Chapter 16 The Dehydrogenase Hypothesis

Conor Woods and Jeremy W. Tomlinson

Abstract Circulating glucocorticoid (GC) levels are controlled by the Hypothalamo–Pituitary–Adrenal (HPA) axis, but within tissues, GC availability is controlled by the isoforms of 11ß (Beta)-Hydroxysteroid Dehydrogenase 11ß (Beta)-HSD that interconvert inactive cortisone and active cortisol. Two isoforms have been identified; in key metabolic target tissues (including liver and adipose), expression of 11B (Beta)-HSD1 predominates that in vivo converts cortisone to cortisol and thus amplifies local GC action. In contrast, in mineralocorticoid target tissues 11β (Beta)-HSD2 is the isoform that is most abundantly expressed. This inactivates cortisol to cortisone and offers protection for the mineralocorticoid receptor form occupation and activation by cortisol. Dysregulated 11β (Beta)-HSD1 activity has been implicated in many metabolic diseases such as obesity and diabetes and inhibition of 11β (Beta)-HSD1 represents a promising therapeutic target. Mutations within the gene encoding 11ß (Beta)-HSD2 cause the Syndrome of Apparent Mineralocorticoid Excess and decreases in activity are linked to hypertension as well as impairment in placental function and neonatal growth. We will discuss the molecular biology and enzymology of 11β (Beta)-HSD and its role in normal physiology and discuss altered 11^β (Beta)-HSD activity in pathological states and the potential for therapeutic targeting.

Keywords 11 β (Beta)-HSD • Endoplasmic reticulum (ER) • Cortisol • Cortisone • H6PDH • NADPH

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Introduction

Glucocorticoids (GC) (cortisol in man and corticosterone in rodents) are crucial in regulating many important physiological functions including glucose and amino acid metabolism, inflammation, immunity and general health and well-being [1]. GCs bind to the glucocorticoid receptor (GR), a member of the nuclear receptor family and alter gene transcription up-regulating anti-inflammatory protein synthesis and reducing expression of pro-inflammatory cytokines.

Since their discovery in the 1940s by Kendall and Hench, GCs have become a mainstay of therapy in many clinical conditions including rheumatoid arthritis, asthma and as a cornerstone in anti-rejection medication regimes in organ transplant recipients.

Whereas circulating GC levels are regulated by the Hypothalamo-pituitaryadrenal (HPA) axis, tissue cortisol metabolism is under the control of the enzymes of the 11ß (Beta)-Hydroxysteroid Dehydrogenase (11ß (Beta)-HSD) system. Two isozymes exist, namely 11 β (Beta)-HSD1 and 11 β (Beta)-HSD2. In recent decades, attention has focussed on tissue cortisol metabolism and the pre-receptor regulation of GCs within tissues such as fat, muscle and liver [2-6]. 11 β (Beta)-HSD1 is a bidirectional enzyme, which in vivo acts primarily as an oxo-reductase, converting inactive cortisone (and inactive 11-dehydrocorticosterone in rodents) to active cortisol. 11 β (Beta)-HSD1 is located in a number of key metabolic tissues including liver, skeletal muscle, gonads and adipose tissue [6, 7] and has key metabolic influence acting as an amplification step in tissue cortisol exposure. Tissue exposure to GC therefore is not only reliant on circulating levels of cortisol but can also be altered by pre receptor regulation via 11β (Beta)-HSD1. Dysregulated 11β (Beta)-HSD1 activity has recently been implicated in many metabolic and inflammatory diseases and is a promising target to alter tissue glucocorticoid exposure. Selective 11ß (Beta)-HSD1 inhibitors exist and have been trialled in early phase 2 studies primarily in type 2 diabetes [8].

11 β (Beta)-HSD2 is expressed in aldosterone selective tissues, mainly in the distal nephrons, colonic epithelium, salivary and sweat glands (mineralocorticoid receptor (MR) expressing tissues) and in the foetus and placenta during gestation. It acts uniquely as a dehydrogenase deactivating cortisol to inactive cortisone, protecting tissues from excess glucocorticoid exposure [6, 9]. 11 β (Beta)-HSD2 deficiency or inhibition causes hypertension secondary to apparent mineralocorticoid excess whereby cortisol activates the MR for which it shares the same affinity as aldosterone. A similar syndrome manifests in people ingesting large amounts of European licorice which contains an inhibitor of this enzyme. 11 β (Beta)-HSD2 is important in gestation with significant activity in both placenta and fetal brain development by preventing early exposure to GCs.

Hypothalamic Pituitary Adrenal Axis and Cortisol Production

GCs, synthesized from cholesterol precursors, are produced primarily by the adrenal cortex (zona fasiculata) under the influence of the HPA axis in a classical circadian rhythm, regulated by negative feedback.

Healthy adults secrete 10–15 mg s cortisol/day [10]. The majority of cortisol is protein bound to cortisol binding globulin (CBG) although some is albumin bound. Estimates suggest that only 5 % of circulating cortisol is "free" and biologically active [11, 12]. The half-life of free cortisol is only a few minutes whereas protein bound cortisol has a much longer half life of between 70 and 120 min [11, 13, 14]. Cortisone has similar unbound concentrations to that of cortisol but has a lower affinity for both CBG and GR [12, 15]. Importantly the biological availability of GCs represents a balance between synthesis/secretion and metabolism/clearance.

Cortisol Metabolism and Clearance

There are multiple pathways involved in cortisol and cortisone metabolism. These include A-ring reduction to from tetrahydrocortisol and it's 5α (alpha)-isomer allotetrahydrocortisol, hydroxylation to 6β (Beta) hydroxycortisol and reduction of the 20-oxo group giving cortols. Among the various distinct pathways, perhaps the most critical of these is the inter-conversion of cortisol and cortisone mediated by the 11 β (Beta)-HSD isozymes. 11 β (Beta)-HSD interconverts cortisol and cortisone by altering the hydroxyl group at C11 (see Fig. 16.1).

Tissue Cortisol Metabolism: 11β (Beta) HSD1

11β (Beta) HSD1: Discovery

Glucocorticoids were discovered in the 1940s and 1950s and were heralded as a potentially curative treatment for many ailments [16]. Kendall et al. published the discovery of what they believed to be a treatment that could reverse rheumatoid arthritis in the 1950s [17]. What they referred to as Compound E; was actually cortisone, a

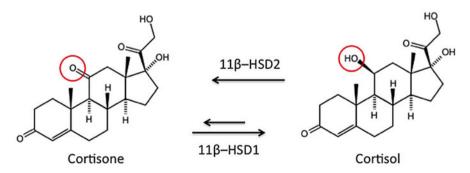


Fig. 16.1 Isoforms of 11β (Beta)-Hydroxysteroid dehydrogenase as pre-receptor regulators of glucocorticoid availability

precursor of cortisol (Compound F). We now know that the cortisone administered, was converted to cortisol by 11 β (Beta)-HSD1. The hydroxyl group at C11 is crucially important for cortisol to be effective. 11 β (Beta)-HSD1 reduces inactive cortisone to active cortisol by adding a hydroxyl group at C11. Thus, unbeknownst to the authors they had actually made two discoveries—the first that the glucocorticoid cortisol has dramatic therapeutic effect on arthritis and second that the enzyme 11 β (Beta)-HSD1 has the ability to convert inactive cortisone to active steroid.

Subsequently to Kendall's paper, cortisol, and not cortisone, was established as the active steroid. It also became apparent that inter-conversion between cortisone and cortisol is possible [18, 19]. This conversion is made possible by the enzyme 11 β (Beta)-HSD1, a member of the dehydrogenase/reductase superfamily. Short-chain Dehydrogenase/Reducatse (SDR) enzymes are NADP(H) dependent enzymes, that are involved primarily in breaking down hormones and chemical messengers [20]. There are more than 3,000 members in the SDR family [7].

Early studies identified 11 β (Beta)-HSD activity in different tissues such as placenta, kidney and liver. However, 11 β (Beta)-HSD did not have the same activity between different tissues. The "set point" of the enzyme varied from tissue to tissue with dehydrogenase activity (cortisol conversion to cortisone) predominating in the placenta and kidney and reductive activity (cortisone conversion to cortisol) predominating in liver [7]. The answer lay in the fact that there are two distinct isozymes of 11 β (Beta)-HSD namely type 1 being predominantly reductive and type 2 acting solely as a dehydrogenase. In addition 11 β (Beta)-HSD type 1 is a bi-directional capable of acting in either direction but dependent on co-factors such as NADPH. *In vivo* in intact tissues the 11 β (Beta)-HSD1 enzyme acts as an oxo-reductase converting cortisone to cortisol. Purified 11 β (Beta)-HSD1 enzyme behaves principally as a dehydrogenase, oxidizing cortisol to cortisone.

Human 11β (Beta)-HSD1

11 β (Beta)-HSD1 was initially purified and cloned from rodents in the 1980s by Carl Monder's group [21, 22]. In humans the gene for 11 β (Beta)-HSD1 (HSD11B1) is located on chromosome 1, is 30 kb in length and has six exons and five introns. It comprises of 292 amino acids and shares 77 % homology with rat amino acid sequence [23]. Human 11 β (Beta)-HSD1, cloned in 2002 [24], exits as a dimer [25] and is bound to the endoplasmic reticulum (ER) with its catalytic domain within the ER lumen [26] (see Fig. 16.2). The catalytic directionality of the enzyme is based on the position of 11 β (Beta)-HSD1 within the ER lumen where it co-localises with Hexose 6 Phosphate Dehydrogenase (H6PD). H6PD generates the reduced co-substrate NADPH. Thus the ratio of NADPH/NADP confers directionality to 11 β (Beta)-HSD1 [27, 28]. Purified 11 β (Beta)-HSD1, in the absence of H6PD, behaves principally as a dehydrogenase, oxidizing cortisol to cortisone. The enzyme 11 β (Beta)-HSD1, has Michaelis Menten (K_m) constants of 1.83±0.06 µm for corticosterone and 17.3±2.2406 µm for cortisol [7].

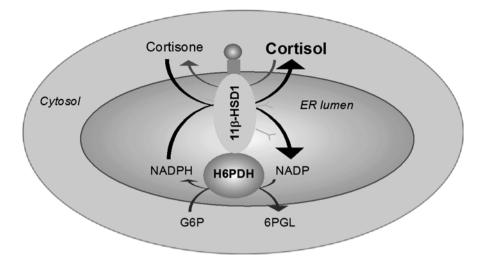


Fig. 16.2 Localization and co-factor dependency of 11β (Beta)-HSD1 within the endoplasmic reticulum

11β (Beta)-HSD1 Ontogeny

The ontogeny of 11 β (Beta)-HSD1 has been studied predominantly in rodents and sheep animal models. The expression of mammalian 11 β (Beta)-HSD1 is predominantly post-natal. In general 11 β (Beta)-HSD1 is detectable in many tissues but there is a lack of activity in early gestation with reductase activity only becoming apparent after delivery and rising steadily throughout infancy [7, 29]. The ontogeny of 11 β (Beta)-HSD1 in humans is not well documented with few published studies. Cortisone therapy is not useful for treating congenital adrenal hyperplasia in early infancy most likely due to absent or significantly reduced liver 11 β (Beta)-HSD1 activity [30]. Both reductase and dehydrogenase activity has been demonstrated in fetal lung tissue [31, 32]. 11 β (Beta)-HSD1 activity remains unchanged throughout childhood in both boys and girls [33]. At puberty there is a reduction in 11 β (Beta)-HSD1 activity in women [33] and this appears to remain. In adults, there is a well described dimorphism in cortisol metabolism between men and women with an apparent reduction in 11 β (Beta)-HSD1 activity in women [34, 35]. However, not all studies have shown this dimorphism [36].

Regulation of 11β (Beta)-HSD1 Expression

11 β (Beta)-HSD1 expression increases steadily until 1 year of age at which time peak levels are reached. 11 β (Beta)-HSD1 is expressed in many tissues including liver, adipose tissue, gonads, GI tract, kidney, eye, anterior pituitary, leukocytes and

bone [7]. 11 β (Beta)-HSD1 expression is highest in liver, brain gonads and adipose tissues [37]. Many factors contribute to alterations in tissue expression of 11 β (Beta)-HSD1. In general glucocorticoids, pro-inflammatory cytokines (TNF α , IL-1 β) peroxisome proliferator-activated receptor γ agonists and CEBPs increase 11 β (Beta)-HSD1 expression whereas Growth Hormone (GH) and liver X receptor agonists inhibit 11 β (Beta)-HSD1 expression [7]. Recently salicylates have been shown to down regulate 11 β (Beta)-HSD1 expression [38] in adipose tissue and improve insulin sensitivity. The effects of sex steroids, insulin and other hormones are variable across tissues and between species. 11 β (Beta)-HSD1 has been shown to be expressed more in adipose tissue in lean women compared to lean men [39]. High dose oestradiol has been shown to repress 11 β (Beta)-HSD1 expression in rat liver and kidney but testosterone did not alter expression [40].

Tissue-Specific Role of 11_β (Beta)-HSD1

Adipose Tissue

11 β (Beta)-HSD1 is highly expressed in human adipose tissue [3, 41]. Broadly speaking, studies have shown similar expression levels of 11 β (Beta)-HSD1 between visceral and subcutaneous compartments [39, 42, 43]. However, H6PD and GR have been shown to have different levels of expression between adipose tissue depots [43].

As in other tissues, 11 β (Beta)-HSD1 acts primarily as a reductase generating active GCs. Its expression and activity are induced by GCs and inflammatory cytokines such as interleukin 1 and TNF α [7, 44–47]. 11 β (Beta)-HSD1 expression and activity increases upon adipocyte differentiation [48] and inhibition of 11 β (Beta)-HSD1 blocks cortisone-induced adipocyte differentiation [49]. 11 β (Beta)-HSD1 activity within adipose tissue depots is controversial with some studies demonstrating increased activity in omental tissue compared to subcutaneous adipose tissue.

Dysregulation of 11 β (Beta)-HSD1 activity has been postulated to be critical in the pathogenesis of metabolic complications associated with obesity. In rodent models increased 11 β (Beta)-HSD1 activity has been demonstrated in visceral adipose tissue of obese rodents compared to wild type [50, 51] although expression in adipose tissue was reduced following high fat feeding in mice [52]. Short term high fat diet reduced the activity of 11 β (Beta)-HSD1 in both subcutaneous and visceral adipose tissue in Wistar rats [53]. Genetically modified rodent models have contributed significantly to our understanding of the role of 11 β (Beta)-HSD1 in adipose tissue. 11 β (Beta)-HSD1 knockout mice are resistant to the metabolic side effects of a high fat diet compared to wild type [54]. Conversely over expression of 11 β (Beta)-HSD1 in adipose tissue leads to weight gain and metabolic complications compared to wild type [55] even on a chow diet.

Clinical studies have yielded varied and sometimes conflicting results. In human adipose tissue, the majority of studies have examined subcutaneous adipose tissue. 11β (Beta)-HSD1 activity in subcutaneous adipose tissue has been shown to be

increased in obese patients compared to non-obese controls [56]. Both 11 β (Beta)-HSD1 activity and expression were shown to correlate positively with BMI by Lindsay et al. in 2003 [57]. Whilst the majority of studies show a positive relationship of 11β (Beta)-HSD1 expression and activity with BMI, this is not the case for all studies. Data with regards to the omental depot are more sparse, but some studies have identified increased expression in women in association with increased omental adiposity [58]. The impact of diet induced obesity on 11β (Beta)-HSD1 activity and expression is not clear. Obese Zucker rats have increased visceral adipose tissue 11β (Beta)-HSD1 expression compared to lean rats [51] however Wistar rats, fed a short term high fat diet show reduced 11 β (Beta)-HSD1 activity [53]. Diet induced obese mice also have reduced expression of 11β (Beta)-HSD1 [59]. In these diet induced obese mice 11ß (Beta)-HSD1 expression is increased by NFk (Kappa) B and reduced by HIF-1. H6PD is also important in the directionality of 11β (Beta)-HSD1 and some studies suggest a role in visceral adipose tissue [60]. Pro inflammatory cytokines and glucocorticoids (increased in obesity) are also known to induce 11β (Beta)-HSD1 expression in adipose tissue [7]. In vitro studies highlight the role of transcription factors (CEBPs) for controlling 11β (Beta)-HSD1 expression in adipocytes [61, 62].

Liver

GCs have key metabolic effects on the liver, augmenting insulin stimulated lipogenesis [63] and reducing lipolysis in hepatic tissue [64]. Indirectly, they increase hepatic lipid accumulation by inducing surrounding adipose tissue lipolysis thus increasing the delivery of free fatty acids to the liver via the portal circulation [65].

11 β (Beta)-HSD1 expression and activity has been extensively studied in hepatic tissue. In animal studies global 11 β (Beta)-HSD1 knock out mice are protected from diet induced hepatic steatosis when fed a high fat diet [54, 55]. Transgenic mice with hepatic 11 β (Beta)-HSD1 overexpression, develop hypertension, hepatic steatosis and dyslipidemia but interestingly do not develop steatohepatitis and minimal insulin resistance [66]. Other factors including HPA axis activation and adipose tissue 11 β (Beta)-HSD1 activity may play a role in the development of hepatosteatitis. Liver specific HSD1-null mice have minimal phenotype (ref) suggesting visceral adipose HSD1 is the key site of action and excess adipose-generated GCs act on the liver via visceral portal drainage.

11 β (Beta)-HSD1 is highly expressed in human liver [23], predominantly in a centripetal pattern histologically with maximum expression around the central vein [67]. Hepatic 11 β (Beta)-HSD1 appears to act exclusively as an oxo-reductase generating active GC [68]. In animal rodent models over expression of 11 β (Beta)-HSD1 in visceral adipose tissue was associated with Non Alcoholic Fatty Liver Disease (NAFLD) [69]. In humans, some studies have demonstrated increased expression of 11 β (Beta)-HSD1 in liver tissue in obese persons with metabolic syndrome [70] however other studies have not confirmed these findings [71, 72]. Liver 11 β (Beta)-HSD1 activity, as measured by urinary steroid metabolite analysis is reduced in obesity compared to non-obese controls [73]. Liver 11 β (Beta)-HSD1 activity (using serum cortisol generation form oral cortisone) is also reduced in

obesity compared to non-obese controls [73]. These reductions in both liver 11 β (Beta)-HSD1 activity and expression contrast with increased activity and expression in adipose tissue in obesity. Ahmed et al. have described a switch in the expression of 11 β (Beta)-HSD1 across the spectrum of liver disease from lower expression of 11 β (Beta)-HSD1 in steatosis to higher levels of 11 β (Beta)-HSD1 expression associated with steatohepatitis [74]. The authors concluded these changes might reflect a response to a more inflammatory phenotype.

Skeletal Muscle

GCs play an important role in protein metabolism. In Cushing's syndrome the effect of excess GCs is clearly seen with myopathy and muscle atrophy. 11 β (Beta)-HSD1 is expressed in skeletal muscle [75, 76], whereas 11 β (Beta)-HSD2 is not. 11 β (Beta)-HSD1 activity has been demonstrated in skeletal muscle [77] and it may have a role in insulin sensitivity and metabolic disease. Increased expression [78] and activity [79] have been demonstrated in rats and humans with type 2 DM.

The exact role of 11 β (Beta)-HSD1 in sarcopaenia remains unexplained. It is thought that increased 11 β (Beta)-HSD1 activity may play a role in allowing increased muscle cortisol exposure. Increased 11 β (Beta)-HSD1 expression in skeletal muscle is associated with reduced muscle strength in older adults compared to younger controls [80].

Skin

Data on tissue cortisol metabolism within skin is only recently becoming an area of interest and investigation. Skin has been shown to be an active site of cortisol production and metabolism [81, 82]. Excess skin exposure to GC's cause skin changes similar to the natural aging process. These include reduced elasticity, reduced collagen and fibroblast numbers, thinning of dermis and epidermis and a general reduction in the repair capacity of skin [83]. Increased exposure to GC's has been postulated as a factor in age related changes, inflammatory and auto-immunity changes seen in skin [84]. It has been postulated that skin changes seen over time are in part as a result of 11β (Beta)-HSD1 activity [82].

Both 11 β (Beta)-HSD1 and 2 are expressed in skin [81, 82, 85]. 11 β (Beta)-HSD2 is expressed in association with the mineralocorticoid receptor on sweat glands however its role (if any) within the dermis and epidermis is debated. In wound healing, 11 β (Beta)-HSD2 expression has been shown to be induced 48 h after tissue injury with subsequent return to basal levels at 96 h [81]. This has been postulated to be a mechanism to reduce local cortisol excess following inflammation.

11 β (Beta)-HSD1 is widely expressed in human and mouse dermis and epidermis [82, 85, 86]. Upon differentiation of keratinocytes 11 β (Beta)-HSD1 expression increases [85], somewhat akin to the changes seen with pre adipocyte differentiation [87]. Interestingly despite reducing levels of expression of 11 β (Beta)-HSD1 in elderly subjects, a paradoxical rise in 11 β (Beta)-HSD1 activity is seen with increasing age in both humans and mice [82]. This gives credence to the concept of age related skin atrophic changes being in part due to increased cortisol exposure secondary to increased 11 β (Beta)-HSD1 activity.

11 β (Beta)-HSD1 has been shown to have a pivotal role in skin repair following injury [88] and tissue remodeling [85]. In mice 11 β (Beta)-HSD1 is contributory to impaired wound healing. Blocking 11 β (Beta)-HSD1 improved wound healing in mice and prevented age induced skin changes [89]. These data suggest that 11 β (Beta)-HSD1 generated local cortisol is critically important in wound healing and in aging skin changes. Inhibitors of 11 β (Beta)-HSD1 inhibitors (topical or oral) may therefore have therapeutic potential.

Cardiovascular System

Both isozymes are expressed in blood vessel walls [90] and heart [91], however oxo reductase directionality (11 β (Beta)-HSD1) predominates in vascular smooth muscle [88]. There is evidence linking 11 β (Beta)-HSD1 activity with atherosclerosis. Mediastinal adipose tissue 11 β (Beta)-HSD1 has been linked with coronary atherosclerosis [92]. The same authors demonstrated increased 11 β (Beta)-HSD1 expression in aortas of obese patients with the metabolic syndrome [93]. 11 β (Beta)-HSD1 inhibition in apoE knockout mice achieved significant reduction atherosclerotic load suggesting a role in plaque formation [94]. Carbenoxolone treatment has been shown to reduce atherosclerosis in mice [95]. 11 β (Beta)-HSD1 knock out mice show improved angiogenesis to infarcted regions possibly through reduced GC regeneration locally [96].

Central Nervous System (CNS)

GCs are required for normal brain development and normal brain function. Excess GCs are associated with alterations in mood, memory and brain function [97, 98]. Both 11 β (Beta)-HSD1 and 2 are expressed in the brain [99–102], but 11 β (Beta)-HSD2 is expressed at lower levels. 11 β (Beta)-HSD1 acts principally as an oxoreductase in brain tissue [103] but interestingly H6PD does not always co-localise with 11 β (Beta)-HSD1 in the CNS [101]. This lack of universal co-localisation suggests other enzymes providing the necessary co factors for 11 β (Beta)-HSD1. Elevated 11 β (Beta)-HSD1 is seen with ageing and is associated with cognitive decline. Altered 11 β (Beta)-HSD1 activity in the CNS is associated with changes in appetite, affective behavior and circadian rhythm [104].

A putative role for 11 β (Beta)-HSD1 in cognitive impairment is postulated along with investigations looking at the link between 11 β (Beta)-HSD1 and human eye disease including thyroid eye disease [105] and glaucoma. 11 β (Beta)-HSD1 inhibitors have been shown to improve cognitive function in elderly persons with T2 DM [106].

Inflammation and Immunity

Glucocorticoids in pharmacological doses are immunosuppressive and produce powerful anti-inflammatory effects [107]. They achieve this by altering gene transcription and altering pro and anti-inflammatory mediators including cytokines and signaling pathways. 11 β (Beta)–HSD1 is believed to play a key role in local inflammation and immune response to stimuli and allergens [108].

11 β (Beta)-HSD1 is expressed on numerous immune cell types in humans including mast cells, mononuclear cells and macrophages, lymphocytes, B cells and dendritic cells [109–112]. 11 β (Beta)-HSD1 is not expressed in non-stimulated monocytes but 11 β (Beta)-HSD1 expression is increased upon differentiation of monocytes cells into macrophages with interleukin 4 and 13 [110]. A further increase in 11 β (Beta)–HSD1 expression is seen when monocytes differentiate into dendritic cells under the influence of interleukin-4 and granulocyte-macrophage colony-stimulating factor. Other pro-inflammatory cytokines do not increase 11 β (Beta)-HSD1. In macrophages, 11 β (Beta)-HSD1 activity increases with exposure to lipopolysaccharide [110]. 11 β (Beta)-HSD1 expression has also been seen in murine lymphocytes and B cells with increases seen when CD4-positive lymphocytes polarize into Th1 or Th2 subsets [111].

Separate to expression, the functional role of 11β (Beta)-HSD1 in immunity remains debated as it is difficult to separate out the specific subsets of immune cells and to be confident of the role (if any) of 11β (Beta)-HSD1. Our understanding of 11β (Beta)-HSD1 expression and activity on human immune cell actions and inflammation is incomplete and warrants further investigation.

There is evidence both in animal models and in human studies that demonstrate that 11 β (Beta)-HSD1 is implicated in immune response to infection and inflammation. In 11 β (Beta)-HSD1 knock out mice defective macrophage phagocytosis of apoptotic neutrophils is seen in peritonitis [113]. KO mice have increased susceptibility to endotoxaemia compared to wild type [114]. Of note, a study in mice demonstrated that in various models of joint and lung inflammation 11 β (Beta)-HSD1 knock out mice developed inflammation earlier, substantially more inflammation and slower resolution of inflammation compared to wild type controls [115]. This study suggests that 11 β (Beta)-HSD1 plays a key role as an anti-inflammatory regulator and raises concerns for drug therapy inhibiting 11 β (Beta)-HSD1.

In humans (and rats) 11 β (Beta)-HSD1 expression is elevated in colitis [116, 117] with a concomitant reduction in 11 β (Beta)-HSD2 expression. Acute exacerbations of inflammatory bowel disease are associated with elevated 11 β (Beta)-HSD1 expression [118] but interestingly 11 β (Beta)-HSD1 activity was elevated in patients who were in remission, suggesting that high local glucocorticoid levels are important in limiting inflammation. In a rat model, chemically induced inflammation increased 11 β (Beta)-HSD1 expression both in colonic tissue and lymphoid tissue [119].

Recently, 11 β (Beta)-HSD1 has been shown to provide tonic inhibition of mast cell deactivation. 11 β (Beta)-HSD1 knock out mice had increased mast cell numbers and a lower threshold for deactivation suggesting that reduced 11 β (Beta)-HSD1 activity increases allergy and anaphylaxis [109].

Bone and Joint

11 β (Beta)-HSD1 is expressed in human bone [120], predominantly in osteoblasts [121].

In rheumatoid arthritis, both 11 β (Beta)-HSD1 and 11 β (Beta)-HSD2 isozymes are expressed in synovial tissue with conflicting reports as to which is the dominant isoform. Global 11 β (Beta)-HSD1 activity is increased in rheumatoid arthritis (as measured by urinary corticosteroid metabolites) compared to non-arthritic controls [122]. Enzyme activity correlates with synovial inflammation severity [123]. In animal models 11 β (Beta)-HSD1 knock out mice develop more severe arthritis and earlier compared to wild type controls [115]and mice treated with 11 β (Beta)-HSD1 inhibitors (Carbenoxolone) developed worse arthritis [124].

Cortisone Reductase Deficiency (CRD) and Apparent Cortisone Reductase Deficiency (ACRD)

Genetic defects in both HSDB1 and H6PD encoding genes demonstrate the effect of alterations in tissue 11 β (Beta)-HSD1 on clinical phenotype. Both Cortisone Reductase Deficiency (CRD) from HSDB1 gene defects and apparent Cortisone Reductase Deficiency (ACRD) from H6PD gene defects show the impact of reduction in tissue 11 β (Beta)-HSD1 activity with low urine cortisol, significantly elevated cortisone with subsequent compensatory increased HPA activity leading to hyperandrogenism, early adrenarche and PCOS in women [7, 125].

CRD was first described in the 1980s. The majority of cases are female and present with clinical and biochemical hyperandrogenism, with males presenting with precocious puberty [7, 37]. CRD has been described as the "human 11 β (Beta)-HSD1 knockout" [7]. The condition is ameliorable to dexamethasone therapy. It shares some clinical and biochemical features and should not be confused with non-classical congenital adrenal hyperplasia.

11 β (Beta)-HSD1 Inhibition in Clinical Studies

Due to evidence that has implicated 11 β (Beta)-HSD1 in the pathogenesis of disease states including obesity, diabetes and the metabolic syndrome, it represents an exciting therapeutic target to limit local GC availability [2, 126]. Many inhibitors of the 11 β (Beta)-HSD enzymes have been described. These include naturally occurring inhibitor compounds such as liquorice derived glycyrrhetinic acid [127], flavanone/hydroxyl flavanones [128], bile acids [129], progesterone metabolites [130] and even coffee [131]. Most of these naturally occurring compounds inhibit both 11 β (Beta)-HSD1 and 11 β (Beta)-HSD2, with subsequent hypertension and hypokalaemia limiting their possible benefits.

Carbenoxolone, a non-selective inhibitor derived from glycyrrhetinic acid, was the first drug to show benefit in human studies [4, 132]. In healthy volunteers it

improved insulin sensitivity and reduced glucose production rates via a reduction in glycogenolysis but not gluconeogenesis in patients with type 2 diabetes [132]. Carbenoxolone has also been shown to reduce local cortisol availability in subcutaneous adipose tissue and inhibits glucocorticoid induced lipolysis [4]. These early small "proof of principal" studies demonstrated that 11 β (Beta)-HSD1 in metabolic disease could be targeted for drug manipulation and importantly that tissue specific effect could be demonstrated, despite a relative lack of specificity of carbenexolone on 11 β (Beta)-HSD1. Several pharmaceutical companies have developed potent selective 11 β (Beta)-HSD1 inhibitors. Indeed 11 β (Beta)-HSD inhibition has become a significant area of investment for many companies [133] and an extensive array of chemical compounds have been patented and reviewed extensively elsewhere [134]. In general, they are highly selective for 11 β (Beta)-HSD1 over 11 β (Beta)-HSD2 and have high potency for inhibition.

A small number of clinical trials have been published with relatively short durations of intervention (all no more than 12 weeks) and have demonstrated improvements in biomarkers including cholesterol profiles, weight, glycaemic control and blood pressure [8, 135, 136]. The first outcome study to be published investigated the addition of an 11 β (Beta)-HSD1 inhibitor (INCN13739) to metformin in patients with type 2 Diabetes. The drug was well tolerated and the 12 week study demonstrated improvements in weight, glycaemic control and lipid profiles in those people that received the inhibitor [135]. Of note, in this trial there was a compensatory increase in the HPA axis with a dose dependent increase in ACTH and subsequent increase in certain androgens. In women there was a small and significant increase in testosterone levels but biologically active testosterone was felt to be unchanged as sex hormone binding globulin also increased. Whilst all blood results remained in the normal reference range these alterations and small elevations in androgens from HPA activation will remain a clinical concern in the long term, especially for women.

In a further study using a different compound (MK0916) again in the setting of type 2 diabetes and metabolic syndrome, modest improvements in glycaemic control with no reduction in fasting glucose levels were seen [8]. Again however there was mild activation of HPA with elevations in adrenal androgen secretion. Overall, the long-term side effects of 11β (Beta)-HSD1 inhibition remain unknown and while the benefits may outweigh any side effects the possible disruption of HPA axis will remain a concern alongside the magnitude of the clinical response.

Tissue Cortisone Metabolism: 11β (Beta)-HSD2

11β (Beta)-HSD2 Discovery

In 1993, Seckl et al. isolated an enzyme with exclusive 11 β (Beta)-HSD dehydrogenase activity from both human placenta and rat kidney [137]. In 1994, Krozoski et al. isolated human 11 β (Beta)-HSD from human kidney [138], identical to the dehydrogenase enzyme found in placenta. This second enzyme was found to be distinct from 11 β (Beta)-HSD1 and was called 11 β (Beta)-HSD2 and is also a member of the SDR family [139]. Both 11 β (Beta)-HSD1 and 11 β (Beta)-HSD2 isozymes are members of the Short-Chain Dehydrogenase/Reductase (SDR) superfamily of enzymes, however each isozyme has a distinct gene sequence with exons found on different chromosomes. There is little similarity or overlap in sequence between isozymes (18 % identity), except for similar co-factor binding regions at the NH₂-terminal [6].

The human 11 β (Beta)-HSD2 gene is located on chromosome 16, has 5 exons and is only 6 kbs in length [140]. Human 11 β (Beta)-HSD2 measures 405 amino acids in length and has a molecular mass of 44 kDa [138]. It is also anchored to the ER and loses its dehydrogenase activity once removed from tissue membranes [141]. 11 β (Beta)-HSD2 universally acts as a dehydrogenase across species [6] and has a Km for cortisol of 50–60 nM and 10–13 nM for cortisone [6, 142]. Mutations in 11 β (Beta)-HSD2 leading to apparent mineralocorticoid excess (AME) and hypertension have been extensively reported [143–145].

11β (Beta)-HSD2 Ontogeny

11 β (Beta)-HSD2 has an important role in fetal development with intra uterine "programming" affecting subsequent adult physiology. 11 β (Beta)-HSD2 plays an important role in gestation in humans and mammals protecting tissues against GC exposure prematurely. 11 β (Beta)-HSD2 is expressed and active in placenta [36, 51] and steadily rises throughout gestation and declines 2 weeks prior to labour [29, 137, 146]. Placenta 11 β (Beta)-HSD2 is localized to the syncytiotrophoblast where it has been described as a barrier to maternal corticosteroid which is considerably more concentrated [6]. GCs play a critical role in the development of fetal organs, in particular, towards the end of pregnancy, in lung tissue. Excess GC exposure in utero is associated with physiological and metabolic complications [147–149]. There is mounting evidence that 11β (Beta)-HSD2 plays a key protective role in normal development of the fetus and in particular brain development [148]. Altered or disrupted 11β (Beta)-HSD2 activity with subsequent excess intra-uterine exposure to glucocorticoid has a "programming" effect on the fetus leading to low birth weight and lifelong physiological consequences such as increased cardiovascular, metabolic and psychiatric complications [150]. This role in the development of the fetus and its role in subsequent lifelong physiology has led some to consider the degree of prenatal GC exposure as a potential prognostic biomarker [6].

Regulation 11 β (Beta)-HSD2 Expression

Unlike 11 β (Beta)-HSD1, there is considerable data published on the epigenetic influence on 11 β (Beta)-HSD2 activity in humans and in rodent models.

The 11β (Beta)-HSD2 gene is susceptible to epigenetic influence, with methylation of the promoter region of particular interest. Increased methylation of this

region has been inversely associated with 11β (Beta)-HSD2 expression and influences the development of hypertension, intrauterine growth, birth weight and neurobehavioural movement [151, 152] in rats. Intrauterine growth retardation has been associated with increased methylation of 11β (Beta)-HSD2 gene promoter with subsequent repression of 11β (Beta)-HSD2 in adult kidneys [153].

Factors that increase 11 β (Beta)-HSD1 expression tend to reduce 11 β (Beta)-HSD2 such as TNF α [154]. Oestrogen increases 11 β (Beta)-HSD2 expression [40, 155]. Vasopressin has been shown to stimulate 11 β (Beta)-HSD2 [156]. Glucocorticoids down-regulate 11 β (Beta)-HSD2 in foetal placenta but not foetal kidney [157]. Dexamethasone up-regulates 11 β (Beta)-HSD2 in lung cells [158]. Hypoxia has been shown to reduce 11 β (Beta)-HSD2 expression [159]. In colonic epithelium, aldosterone increases 11 β (Beta)-HSD2 expression [160].

Tissue-Specific Role of 11β (Beta)-HSD2

CNS

In the adult brain, 11 β (Beta)-HSD2 is expressed in a select few regions [161]. Before birth, 11ß (Beta)-HSD2 is expressed in several additional brain regions, including the thalamus and cerebellum, where this enzyme protects proliferating granule cells from the growth-limiting effects of glucocorticosteroids [162, 163]. In adults, however, mRNA and protein are no longer detectable in these regions. Instead, 11β (Beta)-HSD2 expression is found in just a few small sites in adult mice and rats [161, 164] most prominently in a group of neurons found inside the nucleus of the solitary tract (NTS). NTS neurons with 11β (Beta)-HSD2 expression are the only cells in the brain shown to express both this enzyme and the mineralocorticoid receptor (MR); these "HSD2 neurons" are activated along with salt appetite after dietary sodium deprivation or volume depletion, and may trigger salt appetite. The only other brain sites in which 11β (Beta)-HSD2 expression (mRNA and protein) was identified consistently in adult animals are the subcommissural organ (a circumventricular organ comprised of modified ependymal cells which do not express MR) and a small subdivision of the ventromedial hypothalamic nucleus (neurons that lack MR immunolabeling). Information regarding 11β (Beta)-HSD2 expression remains incomplete for non-neuronal tissues such as the meninges, choroid plexus, ventricular ependyma, and cerebral vasculature.

Cardiovascular System

11 β (Beta)-HSD2 is expressed in vascular endothelium [90]. 11 β (Beta)-HSD2 knockout mice develop endothelial dysfunction [165]. Lack of 11 β (Beta)-HSD2 and MR activation is implicated in generation of severe atherosclerosis in mouse models [166].

Kidney

11 β (Beta)-HSD2 is perhaps best known for its role in the kidney where it protects the mineralocorticoid receptor from excess exposure to GC. 11 β (Beta)-HSD2 is widely expressed in distal nephrons [141]. Although the inherent enzyme ability of 11 β (Beta)-HSD2 to clear cortisol (converting it to cortisone) should not be enough, given concentrations and binding affinities, in reality it protects the mineralocorticoid receptor from GC exposure [6, 167]. Lack of 11 β (Beta)-HSD2 in kidney (AME discussed below), leads to hypertension and other sequelae. 11 β (Beta)-HSD2 activity, measured by urinary metabolite ratios, reduces with age suggesting a role in age related hypertension [168]. Some studies show a role of 11 β (Beta)-HSD2 in hypertension although not all studies are in agreement [169]. In kidney disease reduced 11 β (Beta)-HSD2 activity has been shown in persons with hypertension [170, 171].

Colon

11 β (Beta)-HSD2 is expressed in colonic epithelium [172]. Expression is increased by aldosterone in rats [160]. In Inflammatory bowel disease 11 β (Beta)-HSD2 expression is down regulated in both humans and rats [116]. This is accompanied by an increase in 11 β (Beta)-HSD1 expression and so is presumed to be an attempt to locally control GC exposure to inflamed tissue. Zhang et al. showed that inhibiting 11 β (Beta)-HSD2 reduces colon carcinogenesis by inhibiting COX 2 pathways. The reduction in 11 β (Beta)-HSD2 blocked colorectal adenocarcinoma angiogenesis and metastasis [173].

Salivary Gland and Skin

As mineralocorticoid target tissues, both skin and salivary glands express 11 β (Beta)-HSD2. In the skin expression is mainly restricted to sweat glands [174] 11 β (Beta)-HSD2 is expressed in both parotid and sub-mandibular glands [174, 175]. Measuring salivary cortisone has been postulated a potential biomarker of serum free cortisol [176]. In addition, reduced activity of 11 β (Beta)-HSD2 in sweat glands has also been linked with essential hypertension [177].

Pituitary

In normal anterior pituitary tissue 11 β (Beta)-HSD2 mRNA expression is seen, but immunofluorescence reveals absent 11 β (Beta)-HSD2 isozyme. Interestingly, ACTH secreting tumours induce 11 β (Beta)-HSD2 expression and may in part explain the re-setting of GC feedback control seen in Cushing's disease [99].

Apparent Mineralocorticoid Excess (AME)

AME is a rare clinical disease that presents in early childhood with hypertension, sodium retention, potassium loss, suppressed renin activity and a metabolic alkalosis [178]. This condition was hallmarked by an increased ratio of urinary cortisol to cortisone metabolites as a result of a lack of conversion of cortisol to cortisone. There was also a low level of circulating of mineralocorticoid despite having evidence of apparent excess with metabolic derangement. Usually fatal in childhood adults with AME were discovered and successfully treated with dexamethasone. Physiological replacement with cortisol caused a return of symptoms and signs [179]. Stewart et al. in 1996 demonstrated that AME was caused by a defect in the 11 β (Beta)-HSD2 gene [180] (similar to the effect seen with liquorice ingestion [181]). These and other observations led to the understanding that 11 β (Beta)-HSD2 in the kidney protected the mineralocorticoid receptor from glucocorticoid binding. Under normal circumstances 11 β (Beta)-HSD2 deactivates cortisol to cortisone and thus protects the MR receptor. In AME, with loss of 11 β (Beta)-HSD2 activity unopposed cortisol binds to MR with subsequent clinical mineralocorticoid excess.

The Future of the Dehydrogenase Hypothesis

There is strong evidence that dysregulated 11 β (Beta)-HSD1 activity is involved in many pathological processes including obesity and type 2 diabetes. We have also begun to see early clinical trials that demonstrate clinical benefit in 11 β (Beta)-HSD1 inhibition. However, whether or not these compounds eventually enter the market with a licence to treat diabetes and metabolic disease remains to be seen. However, they may have utility in other conditions including glaucoma, idiopathic intracranial hypertension and low bone mineral density and clinical trials in these areas are ongoing. In addition, recent work based on observations in a patient with Cushing's disease [182] has suggested that 11 β (Beta)-HSD1 may have a role in regulating the phenotype of circulating active GC excess [183]. This not only raises the possibility for the use of 11 β (Beta)-HSD1 inhibitors in the treatment of Cushing's syndrome, but also that basal activity may predict the susceptibility to the adverse effects of exogenous GCs and that 11 β (Beta)-HSD1 inhibitors could ameliorate the adverse effects, without compromise to the desired actions, of therapeutically indicated GCs.

11 β (Beta)-HSD2 has an established role in the regulation of blood pressure and this is highlighted in patients with AME. However, evidence points to an additional role in placental development and function that may have implications for neonatal growth, either directly or through programming. Inhibiting 11 β (Beta)-HSD2 action has been shown to stop colorectal adenocarcinoma spread and development and this warrants further investigation [173]. Importantly though, whilst much attention in recent years has focussed on 11 β (Beta)-HSD1 activity and therapeutic

inhibition, we must not forget that these two enzymes are tightly associated not least of all because the activity of 11 β (Beta)-HSD1 is entirely dependent upon substrate availability (cortisone) that is generated by 11 β (Beta)-HSD2. It is entirely plausible that cortisone availability may represent a rate-limiting step regulating 11 β (Beta)-HSD1 activity through substrate availability and this needs to be further explored [184].

Inhibition of 11 β (Beta)-HSD1 and specifically within tissues is a promising target as a potential disease prevention/modifying pathway. There is strong evidence to date that dysregulated 11 β (Beta)-HSD1 activity is involved in many disease states including obesity and diabetes among many. We have also begun to see early human clinical trials that demonstrate clinical benefit with 11 β (Beta)-HSD1 inhibition. However, as outlined above both the small number, and the short nature of studies to date have not yielded strong robust evidence to use 11 β (Beta)-HSD1 inhibitors. Limited benefits and possible side effects including HPA activation will likely impede and slow the process of these inhibitor compounds from entering phase 3 trials and into clinical use.

One area of possible clinical use, which has not been looked at yet, is to help antagonise the effects of excess glucocorticoid side effects. Published data demonstrate a key role of 11 β (Beta)-HSD1 in contributing to GC side effects in Cushing's [182] and in bone metabolism [184]. 11 β (Beta)-HSD1 global knock out mice treated with excess GC are protected against GC side effects when compared to wild type controls [183]. This suggests that in conditions with excess circulating GC's, it is the re-activation of cortisone to cortisol within tissues by 11 β (Beta)-HSD1, rather than simple cortisol delivery to tissue from the circulation that is the crucial step determining Cushingoid side effects. Data on the role of 11 β (Beta)-HSD1 activity in humans with excess GC's is lacking. GC's are widely prescribed with estimates of 1–2 % [185] of the population taking prescribed steroids for various inflammatory conditions. Despite their efficacy, up to 70 % of patients experience an adverse systemic side-effect profile [186]. Inhibiting 11 β (Beta)-HSD1 activity may play a beneficial role in preventing glucocorticoid side effects therefore making 11 β (Beta)-HSD1 an exciting therapeutic target for patients with Cushing's syndrome.

Lastly, as mentioned previously there is evidence in the literature that 11β (Beta)-HSD1 activity increases with age and is associated with tissue damage and dysfunction. Therefore using tissue specific inhibitors or 11β (Beta)-HSD1 may be a target in conditions such as sarcopaenia and osteoporosis.

Conclusions

Tissue-specific GC metabolism is under the control of the 11 β (Beta)-HSD dehydrogenase/reductase enzymes. 11 β (Beta)-HSD2 exists to primarily protect mineralocorticoid receptor from excess cortisol exposure in tissues such as kidney. 11 β (Beta)-HSD1 exists in many tissues and amplifies tissue exposure to cortisol by

activating cortisol from inert cortisone. Both isoforms have a potent ability to manipulate clinical phenotype entirely independent of circulating GC levels. The complexity of this system and the intricate and finely tuned control that is able to exert at a tissue-specific level to govern ligand access to corticosteroid receptors has highlighted a fundamental shift in our approach. It adds weight to the argument that simple measurement of circulating steroid hormone levels provides an over-simplistic and perhaps misleading view of GC action.

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