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Review

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ABSTRACT

Due to its impact upon health and the economy, the mechanisms that contribute to the pathogenesis of obesity and the metabolic syndrome are under intense scrutiny. In addition to understanding the pathogenesis of disease it is important to design and trial novel therapies. Patients with cortisol excess, Cushing's syndrome, have a phenotype similar to that of the metabolic syndrome and as a result there is much interest the manipulation of glucocorticoid (GC) action as a therapeutic strategy. Intracellular GC levels are regulated by 11 β -hydroxysteriod dehydrogenase (11 β -HSD1) which converts inactive cortisone to cortisol, thereby increasing local GC action. There is an abundance of data implicating 11 β -HSD1 in the pathogenesis of obesity, type 2 diabetes and the metabolic syndrome and 11 β -HSD1 is an attractive therapeutic target. Selective 11 β -HSD1 inhibitors, which do not act upon 11 β -HSD2 (which inactivates cortisol to cortisone) are in development. So far studies have primarily been carried out in rodents, with results showing improvements in metabolic profile. Data are now beginning to emerge from human studies and the results are promising.

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Contents

1.	Introduction	21
2.	GC action	22
3.	The metabolic actions of GCs	22
4.	11β-HSDs and pre-receptor GC metabolism	23
5.	Characterisation of 11β-HSD1 in obesity and insulin resistance	23
	5.1. Rodent models	23
	5.2. Human studies	24
6.	Pharmacological inhibition of 11 β -HSD1	24
	6.1. Rodent studies	24
	6.2. Human studies	24
7.	Conclusions	25
	References	25

1. Introduction

There is an obesity epidemic with 1.5 billion adults worldwide classified as overweight or obese. In obese subjects, standardized mortality ratios are increased in comparison to the lean population [1]. As well as a decreased life expectancy, obesity has a dramatic effect on morbidity, including dyslipidaemia, insulin resistance,

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type 2 diabetes (T2D), hypertension, cardiovascular disease and some cancers [1]. Together this cluster of risk factors comprise the metabolic syndrome. Insulin resistance is a central component and it is hypothesized to drive the disturbances in metabolic homeostasis. As insulin resistance develops, β -cells of the pancreas compensate by producing more insulin, to keep glucose levels within the normal range. T2D occurs when the β -cells cannot increase insulin levels sufficiently to prevent hyperglycaemia. Hyperglycaemia leads to glucose toxicity and further β -cell dysfunction thus escalating the condition [2].

This health issue is not restricted to wealthy nations, with rates of obesity increasing in almost all countries. Over 150 million people worldwide suffer from T2D and it is predicted that this number

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Fig. 1. Schematic representation of glucocorticoid receptor (GR) action. Cortisol binds the GR releasing it from its cytoplasmic complex and allowing it to move into the nucleus. In the nucleus the GR binds specific glucocorticoid receptor elements (GRE) regulating transcription. The GR can also be bound by co-activators and co-repressors, adding a further level of regulation.

will increase dramatically in the future. In addition, there is a worrying increase in the prevalence of childhood obesity and its associated pathologies [3] that include T2D. Due to its impact upon health and the economy, novel therapeutic agents are urgently required.

Patients with cortisol excess, Cushing's syndrome, have a phenotype which shares many aspects with the metabolic syndrome. The similarities include central obesity, impaired glucose tolerance, insulin resistance, T2D and increased cardiac risk of mortality [4]. As a result of these observations there is much interest in the role glucocorticoids (GCs) play in the pathology of obesity, insulin resistance and T2D.

2. GC action

Cortisol release from the adrenal follows a circadian rhythm; levels are at their highest in the morning, just before waking, and then decrease throughout the day. In addition to circadian control, there is a temporary release of cortisol in response to eating, exercise and stress (both physical and psychological). In the circulation, over 90% of cortisol is bound to corticosteroid binding protein. It is only the unbound, free, portion that is able to diffuse into the cell and exert its effects. GCs primarily act by activation of the glucocorticoid receptor (GR) and the regulation of transcription. The GR is a ligand regulated nuclear receptor and a member of the steroid hormone receptor family. The GR is expressed in almost all tissues and its activity is thought to be highly regulated by post-translational modification. Activity has been shown to be regulated by splicing [5], phosphorylation [6], SUMOylation [7] and acetylation and methylation of its chaperone [8]. Upon cortisol binding, the GR moves into the nucleus, binds specific glucocorticoid response elements (GRE) and recruits co-activators and co-repressors which once bound enhance or repress gene transcription [9,10] (Fig. 1). The GR has also been demonstrated to regulate the expression of genes without a GRE [11]. Notably the GR interacts with, and inhibits the activity of, the pro-inflammatory transcription factors NF-KB and AP-1, decreasing the expression of a pro-inflammatory gene profile [12]. In addition, there are non-genomic actions of GCs, these occur within minutes and cannot be disrupted by transcriptional inhibitors. For example, the GC activation of the insulin signalling factor protein kinase B in human vascular endothelial cells [13].

3. The metabolic actions of GCs

GCs are released as part of the stress response and, at least acutely, have a catabolic action making substrates available for mitochondrial oxidation. However chronic systemic GC excess, Cushing's syndrome, leads to an increased fat accumulation, preferentially in the visceral depot, triacylglycerol (TAG) accumulation within the liver, hyperglycaemia and insulin resistance. These physiological and pathophysiological effects of GCs occur due to the action of GCs on a number of tissues.

In the liver GCs enhance glucose output by induction of phosphoenolpyruvate carboxykinase (PEPCK) expression, the rate limiting step of gluconeogenesis, so increasing the *de novo* synthesis of glucose from substrates such as amino acids, fatty acids and lactate. GCs also regulate hepatic lipid metabolism, increasing very low density lipoprotein (VLDL) production rates [14] and TAG synthesis [15].

In the muscle GCs have been shown to directly inhibit insulin action, decreasing insulin stimulated glucose uptake [16,17]. This in combination with increased hepatic glucose output contributes to the hyperglycemia observed with GC excess [17]. GCs also regulate protein metabolism, decreasing amino acid uptake and synthesis via GR mediated inhibition of activating transcription factor 4 (ATF4) expression [18]. ATF4 coordinately turns on expression of genes for amino acid transporters and amino acid biosynthetic enzymes. The impact of this down-regulation is clearly demonstrated by the myopathy of GC excess.

The actions of GCs upon adipose tissue are numerous and complex. They have potent effects to drive pre-adipocyte differentiation [19,20], but there are differential and depot-specific effects upon pre-adipocyte proliferation [19,21]. The effects of GCs upon insulin sensitivity in adipose remains controversial. In some studies [22–25], but not all [23,26], they have been shown to inhibit insulin signalling and glucose uptake. However, GCs appear to work in conjunction with insulin to drive lipogenesis [27] and lipid uptake



Fig. 2. 11β-Hydroxysteroid dehydrogenase type 1 (11β-HSD1) converts the inactive glucocorticoid, cortisone, to the active, cortisol. A closely associated enzyme, hexose-6-phosphate dehydrogenase (H6PDH), generates the co-factor NADPH. Enhanced activity and expression of 11β-HSD1 has been implicated in obesity, metabolic syndrome and type 2 diabetes and modulation of 11β-HSD1 activity is a novel therapeutic target (G6P=glucose-6-phosphate, 6PGL=6-phosphogluconolactonate).

[28–30]. In contrast to this action of enhancing lipid accumulation GCs act, at least acutely, to mobilize lipid at least in part due to an increase in expression of hormone sensitive lipase [31] and adipose triglyceride lipase [32]. GCs also decrease adipose PEPCK expression, the rate limiting step in glyceroneogenesis and fatty acid re-esterification [17,33], which may act to decrease futile fatty acid cycling, further increasing fatty acid output.

GCs also directly impact insulin secretion, where GR over expression specifically in the pancreatic islet decreases insulin secretion [34]. It is thought that they act by both decreasing the expression of GLUT2 in the β -cells [35] and by impairing the efficacy of Ca²⁺⁺ on the secretory process [36].

4. 11β-HSDs and pre-receptor GC metabolism

Circulating GCs are not elevated in patients with Metabolic syndrome [37,38]. However urinary GC metabolites are increased in patients with visceral, but not subcutaneous, obesity [39] indicative of increased cortisol production. Abnormalities in ACTH pulsatile secretion have been observed in obesity; ACTH pulse frequency is increased but amplitude is reduced [40] and there is a reduction in the sensitivity of negative feedback as measured by a dexamethasone suppression test [41]. In addition, viscerally obese patients have a higher cortisol secretion response to ACTH administration [38,40,42].

The availability of GC to bind and activate the GR is controlled by the 11 β -HSD isozymes, 11 β -HSD1 and 11 β -HSD2. Products of separate genes, they have different physiological roles and tissue distribution. 11 β -HSD2 was the first isoform to be identified and is an NAD⁺ dependent dehydrogenase converting the active GC, cortisol, to the inactive, cortisone. It is expressed predominantly in mineralocorticoid target tissues such as the kidney [43] and colon [44]. Cortisol and aldosterone have the same affinity for the GR and the mineralocorticoid receptor (MR) [45] and the role of 11 β -HSD2 in these tissues is to prevent GC activation of the MR [46]. Patients with impaired 11 β -HSD2 activity present with apparent mineralocorticoid excess (AME), where, GCs act as a ligand for the MR causing renal sodium retention and hypertension [47,48].

In contrast, the 11 β -HSD1 isoform is widely distributed, expressed in key metabolic tissues liver, muscle and fat [49]. 11 β -HSD1 is bidirectional, able to act as both an oxoreductase (activating GC) and a dehydrogenase (inactivating GC). However,

in intact cell systems such as hepatocytes [50] and adipocytes [51] activity is predominantly oxoreductive, supported by a higher affinity for cortisone than cortisol [52]. The directionality of 11 β -HSD1 is dependent upon the physiological or developmental status of a particular cell type. In human omental adipose stromal cells, 11 β -HSD1 switches from a dehydrogenase to an oxoreductase upon differentiation [53]. Like the type 2 isozyme 11 β -HSD1 is positioned within the membrane of the ER, however, unlike 11 β -HSD2, its catalytic domain is orientated towards the lumen [54]. The oxoreductase direction of 11 β -HSD1 is due to the high level of the co-factor NADPH present within the ER lumen. This co-factor is generated by the enzyme hexose-6 phosphate dehydrogenase (H6PDH) [55] (Fig. 2). In the liver and adipose of the H6PDH null mouse, 11 β -HSD1 acts predominantly as a dehydrogenase, in contrast to wild type animals [56,57].

5. Characterisation of $11\beta\text{-HSD1}$ in obesity and insulin resistance

As 11β -HSD1 catalyses the conversion of inactive to active GC the tissue specific levels and activity of this enzyme regulate the local, tissue specific, and GC response.

5.1. Rodent models

Recombinant mouse models have added significantly to our understanding of 11β -HSD1 and metabolic disturbance. The 11β -HSD1 knockout mouse has a beneficial metabolic phenotype, resisting diet induced obesity with improved glucose tolerance, insulin sensitivity and decreased plasma FFAs when compared to wild type. Null animals resisted hyperglycaemia, primarily as a result of reduced gluconeogenesis during fasting because of a lack of induction of G6Pase and PEPCK [58–60].

To investigate the effect of tissue specific GC activation upon whole body metabolic homeostasis a number of transgenic (Tg) mice have been generated. 11β -HSD1 was over expressed in adipocytes under the aP2 promoter. These animals had a phenotype comparable to that of central obesity, with increased visceral adiposity and enhanced adipocyte differentiation. Metabolically they were hypertensive, hyperglycaemic, hyperinsulinaemic, glucose intolerant and insulin resistant with raised serum FFAs and triglycerides [61,62]. In a comparative study 11 β -HSD2 was over expressed in adipocytes, leading to adipose specific inactivation of GCs. These animals had decreased fat pad weight and improvements in whole body glucose tolerance and insulin sensitivity [63]. Linked to the apoE promoter, 11 β -HSD1 was over expressed in hepatocytes. The 11 β -HSD1-apoE mouse was hypertensive had dislipidaemia and a fatty liver. In contrast to the 11 β HSD1-aP2 mouse, it was only mildly insulin resistant and did not have altered adipose depot mass [64].

5.2. Human studies

There have been a large number of studies evaluating tissue specific alterations in 11B-HSD1 expression and activity in obesity and insulin resistance. In obesity global 11B-HSD1 activity, as measured by urinary corticosteroid metabolite analysis, is impaired [65]. This predominantly reflects a decrease in hepatic activity and this is thought to be a compensatory mechanism to preserve insulin sensitivity and decrease hepatic glucose output. In contrast this decrease in activity does not occur in type 2 diabetics [66], suggesting that failure to down-regulate 11β-HSD1 activity may contribute to the development of insulin resistance. In the subcutaneous adipose depot studies have predominantly shown 11B-HSD1 expression and activity is positively correlated to obesity and insulin resistance [67-74]. The data from visceral adipose also suggests an increase in expression in obesity [75-79], however, in some studies no association was found [69,80]. Although there have been far fewer studies in muscle, data indicates that 11β-HSD1 expression is increased in myotubes from obese type 2 diabetics, when compared to obese controls [81].

6. Pharmacological inhibition of 11β-HSD1

The most widely used inhibitor in human studies has been carbenoxolone but it is non-selective, inhibiting both 11 β -HSD1 and 11 β -HSD2. Although undoubtedly useful for proof of concept experiments it cannot be used therapeutically due to the hypertensive effects of 11 β -HSD2 inhibition. Many of the major pharmaceutical companies have 11 β -HSD1 programs, however, the published literature on the metabolic consequences of selective 11 β -HSD1 inhibition is predominantly from rodent studies. Some primate studies have been performed and recently data has been released from an early clinical trial in man.

6.1. Rodent studies

There have been numerous rodent studies, using different selective 11β-HSD1 inhibitors. The first selective compound to be described was BVT2733, a benzene sulfonamide which had greater than 200-fold selectivity for inhibition of 11B-HSD1 over 11B-HSD2 [82]. When compared to vehicle, animals administered BVT2733 for 4 days had significant decreases in cholesterol, free fatty acid and triglyceride levels (up to 88%), decrease in food intake (10%) and body weight (5%). There were beneficial changes in carbohydrate metabolism with improved glucose tolerance and insulin sensitivity, and decreases in fasting blood glucose (up to 88%) and insulin (up to 65%). Endogenous glucose production decreased by approximately 60% where hepatic expression of both PEPCK and glucose-6-phosphatase were both decreased. No adverse effects upon liver biochemistry were reported after 7 days of treatment [83,84]. The effectiveness of the BVT2733 compound was compared to thiazolidinedione (TZD) treatment and at the highest doses BVT2733 decreased HbA1c to similar levels observed with Rosiglitazone [85]. Although in rodents BVT 2733 is a potent inhibitor of 11 β -HSD1 (IC₅₀ 96 nM), it is much less effective in humans, with an IC₅₀ of 3341 nM.

Compound 544, an adamantyl triazole, has been tested in several mice models. When administered for 11 days to diet induced obese mice it led to a decrease in body weight and a reduction in fasting insulin, glucose, triglycerides and cholesterol. Similar observations were seen in a mouse model of type 2 diabetes with decreases in fasting insulin, glucose, glucagon, triglycerides and free fatty acids and improvements in glucose tolerance. Interestingly compound 544 treatment slowed atherosclerotic plaque formation in mice with a targeted deletion of apolipoprotein E [86].

Another compound, 2922, was given orally to a mouse model of obesity, insulin resistance, dyslipidemia and atherosclerosis, Ldlr 3KO mice. After 12 weeks of treatment there were improvements in plasma insulin and blood glucose without any changes in circulating lipid profile. As GCs have a potent anti-inflammatory action it has been suggested that the inhibitors may have negative pro-inflammatory effects. In this model 11 β -HSD1 inhibition did not alter circulating pro-inflammatory cytokine levels and did not negatively impact upon atherosclerotic lesion formation [87].

In a study using a 4-heteroarylbicycol[2.2.2]octyltriazole, compound A, rats on high fat, high sucrose diet were treated with the drug for 3 weeks. 11β -HSD1 inhibition impacted beneficially upon lipid metabolism in both the liver and adipose tissue leading to decreased fasting triglycerides and free fatty acids without altering glucose and insulin levels. The lipid content of the liver and brown adipose tissue were decreased, with evidence for increased lipid oxidation. There also appeared to be an adipose depot-specific effect with a decrease in mesenteric fat pad weight, with a concurrent decrease in expression of genes involved in lipogenesis and fatty acid cycling. However, there was no change in size of the more metabolically favorable epididymal fat pad where, in sharp contrast to the mesenteric depot, expression of genes involved in lipogenesis were increased [88,89]. Compound A has also been used in combination with TZD where it further decreased liver lipid levels compared to TZD alone, and tended towards a further decrease in serum lipid levels [90].

In a recent study 4-(phenylsulfonamidomethyl)benzamide, compound 11n, was administered to ob/ob mice. When compared to controls, short term treatment (4 days) led to a 42% reduction in fasting blood glucose and after 8 days a 36% reduction in non-fasting blood glucose. After longer treatment (23 days) there was a 35% decrease in blood glucose, 28% decrease in non-fasting blood glucose and HBA1c was significantly decreased by 0.57% and insulin by 28%, indicating improved glycemic control. There was also improvement in the lipid profile with decreases in serum triglyceride and total cholesterol [91].

6.2. Human studies

In humans, published reports of the metabolic consequences of 11B-HSD inhibition are from the non-selective compound, carbenoxolone. In healthy individuals, carbenoxolone improves whole body insulin sensitivity [92]. In patients with T2D, glucose production rates were decreased principally through a decrease in glycogenolysis, with no apparent effect on gluconeogenesis. In addition, total circulating cholesterol decreased [93]. It has previously been suggested that carbenoxolone was unable to access adipose tissue [74], which is a key target for pharmacological 11β -HSD1 inhibition. However, recent data has shown that it is able to inhibit local cortisol availability in the subcutaneous depot and inhibit GC induced lipolysis [94]. These experiments show that 11β-HSD1 inhibition can, in humans, have beneficial metabolic effects, however, non-selective 11β -HSD inhibition has limited therapeutic use as inhibition of 11β-HSD2 can lead to apparent mineralocorticoid excess, with hypertension, hypokalaemia and fluid retention.

Selective 11β-HSD1 inhibition studies in primates and humans are emerging. Compound PF-915275 has been shown to be an effective 11B-HSD1 inhibitor in both monkeys and humans, as measured by urinary steroid metabolites and from prednisolone generation studies [95,96]. The primate data show a dose dependent decrease in fasting insulin levels after the 8h treatment [95]. In humans, data from Incyte show that their compound, INCB013739, when administered to patients with T2D twice daily for 2 weeks, abolished all conversion of oral cortisone to cortisol. Metabolically, they observed deceased hepatic glucose production rates, without alteration in glucose disposal. Interestingly, data suggests that fasting glucose decreased in the most hyperglycemic patients. In addition, total and LDL cholesterol decreased with no change in HDL-cholesterol or triglyceride levels [97]. The Incyte compound has also been tested in combination with metformin (MET) in T2D patients that were inadequately controlled by MET alone. After 12 weeks of a daily dose the treatment group had lower HbA1c and total cholesterol compared to MET alone, suggesting beneficial effects on both carbohydrate and lipid metabolism. Importantly this study demonstrated that after 12 weeks of treatment the compound was still well tolerated [98].

The results of these studies are encouraging, however, there are further questions that need to be answered in future studies. In normoglycemic mice these compounds are ineffective [84]. In simple obesity 11 β -HSD1 activity is down-regulated and it is possible that these results reflect already basal levels of 11 β -HSD1 expression and activity. Importantly, this down-regulation is not observed in T2D [66] and therefore therapeutic inhibition may be most effective in this group. Inhibition of 11 β -HSD1 in obese individuals and/or those with impaired glucose tolerance may hypothetically reduce the risk of progression of overt T2D and therefore the role of these compounds as agents for disease prevention will need to be considered.

11 β -HSD1 is expressed in pancreatic islets and in isolated rodent islets where inhibition of 11 β -HSD1 decreases local GC regeneration and increases insulin secretion. To date studies of insulin secretion have not been performed in any of the rodent models. However, in the studies published so far fasting insulin levels have been decreased so any impact upon insulin secretion appears to be offset against improvement in insulin sensitivity.

A potential concern of selective 11β-HSD1 inhibition has been the impact upon the hypothalamo-pituitary-adrenal (HPA) axis. Patients with the putative 11β-HSD1-deficient state, apparent cortisone reductase deficiency (ACRD) are unable to activate oral cortisone to cortisol and therefore cortisol clearance is enhanced. As a consequence of this, there is activation of the HPA axis which maintains circulating cortisol levels. Activation of the HPA then drives adrenal androgen excess which is responsible for the phenotype of ACRD (infertility, androgenic alopecia and oligoamenorrhoea in women and precocious puberty in men). Rodents are not a good model for the assessment of adrenal androgens so these issues cannot be thoroughly addressed in rodent models. The human studies to date have used isolated ACTH measurements and therefore provide only limited information. PF-915275 did not significantly increase ACTH levels although there did appear to be a dose dependent increase, but with a high degree of variability [96]. In the two clinical studies with the INCB013739 compound there was no effect on circulating cortisol levels (as expected), but in there was a borderline significant increase in morning ACTH values at 2 weeks (p=0.056). When taken in combination with MET there was a dose dependent increase in morning ACTH by 4 weeks of treatment with no further increase at the twelfth week. Androgens were measured in these patients and there was a dose dependent increase in circulating DHEA-s levels but no alteration in testosterone or androstendione, there was also no change in the levels of sex hormone binding globulin.

7. Conclusions

Phenotypic similarities between obesity and Cushing's syndrome have led to great interest in the potential of therapeutic modulation of GC action. There is a wealth of data that implicates 11 β -HSD1 in the pathogenesis of obesity and insulin resistance and 11 β -HSD1 is an attractive therapeutic target. Selective inhibitors are now available and results are promising with metabolic benefit, most notably in models of insulin resistance and T2D. The available data is principally from rodent studies but emerging data from primates and humans are encouraging, although, more detailed clinical studies are required to address issues of safety and HPA axis activation. It appears that these compounds not only have the potential for insulin sensitisation and glycemic control but also to improve lipid profiles and fat distribution. These multiple actions mean they are an attractive prospect for with T2D suggest added benefit over current therapies.

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