholesterol (HDL-C) without ABCA1/LCAT mutations. HDL-C levels were 39% lower in the ABCA1, LCAT, and low HDL-C groups compared with controls (all P < 0.001). In all groups with low HDL-C levels, urinary excretion of 17-ketogenic steroids was reduced by 33%, 27%, and 32% compared with controls (all P < 0.04). In seven carriers of either type of mutation, adrenocorticotropic hormone (ACTH) stimulation did not reveal differences from normolipidemic controls. In conclusion, this study shows that basal but not stimulated corticosteroid metabolism is attenuated in subjects with low HDL-C, irrespective of its molecular origin. These findings lend support to a role for HDL as a cholesterol donor for basal adrenal steroidogenesis in humans.-Bochem, A. E., A. G. Holleboom, J. A. Romijn, M. Hoekstra, G. M. Dallinga-Thie, M. M. Motazacker, G. K. Hovingh, J. A. Kuivenhoven, and E. S. G. Stroes. High density lipoprotein as a source of cholesterol for adrenal steroidogenesis: a study in individuals with low plasma HDL-C. J. Lipid Res. **2013.** 54: **1698–1704.**

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Published, JLR Papers in Press, March 19, 2013 DOI 10.1194/jlr.P033449 The synthesis and secretion of adrenal steroid hormones for regulating stress responses, electrolyte homeostasis, and maintenance of secondary sexual characteristics depends upon the availability of the precursor cholesterol (1). To secure a continuous cholesterol supply, the adrenal glands can synthesize cholesterol, metabolize intracellular esterified cholesterol, or obtain cholesterol from circulating lipoproteins (1). Although exact data are lacking, plasma lipoproteins have been suggested to contribute more than 75% of all cholesterol required for adrenal steroidogenesis (2, 3), but the current literature gives little insight in associations between plasma lipoprotein levels and adrenal function in humans.

With respect to a role for low-density lipoprotein (LDL), it has been shown that LDL receptor-deficient patients suffering from familial hypercholesterolemia display normal urinary excretion of both 17 hydroxycorticosteroids (17-OHCS) and 17 ketogenic steroid metabolites (17-KS) indicative of unaffected adrenal function (4, 5). In addition, low or absent LDL cholesterol (LDL-C) in carriers of one or two defective *APOB* alleles respectively did not affect basal adrenal function either (6). Combined this suggests that LDL is probably not playing a major role in delivering cholesterol for steroid hormone production in humans. Associations with HDL cholesterol (HDL-C) have thus far only been studied in critically ill patients. In one

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Abbreviations: A, androsteron; ABCA1, ATP binding cassette transporter A1; ACTH, adrenocorticotropic hormone; ALLO, allo-tet-rahydrocortisol; CBG, cortisol binding globulin; D, dehydroepiandrosteron; E, etiocholanolon; HA, 11-hydroxy-androsteron; HDL-C, HDL cholesterol; HE, 11-hydroxy-etiocholanolon; KA, 11-ketoandrosteron; KE, 11-ketoetiocholanolon, 17-KS, 17 ketogenic steroid metabolites; LDL-C, LDL cholesterol; 17-OHCS, 17 hydroxycorticosteroids; P2, pregnaandiol; P3, pregnaantriol; SRB1, scavenger receptor type B1; THE, tetrahydrocortisol; THF, tetrahydrocortisol; THS, 11-deoxytet-rahydrocortisol.

study it was shown that low HDL-C in such patients was associated with attenuated adrenal responses to synthetic adrenocorticotropic hormone (ACTH) (7). In support of this, others reported a high incidence of adrenal failure in critically ill individuals with liver disease, with HDL-C being the only variable predictive of adrenal insufficiency (8).

Using adrenal cells, it has been suggested that HDL is the preferred lipoprotein for cholesterol delivery to the adrenal gland (9) and in accordance, scavenger receptor type B1 (SRB1)-mediated cholesterol uptake from HDL has been shown to be the predominant source of cholesterol in mice (10–13). In line, it has been shown that mice lacking SRB1 display an impaired adrenal glucocorticoid stress response (14, 15), lending support to a major role for HDL as a cholesterol donor in mice. In humans, we also showed that adrenal function was compromised in individuals with a functional mutation in SCARB1, the gene encoding SRB1 (16). While this study showed that cholesterol delivery to the adrenals via the HDL-SRB1 pathway is important for adrenal steroidogenesis in humans, it is unclear whether plasma HDL-C levels are associated with adrenal steroidogenesis in humans. To investigate this, we assessed basal and ACTH-stimulated adrenal cortical function in males with low HDL-C due to mutations in either ATP binding cassette transporter 1 (ABCA1) (17) or lecithin:cholesterol acyltransferase (LCAT) and in subjects with low HDL-C without mutations in ABCA1/LCAT, as well as in normolipidemic controls (18). We hypothesized that in subjects with low HDL-C levels, adrenal function would be compromised irrespective of the molecular origin of the low HDL-C.

METHODS

Recruitment of study participants

Male subjects with HDL-C levels below the fifth percentile were screened for mutations in ABCA1 and LCAT (17, 19), of which the functionality was assessed in previously published studies (17, 20, 21). For the current study, we enrolled 24 carriers of mutations in the ABCA1 gene. We furthermore enrolled 40 male carriers of mutations in the LCAT gene. In addition, subjects with similarly reduced HDL-C levels without mutations in ABCA1 and LCAT were included (n = 11). As a control group, normolipidemic age-matched male individuals were recruited by advertisement. None of the included individuals used medication interfering with steroid metabolism. The study was approved by the institutional review board of the Academic Medical Center, Amsterdam, The Netherlands and all participants provided written informed consent.

Questionnaire and biochemical measurements

Medical history, cardiovascular risk factors, use of medication, and family history of cardiovascular disease were assessed using a questionnaire. Brachial artery blood pressures were measured using an oscillometric blood pressure device (Omron 705IT, Hoofddorp, The Netherlands). Hypertension was defined as *I*) use of antihypertensive medication, or *2*) a systolic blood pressure at visit above 140 mm Hg and/or diastolic blood pressure above 90 mm Hg.

Plasma was obtained after an overnight fast and stored at -80° C. Total cholesterol, LDL-C, HDL-C, and triglyceride levels were analyzed using commercially available enzymatic methods (Randox, Antrim, UK and Wako, Neuss, Germany) on a Cobas Mira autoanalyzer (Roche, Basel, Switzerland). Free cholesterol and total cholesterol were measured before and after precipitation of apoB-containing lipoproteins using phosphotungstic acid (Sigma) and a commercially available enzymatic assay (Diasys) on a Selectra autoanalyzer (Sopachem). ACTH was determined by an immunoluminometric assay (Nichols Institute, Los Angeles, CA). Aldosterone was measured using a radioimmunoassay (Siemens, Los Angeles, CA).

Baseline adrenal steroidogenesis

Urinary excretion of steroid metabolites was analyzed by gas chromatography in 24 h urine samples as previously described (22, 23). Androsteron (A), etiocholanolon (E), dehydroepiandrosteron (D), 11-ketoandrosteron (KA), 11-ketoetiocholanolon (KE), 11-hydroxy-androsteron (HA), 11-hydroxy-etiocholanolon (HE), pregnaandiol (P2), pregnaantriol (P3), 11-deoxytetrahydrocortisol (THS), tetrahydrocortison (THE), tetrahydrocortisol (THF), and allo-tetrahydrocortisol (ALLO) were measured as readout of adrenal steroidogenesis. A, E, D, KA, KE, HA, and HE made up the total 17-KS, whereas THS, THE, THF, ALLO, and P3 were the constituents of the total 17-OHCS. In addition, urinary free cortisol was determined using solid-phase extraction liquid chromatography- tandem mass spectrometry on a Symbiosis Pharma (Spark Holland, Emmen, The Netherlands) Quattro Premier tandem mass spectrometer (Waters, Milford, MA) system. Solid phase extraction was performed on Oasis HLB cartridges (Waters), chromatographic separation was achieved on a Waters Sunfire C18 column (3.5 μ m, 2.1 × 50 mm) using ammonium acetate mM with 0.1% formic acid as mobile phase and acetonitrile as mobile phase B. Limit of detection 5 nmol/l, intra-assay variation <4%, total assay variation <7%.

Stimulated adrenal steroidogenesis

Random subgroups of seven *ABCA1* and seven *LCAT* mutation carriers consented to an ACTH stimulation study (cosyntropin or tetracosactin, 0.25 mg/ml; Novartis Pharma B.V., Arnhem, The Netherlands). After an overnight fast, participants underwent cosyntropin testing at 0900 h. Two baseline blood samples were obtained, 15 min and 1 min before administration of the 1 μ g cosyntropin bolus. Subsequent blood samples were drawn 30 min and 60 min after cosyntropin administration. Plasma cortisol levels were measured by enzyme immunoassay (Siemens Medical Solutions, Los Angeles, CA), and cortisol binding globulin (CBG) levels were measured with a commercial radioimmunoassay (Siemens Medical Solutions). Free cortisol levels were calculated using the method described by Coolens, van Baelen, and Heyns (24).

Statistical analysis

Unpaired Student's *t*-test was performed for analysis of continuous data with a normal distribution. In case of a skewed distribution, data were log-transformed prior to *t*-testing. Categorical data were assessed by χ^2 testing. A *P* value of <0.05 was considered statistically significant. A linear regression model was used to correct for differences in LDL-C and statin use.

RESULTS

Population characteristics

We enrolled 24 and 40 carriers of loss-of-function mutations in either *ABCA1* or *LCAT*. Two of the *ABCA1* mutation

TABLE 1. Characteristics of male ABCA1 and LCAT mutation carriers and matched male controls

	ABCA1					LCAT				
Characteristics	Carriers (n = 24)	Controls (n = 24)	Р	Low HDL-C (n = 11)	Р	Carriers (n = 40)	Controls (n = 40)	Р	Low HDL-C (n = 11)	Р
Demographic										
Age, years	45.2 ± 20.6	45.5 ± 18.5	0.96	44.6 ± 12.0	0.91	46.2 ± 12.8	45.2 ± 14.9	0.84	44.6 ± 12.0	0.71
BMI, kg/m2	25.3 ± 3.3	25.1 ± 5.2	0.92	26.3 ± 8.8	0.81	26.3 ± 2.9	24.9 ± 4.5	0.17	26.3 ± 8.8	0.99
Current smoker, n (%)	8 (33.3)	4 (17.4)	0.21^{a}	3 (27)	0.72^{a}	7 (18.4)	4(10.8)	0.35^{a}	3 (27)	0.52^{a}
Alcohol users, n (%)	17 (71)	19 (79)	0.51^{a}	9 (82)	0.49^{a}	36 (90)	33 (83)	0.33^{a}	9 (82)	0.46^{a}
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Coronary artery disease,	7 (29.2)	1 (4.8)	_	4 (46)	—	9 (23.7)	1 (2.9)	_	4 (36)	—
Diabetes mellitus, n (%)	3(13.6)	0 (0)	0.07^{a}	1(9)	0.71^{a}	1(2.6)	0(0)	0.32^{a}	1(9)	0.34^{a}
Hypertension, n (%)	7(29.2)	3(12.5)	0.16^{a}	1(9)	0.19^{a}	13(32.5)	4 (10)	0.01^{a}	19(8)	0.12^{a}
Statin use, n (%)	8 (33)	2 (8)	0.02^{a}	4 (33)	0.84^{a}	15 (38)	2(5)	< 0.001 ^a	4 (36)	0.80^{a}
Blood pressure, mm Hg	0 (00)	- (-)		- ()		()	- (-)		- ()	
Systolic	136 + 16	132 + 13	0.34	127 + 11	0.29	139 + 17	132 + 13	0.14	127 + 11	0.19
Diastolic	79 + 12	80 + 10	0.75	78 + 11	0.90	81 + 9	80 + 9	0.81	78 + 11	0.57
Biochemical										
Aldosterone, nmol/l	0.20 ± 0.14	0.23 ± 0.12	0.55	0.22 ± 0.19	0.87	0.19 ± 0.11	0.24 ± 0.14	0.12	0.22 ± 0.19	0.68
ACTH, ng/l	23.11 ± 12.62	21.64 ± 15.10	0.25^{b}	n.a.		22.00 ± 14.25	24.57 ± 23.08	0.81^{b}	n.a.	
Total cholesterol, mmol/l	3.9 ± 1.2	5.2 ± 1.3	0.001	4.5 ± 2.1	0.27	4.4 ± 1.1	5.3 ± 1.2	0.001	4.5 ± 2.1	0.95
LDL cholesterol. mmol/l	2.7 ± 1.0	3.6 ± 0.9	0.003	3.2 ± 1.8	0.39	3.2 ± 0.9	3.6 ± 0.9	0.08	3.2 ± 1.8	0.34
HDL cholesterol, mmol/l	0.8 ± 0.3	1.3 ± 0.3	< 0.001	0.8 ± 0.2	< 0.001	0.8 ± 0.3	1.3 ± 0.4	< 0.001	0.8 ± 0.2	< 0.001
Triglycerides, mmol/l	_	_	_	1.5	_	_	_	_	_	_
Median	1.0	1.2	0.65^{b}	1.3-2.1	0.02^{b}	1.25	1.25	0.22^{b}	1.5	0.23^{b}
Interguartile range	0.7 - 1.5	0.8-1.9		_		1.0 - 2.0	0.8 - 1.8	_	1.3 - 2.1	_
Urinary free cortisol, nmol/24 h	60.10 ± 31.66	59.27 ± 39.96	0.62^{b}	n.a.	_	50.33 ± 36.34	57.81 ± 35.19	0.31^{b}	n.a.	—

Blood pressure measurements were available in 21 *ABCA1* mutation carriers and 17 controls and in 34 *LCAT* mutation carriers and 26 controls. Plasma ACTH levels were available in 18 *ABCA1* mutation carriers and 18 controls and in 20 *LCAT* mutation carriers and 31 controls. Urinary cortisol levels were available in 22 *ABCA1* mutation carriers and 22 controls and in 35 *LCAT* mutation carriers and 36 controls. Values are means ± SD unless otherwise indicated. No *t*-test was performed for history of coronary artery disease because referral bias was present. Partial overlap exists between the two control cohorts. *P* is for Student's *t*-test. n.a., not available.

^{*a*}*P* is for χ^2 test.

^bTriglycerides, plasma ACTH, and urinary cortisol were log-transformed prior to *t*-test.

carriers were compound heterozygotes while three of the *LCAT* mutation carriers were homozygotes. None of the participants were referred to our clinic for symptoms of adrenal dysfunction. Individuals were matched to male controls for age. In parallel, we also included subjects with equally low HDL-C levels without mutations in *ABCA1* or *LCAT*. Demographic, clinical, and biochemical characteristics of all low HDL-C groups as well as controls are listed in **Table 1**. As expected, HDL-C levels were 39% lower in carriers of *ABCA1* or *LCAT* mutations and in the low HDL-C group, compared with normolipidemic controls (P < 0.001). LDL-C levels were 25% lower in carriers of *ABCA1* mutations (P = 0.003), 11% lower in carriers of LCAT mutations (n.s.), and 12% lower in the low HDL-C group (P = 0.34 and P = 0.39) compared with controls.

Detailed analysis of the concentrations of cholesteryl esters and free cholesterol in the total cholesterol fraction and the isolated HDL-C fraction, in a subset of patients, is provided in **Table 2**.

Compared with controls, hypertension was more prevalent among *LCAT* mutation carriers (P = 0.01). Statin use was more prevalent in the *ABCA1* and *LCAT* mutation carriers compared with the control group. The low HDL-C group had similar baseline characteristics compared with the ABCA1 and LCAT groups.

Basal adrenal steroidogenesis

Compared with the control group, we identified lower 24 h urinary excretion of 17-KS in carriers of mutations in *ABCA1* (33%, P = 0.003; **Fig. 1A**), in *LCAT* (27%, P = 0.01;

TABLE 2.	Free cholesterol and	cholesteryl este	r values in a subse	t of LCAT	mutation carriers	and controls
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	Controls (n = 29)	Heterozygous carriers (n = 29)	Р	Homozygous carriers (n = 3)	Р
Total cholesterol, mmol/l	5.3 ± 1.2	4.5 ± 1.2	0.002	4.1 ± 0.6	0.09
FC	1.56 ± 0.35	1.4 ± 0.4	0.04	2.1 ± 0.3	0.02
CE	3.65 ± 1.04	3.0 ± 0.9	0.01	2.0 ± 0.6	0.01
FC/CE ratio	0.48 ± 0.30	0.5 ± 0.1	0.73	1.1 ± 0.4	0.001
Cholesterol after apoB precipitation	1.3 ± 0.4	0.7 ± 0.3	0.03	0.04 ± 0.01	< 0.001
FC	1.78 ± 0.71	0.1 ± 0.1	0.13	0.0 ± 0.0	< 0.001
CE	0.7 ± 0.2	0.6 ± 0.2	0.02	0.04 ± 0.01	< 0.001
FC/CE ratio	0.24 ± 0.06	0 ± 0.12	0.50	1.1 ± 0.4	< 0.001

P is for Student's t-test compared with controls. FC, free cholesterol; CE, cholesteryl ester.



Fig. 1. Twenty-four hour urinary steroid excretion in male *ABCA1* (A) and *LCAT* (B) mutation carriers compared with age-matched male controls. Data are presented as mean \pm SD. p¹, the uncorrected *P* value for Student's *t*-test; p², the *P* value corrected for LDL-C; p³, the *P* value corrected for statin use.

Fig. 1B), and in the low HDL-C group (30%, P = 0.04 and 34%, P = 0.02; Fig. 1). These differences remained statistically significant after correcting for differences in plasma LDL-C and statin use using a linear regression model (P = 0.01 and P = 0.02 for carriers of *ABCA1* mutations or *LCAT* mutations vs. controls, respectively). The mean 17-KS steroid excretion of 14.1 μ mol/24 h in the *ABCA1* mutation carriers was below the reference values for 17-KS in the appropriate age group (23).

The lower levels of urinary 17-OHCS in both *ABCA1* and *LCAT* mutation carriers, as well as the low HDL-C group, did not reach statistical significance (P = 0.11 and P = 0.15, respectively; Fig. 1A, B). The full panel of steroid metabolites is presented in **Fig. 2A, B**. No gene-dose effect was observed when comparing the two compound heterozygous *ABCA1* mutation carriers or the three homozygous *LCAT* mutation carriers to heterozygous carriers.

Free urinary cortisol was not significantly different between ABCA1 mutation carriers and controls (60.10 ± 31.66 vs. 59.27 ± 39.96 nmol/24 h, P = 0.62) or LCAT mutation carriers and controls (50.33 ± 36.34 vs. 57.81 ± 35.19 , P = 0.31).

Plasma levels of ACTH did not differ significantly between *ABCA1* mutation carriers and controls (23.11 ± 12.62 vs. 21.64 ± 15.10, P = 0.25) or *LCAT* mutation carriers and controls (22.00 ± 14.25 vs. 24.57 ± 23.08, P = 0.81).

Adrenal response to cosyntropin

In a cosyntropin stimulation test, the cortisol response to physiological levels of ACTH was measured as an approach to assess adrenal cortical function (25, 26). The peak serum cortisol response to ACTH was comparable in *ABCA1* and *LCAT* mutation carriers and did not differ from controls (P=0.10 and P=0.87 respectively, **Fig. 3A, B**). Also peak plasma levels of free cortisol, taking into account possible differences in CBG levels (16), were not different. Plasma lipid levels did not differ significantly before and after cosyntropin testing (data not shown).

DISCUSSION

The present study demonstrates that under normal conditions male individuals with low levels of HDL-C, irrespective of molecular origin, have significantly lower urinary excretion of total 17-KS compared with normolipidemic controls, indicative of disturbed basal corticosteroid synthesis. A parallel reduction in hydroxycorticosteroid excretion in both groups did not reach statistical significance which may suggest a preferential adrenal pathway for HDL driven steroidogenesis. Mutations in either ABCA1 or LCAT did not, however, affect the response to ACTH, while physical examination and a standardized questionnaire for signs of adrenal insufficiency (16) did not reveal symptoms of clinically relevant adrenal dysfunction. These findings are in line with those in mice, and tissue culture showing that the supply of plasma lipoprotein-derived cholesterol is used for basal adrenal steroidogenesis, but that this pathway is not able to respond to acute stress (27-29). For a proper quick response, endogenous adrenal cortisol reserves are secreted upon stimulation. Because steroidogenic tissues are rapidly depleted of cortisol following stimulation, replenishing is thought to occur via uptake of cholesterol from lipoproteins.

We previously reported that individuals with reduced SRB1 function displayed mild adrenal insufficiency in addition to a reduced urinary excretion of steroid hormones (16). Thus reduced SRB1 function on adrenal cells has a larger impact on adrenal steroidogenesis when compared with low levels of plasma HDL-C caused by reduced ABCA1 or LCAT function.

There were no differences in urinary excretion of free cortisol. However, particularly in the lower ranges the quantification of free cortisol has been shown to be less reliable as compared with the quantification of full urinary steroid metabolites in diagnosing the presence of hypocortisolism (30, 31).

In accordance with LCAT's mechanism of action, cholesteryl ester concentration is lower in LCAT mutation carriers, with the most pronounced effect in homozygous carriers. Because adrenal steroidogenesis was not further decreased in homozygous carriers compared with heterozygous carriers, there is no indication that the adrenal gland has a preference for cholesteryl esters as a substrate for steroidogenesis. This is supported by the fact that a similar reduction in steroid metabolites was observed in subjects with low HDL-C not related to defects in *LCAT*.

Plasma levels of ACTH did not differ between the low HDL-C group and controls. The latter most likely reflects



Fig. 2. Urinary steroid metabolites in male *ABCA1* (A) and *LCAT* (B) mutation carriers compared with age-matched male controls. Data are presented as mean \pm SD.

the fact that the decrease in urinary steroid metabolites is only mild, thereby precluding a compensatory ACTH increase. In line, none of the study participants reported signs of hypocortisolism. We would like to discuss three limitations of our study. First, adrenal cells express *ABCA1* (32). Thus reduced adrenal *ABCA1* expression could conceptually lead to accumulation of cholesterol in adrenal cells, compromising



Fig. 3. Peak serum cortisol increase after cosyntropin administration in male *ABCA1* (A) and *LCAT* (B) mutation carriers versus male controls. Data are presented as mean \pm SD. *P* values for Student's *t*-test.

intra-adrenal signaling and steroidogenesis due to cholesterol toxicity as proposed for pancreatic β -cell dysfunction in individuals carrying *ABCA1* mutations (33, 34). Although we cannot exclude this possibility, mutations in *LCAT* as well as low HDL-C in noncarrier individuals are associated with similar reductions in HDL-C and urinary steroid metabolite excretion, making an ABCA1-specific effect implausible.

A second limitation is that carriers of ABCA1 mutations showed significantly lower levels of plasma LDL-C levels in addition to low levels of HDL-C. This could mean that combined low levels of HDL-C and LDL-C account for the effects observed, but as already discussed, a role for LDL-C in human adrenal corticoid production is unlikely (4–6). In line, statistical corrections for the observed reductions in LDL-C did not affect outcome.

Third, in the urinary steroid metabolites, the 17-KS were particularly decreased in carriers compared with controls. Because 17-KS are metabolites derived largely from the androgenic pathway, this finding hints toward the preferential use of HDL-derived cholesterol in adrenal steroidogenesis. Further studies are needed to dissect whether, and to what extent, HDL-derived cholesterol contributes to steroidogenesis in either adrenal or testicular steroid production.

In conclusion, we demonstrate that basal adrenal steroidogenesis is compromised in males with low levels of plasma HDL-C, establishing a role for HDL-derived cholesterol in adrenal steroidogenesis in humans.

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