Steroid-converting enzymes in human adipose tissues and fat deposition with a focus on AKR1C enzymes

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Abstract. – Adipocytes express various enzymes, such as aldo-keto reductases (AKR1C), 11β-hydroxysteroid dehydrogenase (11β-HSD), aromatase, 5α-reductases, 3β-HSD, and 17β-HSDs involved in steroid hormone metabolism in adipose tissues. Increased activity of AKR1C enzymes and their expression in mature adipocytes might indicate the association of these enzymes with subcutaneous adipose tissue deposition. The inactivation of androgens by AKR1C enzymes increases adipogenesis and fat mass, particularly subcutaneous fat. AKR1C also causes reduction of estrone, a weak estrogen, to produce 17β-estradiol, a potent estrogen and, in addition, it plays a role in progesterone metabolism. Functional impairments of adipose tissue and imbalance of steroid biosynthesis could lead to metabolic disturbances. In this review, we will focus on the enzymes involved in steroid metabolism and fat tissue deposition.

Key Words:

Adiposity, AKR1C, Steroid converting enzymes, Adipose tissue, Steroid metabolism.

Introduction

Adipose tissue constitutes an important site for steroid hormone synthesis, metabolism, and storage¹⁻³.

Plasma dehydroepiandrosterone (DHEA), DHEA sulfate (DHEA-S), androstenedione and testosterone are taken up and transformed to active hormones in adipose tissue by various steroid-converting enzymes⁴. The steroid biosynthetic pathway in adipose tissue depends on the relative expression or activity of steroidogenic enzymes⁴.

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Steroid metabolism involves the cytochrome P450 monooxygenases superfamily, aldo-keto reductases (AKRs), short-chain dehydrogenase/reductase oxidoreductases, polyprenol reductases, uridine diphosphate glucuronosyl transferases, catechol-O-methyl transferases, sulfotransferases⁵⁻⁷, hydroxysteroid dehydrogenases (HSD), like 11 β -HSD type 1, 11 β -HSD type 2, 3 β -HSD, 17β-HSDs, and aromatase (Figure 1). These enzymes are important for steroid biosynthesis and are expressed in preadipocytes and adipocytes⁸. In particular, the most important enzymes for the pathophysiology of adipose tissue are aldo-keto reductases, hydroxysteroid dehydrogenases (HSD), and aromatase, since they regulate the homeostasis of steroid hormones in the adipocytes9. In this review, we will focus on the enzymes involved in both steroid metabolism and fat tissue deposition.

11β-Hydroxysteroid Dehydrogenase

Hydroxysteroid dehydrogenase enzymes are known to catalyze hydroxysteroid dehydrogenation. In addition, these enzymes catalyze the reverse reaction as ketosteroid reductases¹⁰.

The 11 β -HSD type 1 enzyme *in vivo* functions as a reductase that generates active glucocorticoids. In human visceral adipose tissue, 11 β -HSD1 converts inactive cortisone to active cortisol levels¹¹ to higher levels than in subcutaneous fat¹².

11 β -HSD1 knockout mice show lower weight and fat and excellent glucose tolerance, whereas moderate overexpression of the 11 β -HSD1 encoding gene in adipose tissue leads to abdominal obesity and metabolic syndrome¹³.

11 β -HSD2 expression in subcutaneous adipose tissue (SAT) has a negative association with body mass index¹⁴ and is expressed to higher levels in



Figure 1. Overview of steroidogenesis pathways. All steroid hormones are derived from a common precursor (cholesterol) through sequential steps involving several steroidogenic enzymes. Cholesterol is introduced into the cells via membrane receptor interactions and further internalized in vesicles and fused with lysosomes. Free cholesterol is released in cells by the action of lysosomal hydrolases which is then converted to pregnenolone in the mitochondria. Pregnolone is metabolized via two pathways: Δ^5 -hydroxy steroid pathway leads to the synthesis of 17 α -hydroxypregnenolone, dehydroepiandrosterone and androstenediol, and Δ^4 -ketosteroid pathway leads to the synthesis of 17 α -hydroxyprogesterone and androstenedione. Androstenedione is further activated to testosterone via catalytic action of AKR1C3 and 5 α -reductase. Androstane-3 α ,17 β -diol is a metabolite of dihydroxytestosterone. Alternatively, androstenedione and testosterone can be aromatized by CYP19A1 to form the estrogenic steroid hormones estrone and are dependent on CYP17A1 activity. Progesterone is also reduced to its less active form 20 α -hydroxyprogesterone by AKR1C1 activity. 3 β HSD = 3 β -hydroxysteroid dehydrogenase; 21OH = 21-hydroxylase; 11 α -hydroxylase; 17 α OH = 17 α -hydroxylase.

SAT of obese rats compared to lean controls¹⁵. 11 β -HSD2 overexpression leads to resistance to diet-induced obesity by decreasing food intake and increasing energy expenditure through inactivation of glucocorticoids and/or by inhibiting access to their receptors¹⁶.

Aromatase

The ovaries and adipose tissue convert androstenedione and testosterone into estrogens through P450 aromatase activity. Aromatase activity is associated with body weight in both preand post-menopausal females, and when knocked out¹⁷, both female and male mice show obesity with increased visceral fat¹⁷.

In adipose tissue, cytokines, including IL-6 and TNF- α , increase aromatase activity and transcription of the aromatase gene¹⁸. In opposition, a pulse of peroxisome proliferator activated receptor gamma agonist (PPARgA) to preadipocyte cultures of human breast cells decreases both transcription and activity of aromatase¹⁹. In the absence of PPARgA, subcutaneous abdominal preadipocyte expression of the P450 aromatase gene increases several days after induction of differentiation^{9,20}.

17β-Hydroxysteroid Dehydrogenases

17β-HSD enzymes specifically catalyze estrone to estradiol conversion in human adipose tissue and in preadipocyte cultures²¹. *In vitro*, preadipocyte differentiation to lipid-storage cells increases the activity of 17β-HSD enzyme²¹.

In humans, of fourteen isoenzymes, 17β-HSD isoenzyme type 12 plays a significant role in the formation of estrogen by catalyzing conversion of estrone to estradiol, with relatively higher expression levels in organs related to lipid metabolism including liver, heart, skeletal muscle, and kidney. Additionally, the 17β-HSD type 12 isoenzyme has a significant higher expression in endocrine-related organs, like the pituitary gland, pancreas, testis, adrenal gland, placenta, and the gastrointestinal tract, thus suggesting its regulatory role in fatty acid synthesis and steroid metabolism^{9,22}.

Glucocorticoid-Mediated Steroid Converting Enzymes

Glucocorticoid hormones are a class of corticosteroids secreted by the adrenal cortex. They are required for the regulation of different homeostatic and metabolic functions in the body. The physiological functions of glucocorticoid hormones are regulated by 11 β -HSDs that catalyze the interconversion of active cortisol and corticosterone with the inactive counterparts, cortisone and 11-dehydrocorticosterone. The active glucocorticoids bind the glucocorticoid receptor, a ligand-dependent transcription factor²³.

During fasting, glucocorticoids stimulate lipolysis in adipocytes, resulting in the production of glycerol for gluconeogenesis, and free fatty acids for energy production through oxidation²⁴⁻²⁶.

Estrogen-Mediated Steroid Converting Enzymes

Estrogens, estradiol, estriol and estrone, have a direct impact on adipose tissue metabolism and function²⁷. Enzymes involved in estradiol synthesis also modulate local and whole-body estrogen availability²⁸. Knockout mice for estrogen receptor α (ER α) have increased adiposity⁸. In agreement, variants in ER- α and ER β encoding genes are associated with increased body fat mass and visceral fat accumulation in females. Moreover, low levels of estrogens might also stimulate preadipocyte proliferation, especially in females^{29,30}. In white adipose tissue, lipid metabolism is regulated by estrogens through ER α , ER β and G protein coupled-estrogen receptors.

Progesterone-Mediated Steroid Converting Enzymes

Progesterone might stimulate fat deposition by enhancing lipid synthesis, lipoprotein lipase activity, and steroid-mediated preadipocytes differentiation. Some researchers⁹ have suggested a role for progesterone in the gynoid fat distribution pattern of females due to its anti-glucocorticoid activity in abdominal adipose tissue. In support of this hypothesis, progesterone inhibits glucocorticoid-mediated fat cell differentiation, body fat accumulation or lipogenesis in the omental adipose tissue³².

In cultured preadipocytes of rodents, progesterone enhances the expression of the sterol regulatory element binding transcription factor 1 (*Srebf1*) gene that subsequently controls fatty acid synthase transcription³³. After progesterone treatment, the levels of resistin and leptin mRNAs increase, whereas the expression of adiponectin decreases in inguinal white adipose tissue of female rats³⁴. Male rats treated with progesterone do not show any effect upon the expression of adiponectin, leptin and resistin in inguinal white adipose tissue because they possess low levels of progesterone receptors³⁴.

Androgen-Regulated Steroid Converting Enzymes

Androgens regulate the pattern of body fat distribution in males. Low plasma testosterone levels are often observed with increased visceral fat accumulation and abdominal obesity. Furthermore, androgen-based treatment of hypogonadal men results in the decrease of abdominal fat accumulation³⁵. Similarly, research studies in males have revealed substantial negative association of DHEA levels with abdominal fat accumulation, indicating that lower levels of DHEA are associated with increased abdominal fat accumulation³⁶.

AKR1 Enzymes

Hydroxysteroid dehydrogenases regulate the synthesis and inactivation of steroid hormones. These enzymes either belong to the short-chain dehydrogenase/reductase superfamily or aldo-ke-to reductase (AKR) superfamily³⁷.

In humans, 13 AKR proteins have been identified to date: the aldehyde reductase AKR1A1; the aldose reductases AKR1B1 and AKR1B10; the hydroxysteroid dehydrogenases AKR1C1, AKR1C2, AKR1C3, and AKR1C4; the Δ 4-3-ketosteroid-5- β -reductase AKR1D1; the Kv β proteins AKR6A3, AKR6A5, and AKR6A9; and the aflatoxin reductases AKR7A2 and AKR7A3³⁸. The three-dimensional structures of the above enzymes, except for AKR6A3 and AKR6A9, have been experimentally resolved, showing a conserved motif of eight α -helices and eight parallel β -strands that alternate along the peptide backbone, the typical fold of the TIM barrel³⁹, with the central cavity hosting the nicotinamide moiety of NADP(H); other than the cofactor, the structures also present the binding modes of several different steroid ligands⁴⁰.

Role of AKR1C Enzymes

AKR1Cs enzymes are expressed in different tissues. AKR1C1 is mainly expressed in testis, kidneys and liver; AKR1C2 is mainly expressed in prostate, mammary gland and liver; AKR1C3 shows higher expression in brain, testis, liver, placenta, and kidneys; and AKR1C4 is particularly expressed in the liver⁴¹. Furthermore, AKR1C1 is highly expressed in the adipose tissue and its activity is induced by adipocyte differentiation⁴¹. In both males and females, AKR1C1 expression levels are relatively higher in SAT than in omental adipose tissue⁴¹. The AKR1C enzymes play significant roles in the metabolism of prostaglandins (AKR1C3), steroid hormones (AKR1C1-AKR1C3), and bile acids and xenobiotics/drug detoxification (AKR1C4)⁴² (Table I)^{8,32,43-47}.

The aldo-ketoreductase 1C family member, AKR1C1, exhibits 17-oxoreductase activity that is involved in testosterone synthesis from 4-dione, 20-oxoreductase activity that inactivates progesterone, and 3-oxoreductase activity that

Table I. Publications reporting the activity of AKR1C enzymes in subcutaneous adipose tissue (SAT).

AKR1C enzyme	Activity	Sampling	Tissue expression	Reference
C1	2α-HSD	Women undergoing abdominal hysterectomies	Subcutaneous expression higher than omental	43
C1-C2-C3	11α-HSD-1; 3α-HSD; 17β-HSD	Women with metabolic disorders and obesity	Significantly higher in SAT, mostly for AKR1C3	8
C2	3α-HSD, 5α-HSD	Morbidly obese men undergoing biliopancreatic derivation surgery and lean to obese men undergoing	Activity significantly higher in obese men	
		general abdominal surgery		44
C3	5α-HSD	Women with PCOS	Higher expression in serum of PCOS women than in control	45
C3	17β-HSD	Women with simple obesity	AKR1C3 activity higher in SAT	46
C2-C3	11α-HSD-1; 3α-HSD; 17β-HSD	Men and women with idiopathic obesity	High expression in human SAT	47
C1	20-HSD	<i>Ex vivo</i> adipocytes isolated from women	Significantly higher expression in mature adipocytes than in preadipocytes	32

PCOS = polycystic ovary syndrome.

inactivates dihydrotestosterone⁴⁸. It has been reported that women with increased accumulation of visceral fat have higher expression of AKR1C1 mRNA and an increased 20-oxoreductase activity within omental adipose tissue^{43,49}.

In addition, AKR1C1, AKR1C2 and AKR1C3 catalyze the reduction of progesterone to produce 20α -hydroxyprogesterone, a less potent progestogen⁵⁰. Primarily, AKR1C1 catalyzes the inactivation of progesterone by converting it into 20-progesterone through its 20- α -hydroxysteroid dehydrogenase activity. AKR1C1 is expressed in SAT and in the omental adipose tissue in females, whereas it is not a prominent contributor of adipose androgen in males⁴⁶. Thus, AKR1C1 lowers progesterone and 5α -tetrahydroprogesterone levels in peripheral tissue⁴⁶. Progesterone is important for the inhibition of cell proliferation, stimulation of endometrial cell differentiation and pregnancy maintenance⁵¹⁻⁵³.

AKR1C enzymes also function as a 17-ketosteroid reductase in peripheral tissues, reducing estrone, a weak estrogen, to produce 17β -estradiol, a potent estrogen. AKR1C3 enzyme is the most efficient enzyme for this reduction reaction⁴⁶.

Role of AKR1C in Subcutaneous Adipose Tissue (SAT) Accumulation

Increased activity and expression of AKR1C enzymes in mature adipocytes might be associated with adipose tissue accumulation⁵⁴.

AKR1C2 and AKR1C3 exhibit fine regulatory effects on the availability of androgens within adipose tissue⁵⁵ while glucocorticoids reverse the effects of androgens on adipocyte differentiation. In fact, glucocorticoids eliminate androgen inhibitory action on adipogenesis, probably by increasing androgen inactivation mediated by AKR1C. This mechanism might contribute to individual differences in body fat distribution and composition; thus, reduced androgen availability at a local level allows for glucocorticoid-induced adipocytes differentiation⁵⁴.

AKR1C2 is the enzyme that plays a significant role in this crosstalk between androgens and glucocorticoids which involves regulation of lipid accumulation and adipogenesis^{56,57}.

Increased AKR1C2 expression or activity induces adipocyte differentiation by dihydrotestosterone inactivation, whereas AKR1C2-mediated androgen inactivation induced by glucocorticoids promotes adipogenesis in human subcutaneous preadipocytes. Previous studies revealed that expression of the AKR1C2 protein is increased after the maintenance or loss of weight and this increase is linked with changes in BMI, weight, plasma low density lipoprotein and waist circumference⁵⁶⁻⁵⁹.

Stimulation of AKR1C2 expression and glucocorticoid-mediated dihydrotestosterone inactivation in preadipocytes might eliminate androgen inhibitory effects on adipogenesis favoring progression of adipogenesis⁶⁰. Many scholars⁶⁰ have described further interactions between androgens and the glucocorticoid signaling pathways within adipose tissue. Such hormonal signal interactions at local levels might be an important modulators of body fat distribution patterns^{9,61}. In **Supplementary Table I**, steroid converting enzymes involved in human adipose tissue homeostasis are listed with functional polymorphisms that modulate their activity.

AKR1C Enzymes in Androgen Metabolism

The expression of AKR1C and dihydrotestosterone inactivation take place in visceral and subcutaneous adipose tissue, and inactivation rates of androgen are much higher in obese individuals⁴⁴. Furthermore, the expression of AKR1C increases with the increase in mass of adipose tissue, particularly, in subcutaneous fat, leading to higher inactivation rates of androgens⁴⁴. Additionally, in adipose tissue, AKR1C enzymes converts dihydrotestosterone, a stronger androgen into an inactive metabolite⁴⁴.

The expression of all isoforms of AKR1C increases with an increase of visceral adiposity. It has been proposed that androgens within the adipose tissue mediate central fat accumulation, preferentially causing android fat distribution⁶².

Role of AKR1Cs in Androgen Activation/Inactivation

AKR1C2 is primarily involved in the inactivation of androgen by the conversion of the potent androgen dihydrotestosterone into the weaker 3-diol by its 3-reductase activity⁶³. Androgens cause negative effects on lipid synthesis and adipogenesis by upregulating androgen receptors for catecholamine, consequently increasing lipolysis⁶³. Androgens can also modulate abdominal adipocyte accumulation by decreasing the activity of lipoprotein lipase, required for adipocyte intracellular fatty acid esterification. Hence, androgen and adipose tissue have a bidirectional and reciprocal impact on each other⁶³.

In a very interesting study⁶⁴, the Authors observed an increase in 5-dihydrotestosterone inactivation by AKR1C2 enzyme in omental adipose tissue from females with visceral obesity and proposed that the local inactivation of androgen is the main reaction catalyzed by AKR1C2 in the abdominal tissue of females. Similarly, androgen mediated inactivation of AKR1C2 activity has been observed in isolated adipocytes and in primary stromal cells. The AKR1C2 enzyme appears to have higher activity in SAT than in omental adipose tissue where inactivation of androgen is linked with obesity^{46,56}. These findings were further supported by a decrease in dihydrotestosterone levels in SAT as compared to omental adipose tissue. Subcutaneous fat is the main region of AKR1C mediated androgen metabolism both in females and males^{65,66}.

AKR1C3 inactivates progesterone to 20-hydroxyprogesterone and activates androgen receptor activity by converting androstenedione to testosterone⁶⁷. AKR1C3 expression is induced by the differentiation of adipocytes⁶⁷. In addition, the expression of AKR1C3 is increased in obese individuals, particularly in the SAT as compared to omental adipose tissue⁶⁷.

Adipocyte size could also affect the expression of AKR1C3. In fact, AKR1C3 has higher levels of expression in larger adipocytes than in smaller ones from the same subject⁴⁵.

AKR1Cs Effects on Neurosteroids

Neuroactive steroids are considered natural endogenous steroid hormone metabolites that exert non-genomic and rapid effects on neurotransmitter receptors present on the membrane. Synthesis of neurosteroids mostly involves steroidal or cholesterol precursors⁶⁸.

AKR1C2 induces the synthesis of neurosteroids, whereas AKR1C1 reduces the concentrations of neurosteroids in the human brain through 3α , 5α -tetrahydroprogesterone inactivation and elimination of the precursors of synthetic pathways⁶⁹. AKR1C isozymes preferentially work as reductases and regulate the inactive and active androgen, progestin, and estrogen concentrations in target tissues⁶⁹.

AKR1C1 also decreases the neurosteroid cellular concentrations by 5α -dihydroprogesterone and progesterone elimination from neurosteroids synthetic pathways along with the inactivation of 3α , 5α -tetrahydroprogesterone⁴⁵. Additionally, AKR1C1 is significantly involved

in the production and inactivation of the neuroactive allopregnanolone 3α , 5α -tetrahydroprogesterone that allosterically modulates the activity of gamma aminobutyric acid type A (GABAA) receptors, thereby causing analgesic, anesthetic, anticonvulsant and anxiolytic effects⁷⁰.

AKR1C Effects on Urinary Metabolites

Several urinary steroid metabolites, like DHEA, androstanediol, 20β-dihydroxycortisone, cortisol, estriol, other estrogens and glucocorticoid metabolites are increased in disorders like polycystic ovary syndrome (PCOS)⁷⁰. The highest increase was found for DHEA, the precursor for both adrenal and ovarian androgens, indicating a pathological mechanism in PCOS that targets both organs and/or overall steroidogenesis⁷¹. One study⁷² reported an increase in the activity of AKR1C1 in women with PCOS, while other studies revealed reduced activities of AKR1C1 and 20β-HSD along with an increase of 3α-HSD activity evaluated by tetrahydrocortisol and α -tetrahydrocortisol conversion to 20α -dihydrocortisol⁷³. Table II lists the urinary metabolites associated with AKR1C1 activity.

Conclusions

Adipose tissue is known to have endocrine properties and synthesize steroid metabolizing enzymes, like AKR1 enzymes, 11β-HSD, aromatase, and 17β -HSD. Adipose tissue is recognized as a substantial site for the action and transformation of steroid hormones. AKR1C enzymes are involved in the inactivation of androgen and progesterone which induces adipogenesis, and accumulation, proliferation, and differentiation of adipocytes. Genetic analyses have identified genes crucial for steroid metabolism that are linked with subcutaneous fat accumulation and lipedema⁷⁴. These steroid-converting enzymes mediate the transformation of specific hormones into other hormones that are significantly involved in the metabolic pathways of adipose tissue. Further studies are required to elucidate the complexity of this enzymatic network and its multiple effects on adipose tissue functions.

Conflict of Interest

The Authors declare that they have no conflict of interests.

Table II. AKR1C1 metabolites with potential clinical relevant

3α -Hydroxy- 5α -pregnan-20-oneAllopregnanolone 3α -Hydroxy- 5β -pregnan-20-onePregnanolone 3β -Hydroxy- 5α -pregnan-20-oneIsopregnanolone 3β -Hydroxy- 5β -pregnan-20-oneEpipregnanolone 3β -Hydroxy- 5β -pregnan-20-oneEpipregnanolone 5α -Pregnane-3,20-dione 5α -Dihydroprogesterone 5α -Pregnane-3,20-dione 5β -Dihydroprogesterone 5β -Pregnane-3,20-dione 5β -Dihydroprogesterone 20α -Hydroxy-pregn-4-ene-3-one 20α -dihydroprogesterone 5α -Pregnane- 3α ,20 α -diolAllopregnanediol 5β -Pregnane- 3α ,20 α -diolPregnanediol 5α -Androstan- 17β -ol-3-one 5α -Dihydrotestosterone 5α -Androstan- 17β -diol 3α -Androstanediol 5α -Androstane- 3α ,17 β -diol 3α -Androstanediol 21 -hydroxy- 5α -pregnan-20-one 5α -Dihydrodeoxycorticosterone 3α ,21-dihydroxy- 5α -pregnan-20-one 3α -Sa-Tetrahydrodeoxycorticosterone 3α ,21-dihydroxy- 5α -pregnanoloneAllotetrahydrodeoxycorticosterone 15 -keto- 13 ,14-dihydrox-PGF2 α PGFM	Molecule	Common name
8-iso-Prostaglandin E2g 8-iso-PGE2g	3α -Hydroxy- 5α -pregnan-20-one 3α -Hydroxy- 5β -pregnan-20-one 3β -Hydroxy- 5α -pregnan-20-one 3β -Hydroxy- 5β -pregnan-20-one 5α -Pregnane-3,20-dione 5β -Pregnane-3,20-dione 20α -Hydroxy-pregn-4-ene-3-one 5α -Pregnane- 3α ,20 α -diol 5β -Pregnane- 3α ,20 α -diol 5β -Pregnane- 3α ,20 α -diol 5α -Androstan- 17β -ol-3-one 5α -androstane- 3α ,17 β -diol 21-hydroxy- 5α -pregnan-20-one 3α ,21-dihydroxy- 5α -pregnan-20-one Pregnanetriol/17-hydroxypregnanolone 15 -keto-13,14-dihydro-PGF2 α &	AllopregnanolonePregnanoloneIsopregnanoloneEpipregnanoloneEpipregnanolone5α-DihydroprogesteroneSβ-Dihydroprogesterone20α-dihydroprogesteroneAllopregnanediolPregnanediolSα-Dihydrotestosterone3α-Androstanediol5α-Dihydrodeoxycorticosterone3a, Sa-Tetrahydrodeoxycorticosterone3a, Sa-TetrahydrodeoxycorticosteroneAllotetrahydrodeoxycorticosteroneAllotetrahydrodeoxycorticosteroneAllotetrahydrodeoxycorticosterone/17-hydroxypregnanolonePGFM&-pGF2α

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