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Reciprocal Regulation of 11 β -Hydroxysteroid Dehydrogenase 1 and Glucocorticoid Receptor Expression by Dexamethasone Inhibits Human Coronary Artery Smooth Muscle Cell Proliferation *In Vitro*

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Abstract

The actions of glucocorticoids are mediated, in part, by 11 β -hydroxysteroid dehydrogenase 1 (11 β -HSD1), which amplifies their effects at the pre-receptor level by converting cortisone to cortisol. Glucocorticoids, such as dexamethasone, inhibit vascular smooth muscle cell proliferation; however, the role of 11 β -HSD1 in this response remains unknown. Accordingly, we treated human coronary artery smooth muscle cells (HCSMC) with dexamethasone (10⁻⁹–10⁻⁶ mol/L) and found that after 72 h, dexamethasone increased 11 β -HSD1 expression (14.16 \pm 1.6 fold, p<0.001) and activity (6.21 \pm 1.2 fold, p<0.001) in a dose- and time-dependent manner, which was dependent upon glucocorticoid receptor (GR) activation and C/EBP β and C/EBP δ signaling. As glucocorticoids are known to negatively regulate GR expression, we examined the effect of decreasing 11 β -HSD1 expression on GR expression. In HCSMC transfected with 11 β -HSD1 siRNA, GR expression was increased; this effect was associated with protein kinase A activation and CREB phosphorylation. To examine the role of 11 β -HSD1 in HCSMC proliferation, we decreased 11 β -HSD1 expression and stimulated cells with platelet-derived growth factor (PDGF) (10 ng/ml). Decreased 11 β -HSD1 expression was associated with increased cell proliferation in the absence of PDGF compared to scrambled control-transfected cells (236.10 \pm 13.11 %, n=4, p<0.001) and this effect was augmented by PDGF. Furthermore, the inhibitory effect of dexamethasone on cellular proliferation was abrogated in 11 β -HSD1 siRNA-transfected HCSMC. Downregulation of 11 β -HSD1 was associated with decreased p27^{kip1} expression and increased phosphorylated retinoblastoma protein, consistent with a proliferative response. These findings suggest that 11 β -HSD1 plays a role in the effects of glucocorticoids on vascular smooth muscle cell phenotype.

Keywords

11 β -hydroxysteroid dehydrogenase 1; dexamethasone; glucocorticoids; glucocorticoid receptor; proliferation; vascular smooth muscle cells

INTRODUCTION

Glucocorticoids have been shown to modulate vascular reactivity, the response to injury, and atherogenesis. These effects are mediated by glucocorticoid binding to, and activation of, cognate intracellular glucocorticoid receptors (GR) [1]. Glucocorticoids limit vascular remodeling and neointimal proliferation following vascular injury in part, by inhibiting vascular smooth muscle cell proliferation [2–6]. Local application of dexamethasone to rat carotid arteries following balloon injury has been shown to abrogate neointimal proliferation compared to untreated vessels [7]. Similar studies performed in mouse femoral arteries using a drug-eluting polymer cuff revealed that this finding was associated with vascular atrophy owing to a decreased number of vascular smooth muscle cells [8]. Mechanistically, dexamethasone has been shown to affect these responses by inducing cell cycle arrest at the G1 phase in vascular smooth muscle cells [9,10].

In mammalian blood vessels, local glucocorticoid concentrations are regulated at the pre-receptor level by the two isozymes of 11 β -hydroxysteroid dehydrogenases (11 β -HSDs), 11 β -HSD1 and 11 β -HSD2 [11]. The 11 β -HSDs are microsomal enzymes of the short-chain alcohol dehydrogenase superfamily that interconvert active glucocorticoids to their inert 11-keto forms [11]. While 11 β -HSD2 is expressed in the vascular endothelium and converts active glucocorticoids to their inactive 11-ketosteroids, 11 β -HSD1 is a low-affinity NADP(H)-dependent, predominant reductase that regenerates active 11 β -hydroxy forms and is expressed mainly in vascular smooth muscle cells [12–14]. Although not previously demonstrated in vascular smooth muscle cells, in adipocytes and transformed cell lines, it has been shown that glucocorticoids up-regulate expression of 11 β -HSD1 through an indirect mechanism that involves members of the CCAAT/enhancer-binding protein (C/EBP) family of transcription factors [15–23]. In this manner, glucocorticoids regulate 11 β -HSD1 expression and, in turn, 11 β -HSD1 serves to amplify the local response to glucocorticoids [12,24–26].

Experimental evidence has linked 11 β -HSD1 to the effects of glucocorticoids on vascular remodeling. Increased 11 β -HSD1 activity resulted in elevated glucocorticoid concentrations and inhibition of angiogenesis in murine injury and repair models while inhibition of 11 β -HSD1 was found to decrease atherosclerosis in ApoE knockout mice through a mechanism that did not involve altering lipid profiles [27,28]. To date, the role of 11 β -HSD1 in mediating the effects of glucocorticoids in vascular smooth muscle cells and the mechanism(s) by which this occurs remains incompletely characterized. In the present study, therefore, we hypothesized that glucocorticoids increase 11 β -HSD1 expression in vascular smooth muscle cells and that 11 β -HSD1 expression was necessary to mediate the response to mitogenic stimuli.

MATERIALS AND METHODS

Cell culture

Human coronary artery smooth muscle cells (HCASMC) (Lonza, Walkersville, MD and Promocell, Heidelberg, Germany) were maintained in smooth muscle cell basal medium (SmBM) supplemented with SmGM SingleQuots (Lonza), without antibiotics, at 37°C in 5% CO₂. Cells were passaged twice weekly using 0.5% trypsin/EDTA, and experiments were performed on cells from passages 2–5. In selected studies, to decrease 11 β -HSD1 expression, HCASMC were transfected with 40 nmol/L Stealth RNAiTM to 11 β -HSD1 mRNA (Invitrogen, Carlsbad, CA) or scrambled control (Invitrogen) using Lipofectamine RNAi MAX (Invitrogen) in OptiMEM media (Invitrogen). After 4 h, transfection media was replaced with full growth media, and experiments were performed after 48 h. To decrease glucocorticoid receptor (GR) expression, HCASMC were transfected with 40 nmol/L Stealth

RNAiTM to glucocorticoid receptor mRNA (Invitrogen) using the same protocol as for 11 β -HSD1 transfection. To decrease C/EBP β and C/EBP δ expression, HCASMC were transfected with 40 nmol/L RNAi to C/EBP β mRNA (Ambion, Foster City, CA) or C/EBP δ mRNA (Ambion), respectively, using the same protocol as for 11 β -HSD1 transfection.

RNA extraction and qRT-PCR

Total RNA from HCASMC were extracted with the RNeasy kit (Qiagen, Germantown, MD), following the manufacturer's instructions. Samples were quantified and checked for purity and quality by A260/A280 measurements. cDNA was synthesized from 1–5 μ g of each total RNA sample with random hexamers using the Superscript III First-Strand Synthesis System for RT-PCR (Invitrogen), according to the manufacturer's instructions. Quantitative real-time PCR (qRT-PCR) was performed on an Applied Biosystems PRISM 7900 HT Sequence Detector (Foster City, CA). TaqMan[®] Gene Expression Assays for 11 β -HSD1, GR, C/EBP β , C/EBP δ , and GAPDH were purchased from Applied Biosystems. PCR products were analyzed using the $\Delta\Delta C_t$ method with GAPDH as the endogenous control. Cycle parameters were 95°C for 15 min followed by 40 cycles at 95°C for 15 sec, 58°C for 1 min, and 72°C for 1 min.

Western immunoblotting

Cells were lysed directly in RIPA buffer (sodium phosphate pH 7.2 10 mM, sodium chloride 150 mM, Triton \times 100 1%, sodium deoxycholate 0.5%, SDS 0.1%) supplemented with protease and phosphatase inhibitors and stored at -80°C prior to analysis. Protein samples (10–35 μ g) were separated on 4–15% SDS-polyacrylamide gels and transferred to PVDF membranes. The membranes were incubated with antibodies to 11 β -HSD1, 11 β -HSD2, mineralocorticoid receptor, anti-C/EBP β (Abcam, Cambridge, MA); GR, C/EBP δ , phospho-protein kinase A, protein kinase A α -subunit, cAMP Response Element Binding Protein (CREB), hexose-6-phosphate dehydrogenase (Santa Cruz, Santa Cruz, CA); or phospho-GR, phospho-CREB, p27^{kip1}, phospho-Retinoblastoma protein (Rb)(Ser 608), (Cell Signaling, Danvers, MA) overnight at 4°C, and visualized using the ECL detection system (Amersham Biosciences), according to the manufacturer's instructions. The membranes were then stripped and reprobed with either a polyclonal rabbit anti- β -actin antibody (Sigma-Aldrich) or a polyclonal rabbit anti- β -tubulin antibody (Santa Cruz). A VersaDoc (BioRad) scanning system was used to quantitate band density.

11 β -Hydroxysteroid dehydrogenase activity assay

11 β -HSD1 activity was measured as previously described [29] with minor modifications. Briefly, HCASMC were incubated with 100 nmol/L cortisone and [1,2(*n*)³H]-cortisone (Amersham) for 8h, after which steroids were extracted with dichloromethane and resolved by thin-layer chromatography on silica plates. Plates were imaged with iodine vapor, cortisone and hydrocortisone were identified and isolated from the plate, and the corresponding radioactivity was determined by scintillation counting (Beckman Coulter). Results were standardized by total cellular protein levels.

Protein Kinase A activity assay

Protein kinase A activity was measured using PepTag[®] Non-Radioactive cAMP-Dependent Protein Kinase Assay (Promega) according to the manufacturer's instructions.

Proliferation Assay

HCASMC were serum starved for 48h in 0.2% fetal calf serum to synchronize the cells. After this time, cells were stimulated with 5% FCS and 10 ng/ml PDGF-BB in the presence or absence of dexamethasone (10^{-7} mol/L). Cell proliferation was determined by measuring

BrdU incorporation into the DNA of dividing cells using the BrdU Cell Proliferation Assay (Calbiochem, San Diego, CA), according to the manufacturer's instructions. Results were standardized by cell number. Cell viability was determined using the CellTiter 96 AQueous Non-Radioactive Cell Proliferation Assay (Promega, Madison, Wisconsin) according to the manufacturer's directions.

Cell cytotoxicity and apoptosis assays

The ratio of dead-to-live cells was determined using the MultiTox-Fluor Cytotoxicity Assay (Promega), and caspase 3, 7 activity was quantified using the Caspase-Glo 3/7 Assay (Promega), according to the manufacturer's instructions.

Murine aorta isolation, immunohistochemistry, and immunofluorescence

All procedures were approved by the Institutional Animal Care and Use Committee at Brigham and Women's Hospital and Harvard Medical School. Aortas were harvested from C57Bl/6 male mice (Charles River) age 12–16 weeks. The vessels were irrigated with ice-cold phosphate-buffered saline until free of blood, dissected free of adventitia, and cut into 5 mm rings. The rings were placed in smooth muscle cell basal medium (Lonza) and incubated at 37°C in 5% CO₂. To decrease 11βHSD-1 expression, 80 nmol/L Stealth RNAi™ to 11β-HSD1 mRNA (Invitrogen, Carlsbad, CA) or scrambled control (Invitrogen) and Lipofectamine RNAi MAX (Invitrogen) was added to the medium. At selected time points, the aortas were embedded in paraffin and 5 μm sections were analyzed by hematoxylin & eosin staining, immunohistochemistry or immunofluorescence as described previously [30,31] using antibodies to 11β-HSD1 (abcam) and Ki67 (BD Biosciences), respectively.

Data analysis

All experiments were performed a minimum of three times at least in duplicate and data presented are expressed as mean ± standard error of the mean (SEM). The statistical significance of differences was determined by ANOVA followed by Bonferroni's pairwise comparison using GraphPad Prism 3.0 software (GraphPad Software, Inc, La Jolla, CA). A p value of <0.05 was considered significant.

RESULTS

Dexamethasone increases 11β-HSD1 expression and activity

To determine the effect of dexamethasone on 11β-HSD1 expression in HCSCMC, we first assessed steady-state levels of 11β-HSD1 mRNA. Dexamethasone increased 11β-HSD1 mRNA in a dose-dependent manner with the maximum effect observed at 10⁻⁷ mol/L (6.5 ± 0.7-fold increase at 18 h vs. control, n=3, p<0.01) (Fig. 1a). As we saw no further increase in 11β-HSD1 mRNA levels with higher concentrations of dexamethasone, we elected to perform experiments on HCSCMC treated with dexamethasone 10⁻⁷ mol/L. Dexamethasone also increased 11β-HSD1 mRNA in a time-dependent manner with the maximum effect observed at 72 h (116 ± 36-fold increase vs. control, n=3, p<0.01) (Fig. 1b); no further increase in mRNA levels were observed at 96 h. This finding was associated with an increase in 11β-HSD1 protein levels (14.6 ± 1.6-fold increase vs. control, n=3, p<0.001) (Fig. 1c) and 11β-HSD1 reductase activity (Fig. 1d). We found that under these treatment conditions, there was no effect of dexamethasone on 11β-HSD2 or hexose-6-phosphate dehydrogenase protein expression (Supplementary Fig. 1a, 1b).

The glucocorticoid receptor, C/EBP β , and C/EBP δ modulate 11 β -HSD1 expression

To examine the role of glucocorticoid receptor (GR) activation on the dexamethasone-mediated increase in 11 β -HSD1 expression, we first examined GR expression and phosphorylation. While dexamethasone treatment for 24 h decreased total GR protein levels by $45.3 \pm 8.6\%$ compared to control cells ($n=3$, $p<0.01$) (Fig. 2a), phosphorylation at Ser 211 was increased by $52.4 \pm 17.3\%$ compared to control ($n=3$, $p<0.05$) (Fig. 2b), indicating activation of the receptor. This increase in phospho-GR was transient, peaking at 24 h. In contrast, we found that under these conditions, dexamethasone had no effect on MR expression in HCSMC (Supplementary Fig. 1c).

We next examined the effect of GR activation on expression of members of the CCAAT/enhancer-binding protein (C/EBP) family of transcription factors, as these have been shown to modulate 11 β -HSD1 enzyme expression in other cell types [20–23]. Dexamethasone increased C/EBP β ($53.2 \pm 11\%$, vs. control, $n=3$, $p<0.05$) (Fig. 2c) mRNA levels and transiently increased C/EBP β protein levels ($80.2 \pm 4.7\%$ vs. control, $n=3$, $p<0.01$) (Fig. 2d). Dexamethasone also increased C/EBP δ mRNA levels ($241.1 \pm 19\%$ vs. control, $n=5$, $p<0.001$) (Fig. 2e), as well as C/EBP δ protein levels ($223.9 \pm 6\%$ vs. control, $p<0.001$) (Fig. 2f).

Next, to demonstrate that GR activation was necessary for the observed increase in 11 β -HSD1 expression, we transfected HCSMC with an siRNA to GR mRNA to decrease protein expression by 75.6% ($n=3$, $p<0.001$) compared to scrambled control-transfected cells, and examined the effect of GR downregulation on 11 β -HSD1 expression. When HCSMC transfected with GR siRNA were treated with dexamethasone, we found a 46.5-fold ($n=3$, $p<0.001$) decrease in 11 β -HSD1 mRNA (Fig. 3a) and an 8.7-fold decrease ($n=3$, $p<0.001$) in 11 β -HSD1 protein levels (Fig. 3b) compared to scrambled control-transfected cells treated with dexamethasone. These data indicate that GR activation is necessary for the dexamethasone-dependent increase in 11 β -HSD1 expression.

Similarly, to determine if the transcription factors C/EBP β or C/EBP δ , which are activated by the GR, were necessary for the observed increase in 11 β -HSD1 expression, we first transfected HCSMC with an siRNA to C/EBP β to decrease C/EBP β protein levels by 71.5% ($n=3$, $p<0.001$) compared to scrambled control-transfected cells. Downregulation of C/EBP β attenuated the effect of dexamethasone on 11 β -HSD1 mRNA (4.6-fold, $n=3$, $p<0.001$) and protein levels (7.4-fold, $n=3$, $p<0.001$) compared to scrambled control-transfected cells (Fig. 3c, 3d). We next transfected HCSMC with an siRNA to C/EBP δ , which decreased C/EBP δ protein expression by 80.3% ($n=3$, $p<0.01$), and found that downregulation of C/EBP δ inhibited the dexamethasone-mediated increase of 11 β -HSD1 mRNA (2.1-fold, $n=3$, $p<0.001$) and protein levels (2.0-fold, $n=3$, $p=0.01$) compared to scrambled control-transfected cells (Fig. 3c, 3d). Together, these data indicate that both C/EBP β and C/EBP δ are necessary to increase 11 β -HSD1 mRNA and protein levels in response to dexamethasone.

11 β -HSD1 is involved in the regulation of GR expression

As GR expression is negatively regulated by its ligand, and 11 β -HSD1 activity is an important pre-receptor mechanism to generate GR ligand, we also examined the effect of decreasing the expression of 11 β -HSD1 on GR expression. When we transfected HCSMC with an siRNA to 11 β -HSD1 to decrease protein expression by 64% ($p<0.01$), we found that compared to scrambled control-transfected cells, downregulation of 11 β -HSD1 resulted in an increase in GR protein expression (Fig. 4a). Furthermore, while dexamethasone decreased GR protein expression in scrambled control-transfected cells, this response was abrogated in 11 β -HSD1 siRNA-transfected HCSMC suggesting that 11 β -HSD1 plays a role in the dexamethasone-mediated downregulation of GR expression (Fig. 4a).

The cyclic AMP response element binding protein (CREB) has been shown to regulate transcription of the GR gene in response to glucocorticoid levels [26]. To determine if 11 β -HSD1 modulated GR expression by influencing activation of CREB, we first examined expression and activity of protein kinase A, which is known to phosphorylate and activate CREB. In scrambled control-transfected HCSMC, dexamethasone decreased phosphorylation of protein kinase A, which was associated with a decrease in protein kinase A activity (Fig. 4b, 4c). Interestingly, 11 β -HSD1 siRNA-transfected HCSMC demonstrated increased protein kinase A phosphorylation and activity compared to scrambled control-transfected cells, and this effect was not influenced significantly by dexamethasone (Fig. 4b, 4c). In line with these findings, dexamethasone treatment decreased CREB phosphorylation in scrambled control-transfected cells, while HCSMC with decreased 11 β -HSD1 expression demonstrated increased CREB phosphorylation (Fig. 4d). These findings suggest that 11 β -HSD1 expression influences CREB activation as a mechanism by which to regulate GR expression.

Role of 11 β -HSD1 in PDGF-stimulated HCSMC proliferation

Glucocorticoids have been shown to regulate vascular smooth muscle cell proliferation; however, the role of 11 β -HSD1 in this process remains incompletely characterized. We, therefore, examined the effect of dexamethasone on HCSMC proliferation in cells with normal 11 β -HSD1 expression and decreased 11 β -HSD1 expression. After 48 h of serum starvation, HCSMC were stimulated with 5% serum and recombinant human PDGF-BB (10 ng/ml) in the presence or absence of dexamethasone (10^{-7} mol/L). In scrambled control-transfected HCSMC, PDGF-BB increased proliferation as determined by BrdU incorporation (100.38 ± 14.72 vs. 581.32 ± 11.87 arb. units/cell number, $n=4$, $p<0.001$), while the addition of dexamethasone to PDGF-BB-stimulated cells decreased cell proliferation (253.42 ± 7.15 arb. units/cell number, $n=4$, $p<0.001$). In 11 β -HSD1 siRNA-transfected HCSMC, there was an increase in proliferation compared to scrambled control-transfected cells under basal conditions (100.38 ± 14.72 vs. 237.10 ± 12.72 arb. units/cell number, $n=4$, $p<0.001$). While PDGF-BB stimulation increased proliferation significantly in 11 β -HSD1 siRNA-transfected HCSMC (237.10 ± 12.72 vs. 734.03 ± 29.37 arb. units/cell number, $n=4$, $p<0.001$), the addition of dexamethasone had a somewhat attenuated effect in inhibiting proliferation (734.03 ± 29.37 vs. 437.07 ± 36.83 arb. units/cell number, $n=4$, $p<0.001$) compared to what was observed in scrambled control-transfected cells (Fig. 5). To demonstrate the potential *in vivo* relevance of these findings, we harvested aortas from C57Bl/6 wild-type mice ($n=12$), sectioned the vessels to 5 mm rings, transfected them with siRNA to 11 β -HSD1 to decrease protein expression by 52% ($p<0.01$) or a scrambled control (Fig. 6a), and maintained them in culture for up to 72 h. In selected sections, we stimulated the rings with recombinant rat PDGF-BB (10 ng/ml) in the presence or absence of dexamethasone (10^{-7} mol/L). Following stimulation with PDGF-BB, co-incubation with dexamethasone decreased cell proliferation, determined by the average number of Ki67 positive fluorescent cells/10 high powered fields (hpf), in scrambled control-transfected aortas (18.21 ± 5.45 vs. 7.67 ± 3.66 cells/hpf, $p<0.01$). Similar to what was observed with HCSMC, in PDGF-stimulated aorta sections transfected with 11 β -HSD1 siRNA that were co-incubated with dexamethasone, there was an attenuated effect in inhibiting proliferation (30.11 ± 3.87 vs. 24.55 ± 4.33 cells/hpf, $p=0.07$) (Fig. 6b).

We next sought to determine a mechanism by which to explain the observed differences in proliferation between HCSMC with normal levels of 11 β -HSD1 expression and those with decreased 11 β -HSD1 expression. We found that there was no difference between the ratio of dead-to-live cells between scrambled control-transfected HCSMC and those with decreased 11 β -HSD1 expression (0.93 ± 0.08 vs. 0.93 ± 0.15 , $n=4$, $p=NS$), and we found no difference in apoptosis between the two groups (17.54 ± 6.80 vs. 17.53 ± 4.68 relative light

units, n=4, p=NS). These results suggest that the observed differences in cell proliferation could not be attributed to an increase in cell death or apoptosis.

Next, we examined expression of p27^{kip1} and its downstream target phosphorylated retinoblastoma (Rb) protein as both of these regulate the G1-phase of the cell cycle, are subject to control by glucocorticoids, and have been implicated in mediating vascular smooth muscle cell proliferation. In accordance with previously published reports, we found in scrambled control-transfected cells that PDGF-BB decreased p27^{kip1} expression and increased phosphorylation of Rb at Ser608, consistent with progression through the cell cycle and proliferation; the addition of dexamethasone to PDGF-stimulated cells increased p27^{kip1} expression and decreased phosphorylation of Rb, indicating a cell cycle block and inhibition of proliferation. In contrast, in 11 β -HSD1 siRNA-transfected HCSMC, p27^{kip1} expression was decreased and phosphorylation of Rb was increased compared to scrambled-control transfected cells under basal conditions. Here, stimulation with PDGF resulted in a modest, further decrease in p27^{kip1} expression and the addition of dexamethasone resulted in a minor increase in p27^{kip1} expression and corresponding decrease in phospho-Rb (Fig. 7a, 7b). Taken together, these data demonstrate that 11 β -HSD1 participates in glucocorticoid-mediated inhibition of HCSMC proliferation in response to PDGF, in part, by modulating p27^{kip1} expression and Rb phosphorylation to facilitate the cell cycle block at G1.

DISCUSSION

In these studies, we found that dexamethasone increases 11 β -HSD1 expression and activity in HCSMC in a GR-dependent manner to regulate cell proliferation through p27^{kip1} and Rb. We also identified a role for 11 β -HSD1 in the regulation of GR expression. There are several prior reports that demonstrate that glucocorticoids increase 11 β -HSD1 expression in non-vascular cells and one study showing an increase in 11 β -HSD1 mRNA levels in rat aortic smooth muscle cells [32,33]. Our studies now extend these observations to human vascular smooth muscle cells. Furthermore, we demonstrate that 11 β -HSD1 expression in response to glucocorticoids is dependent upon GR activation as evidenced by the lack of increase in 11 β -HSD1 expression in HCSMC transfected with an siRNA to GR.

The time course of upregulation of 11 β -HSD1 expression observed in our studies is consistent with indirect regulation of the 11 β -HSD1 promoter by glucocorticoids, paralleling previous reports in human fibroblasts and in A549 cells [15,23]. The transcriptional activity of glucocorticoids has been linked to the C/EBP family of transcription factors that interact with either the GR or target gene promoters to promote transcription. In addition, C/EBPs, primarily C/EBP β , mediate the effect of glucocorticoids by inducing transcription of genes that lack GR binding sites [34–36]. We find that C/EBP β , and to a lesser extent C/EBP δ , are important for the dexamethasone-mediated increase in 11 β -HSD1 expression in HCSMC. Our findings are in accord with previously published studies that established the central role of C/EBPs in the regulation of 11 β -HSD1 transcription in liver, lung, and adipose cells [20–23].

It is also recognized that glucocorticoids negatively regulate expression of the GR [37,38]. In our studies, we demonstrate 11 β -HSD1 mediates dexamethasone-mediated downregulation of GR expression in HCSMC. Here, we showed that decreased 11 β -HSD1 expression, which would decrease levels of active glucocorticoids, was associated with an increase in GR expression compared to control cells and cells exposed to dexamethasone, indicating that 11 β -HSD1 plays a role in the regulation of GR expression under ambient conditions. It has been demonstrated that GR expression is controlled at the transcriptional level by the phosphorylation state of CREB [39]. We found that downregulation of GR was associated with decreased CREB phosphorylation in cells with normal levels of 11 β -HSD1.

Interestingly, we also observed that when we decreased 11 β -HSD1 expression, which resulted in an increase in GR expression, there was an associated increase in phospho-CREB levels. We further examined protein kinase A activation as a mechanism of CREB phosphorylation, as a prior study demonstrated that GR transcription is dependent on protein kinase A signaling [40]. Here, we found that increased activation of protein kinase A was associated with the increase in CREB phosphorylation.

As glucocorticoids modulate vascular smooth muscle cell phenotype, we examined the role of 11 β -HSD1 in regulating the effect of dexamethasone on vascular smooth muscle cell proliferation. While prolonged exposure to glucocorticoids inhibits proliferation of cultured vascular smooth muscle cells by inducing a cell cycle arrest at the G1 phase [9,10,41,42], the role of 11 β -HSD1 in mediating this effect is less well defined. In prior studies performed in an osteosarcoma cell line transfected with 11 β -HSD1, expression of 11 β -HSD1 was associated with decreased cell proliferation that was attributed to metabolism of endogenous glucocorticoids to increase cortisol levels [43]. This observation supports our findings that HCSMC with decreased 11 β -HSD1 expression demonstrate increased proliferation under basal conditions and in response to PDGF as compared to cells with normal 11 β -HSD1 expression. The diminished inhibitory effect of dexamethasone in cells with decreased 11 β -HSD1 expression suggests that endogenous metabolism of glucocorticoids is involved in modulating cell proliferation. Furthermore, the translational relevance of these findings was confirmed *ex vivo* in murine aortas with decreased 11 β -HSD1 expression.

The precise mechanism by which glucocorticoids inhibit cell proliferation is not yet known; however, it has been shown that glucocorticoids influence the expression of cyclin-dependent kinases and cyclin-dependent kinase inhibitors, including p27^{kip1} [10]. Accordingly, we investigated the role of 11 β -HSD1 in modulating the expression of p27^{kip1} and the phosphorylation state of Rb. Rb, which lies downstream of p27^{kip1}, has also been linked to the inhibitory effects of dexamethasone on vascular smooth muscle cell proliferation [10,41,44]. Consistent with our findings on the effects of 11 β -HSD1 expression on proliferation, we found that decreased 11 β -HSD1 expression was associated with a downregulation of the cyclin-dependent kinase inhibitor p27^{kip1}, leading to increased phosphorylation of Rb to promote exit from G1 and cell proliferation.

Taken together, our findings indicate that glucocorticoids stimulate a “feed-forward” mechanism to enhance local generation of active glucocorticoids by upregulating the expression and activity of 11 β -HSD1 in HCSMC. This increase in 11 β -HSD1 expression further serves to regulate the actions of GR activation by modulating GR expression through a protein kinase A–CREB-dependent mechanism. Although this may seem counterintuitive, such a signaling mechanism may serve to redirect glucocorticoids to activate the mineralocorticoid receptor in cells and tissues that don’t express 11 β -HSD2. In vascular smooth muscle cells, evidence exists that this mechanism is operative [45]. In HCSMC, we also demonstrate that 11 β -HSD1 expression is also necessary to mediate cellular proliferation under basal and growth factor-stimulated conditions. These findings suggest that 11 β -HSD1 may serve as a target for therapeutic intervention in vascular disease states characterized by excess vascular smooth muscle cell proliferation, such as restenosis. In fact, selective 11 β -HSD1 inhibition has already been shown to limit atherosclerotic neointima formation, and 11 β -HSD1^{-/-} mice have a cardioprotective phenotype that is characterized by increased angiogenesis following injury [27,28]. Our findings suggest further that systemic administration of novel 11 β -HSD1 inhibitors that are currently under investigation for obesity and diabetes may have effects on the vasculature [28,46–49].

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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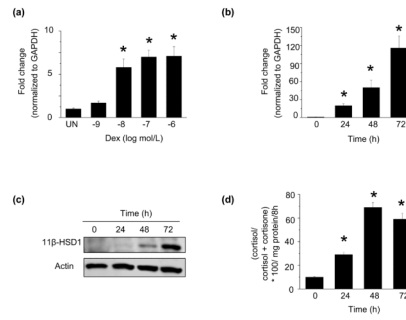


Fig. 1. Dexamethasone increases 11β-HSD1 expression and activity

Human coronary artery smooth muscle cells (HCSMC) (a) were treated with increasing concentrations of dexamethasone and 11β-HSD1 mRNA levels were determined by qRT-PCR after 18 h (n=3). HCSMC were treated with dexamethasone (10^{-7} mol/L) for 0–72 h and 11β-HSD1 (b) mRNA levels were determined by qRT-PCR (n=3), (c) protein expression was examined by Western immunoblotting (n=3), and (d) 11β-HSD1 activity was measured. (n=3). Data are presented as mean ± SEM. A representative blot is shown. *p<0.01 vs. UN, 0. UN, treated; Dex, dexamethasone

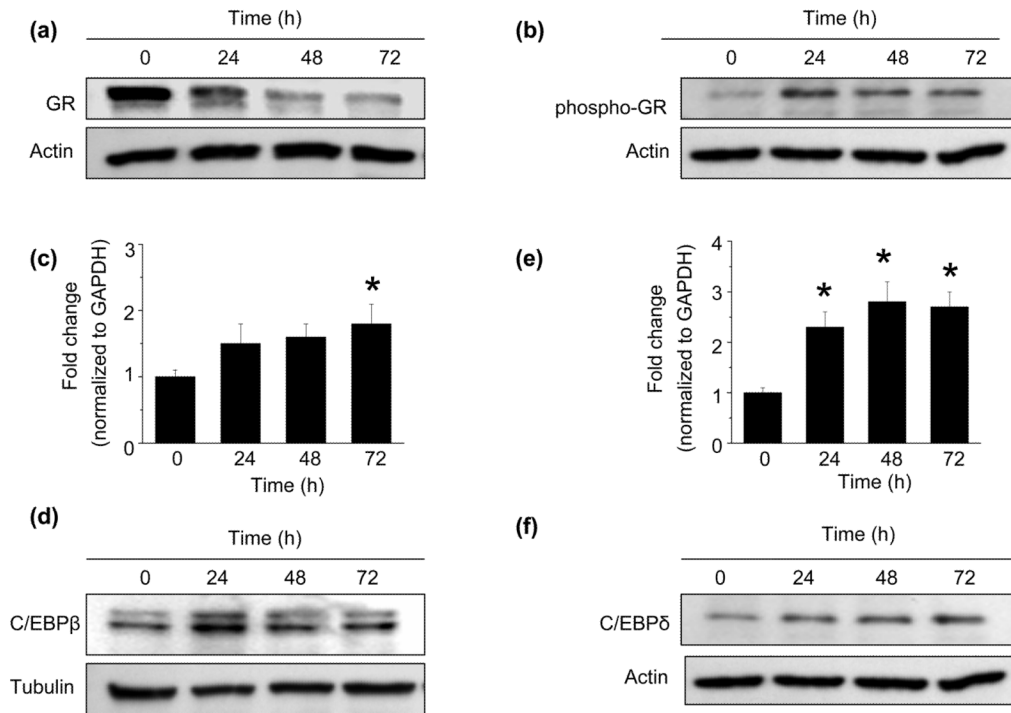


Fig. 2. The effect of dexamethasone on glucocorticoid receptor, C/EBPβ and C/EBPδ expression
 Human coronary artery smooth muscle cells (HCSMC) were treated with 10^{-7} mol/L dexamethasone for 0–72 h and expression of (a) the glucocorticoid receptor (GR) (n=3) and (b) the phospho-GR (n=3) were determined by Western immunoblotting. At the same timepoints, (c) C/EBPβ and (d) C/EBPδ mRNA levels were determined by qRT-PCR (n=3) and corresponding (e) C/EBPβ and (f) C/EBPδ protein levels were assessed by Western blotting (n=3). Data are presented as mean ± SEM. Representative blots are shown. *p<0.05 vs. 0

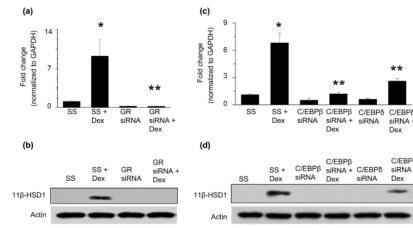


Fig. 3. The glucocorticoid receptor, C/EBPβ, and C/EBPδ regulate the dexamethasone-mediated increase in 11β-HSD1 expression

Human coronary artery smooth muscle cells (HCSMC) were transfected with an siRNA to decrease glucocorticoid receptor (GR) expression or a scrambled control (SS), treated with dexamethasone (10^{-7} mol/L), and (a) 11β-HSD1 mRNA levels were determined by qRT-PCR (n=3) and (b) protein expression by Western immunoblotting (n=3). HCSMC were transfected with an siRNA to decrease C/EBPβ or C/EBPδ expression, respectively, or a scrambled control, treated with dexamethasone (10^{-7} mol/L), and (c) 11β-HSD1 mRNA levels were determined by qRT-PCR (n=3) and (d) protein expression by Western immunoblotting (n=3). mRNA and protein levels were determined. Data are presented as mean ± SEM and representative blots are shown. *p<0.001 vs. SS, **p<0.001 vs. SS + Dex

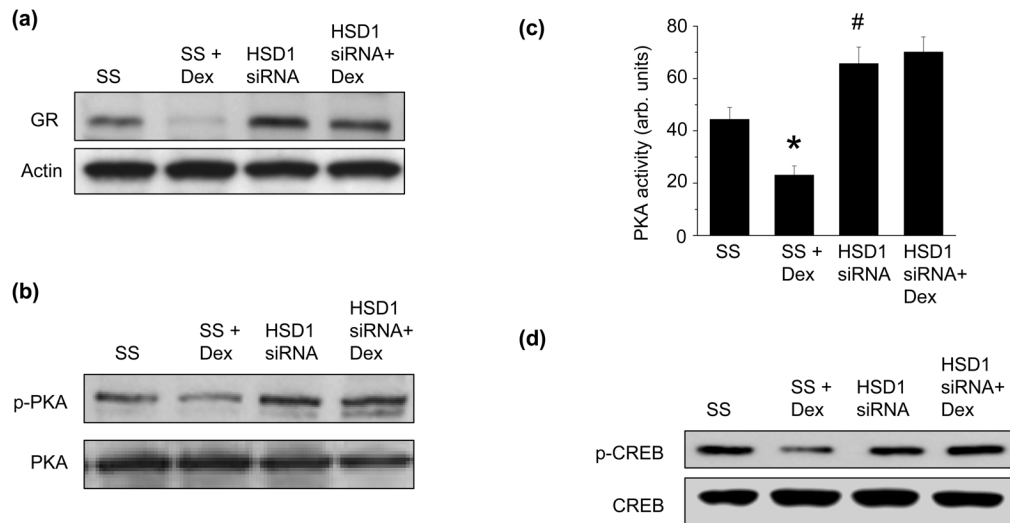


Fig. 4. 11β-HSD1 regulates glucocorticoid receptor expression

HCSMC were transfected with an siRNA to decrease 11β-HSD1 expression, treated with dexamethasone (10^{-7} mol/L), and glucocorticoid receptor (GR) (a) protein expression was determined by Western immunoblotting (n=3). (b) Protein kinase A (PKA) and phospho-PKA expression were assessed by Western blotting (n=3), and (c) PKA activity was measured (n=4). (d) Expression of the cyclic AMP response element binding protein (CREB) and phospho-CREB were assayed by Western blotting (n=3). Data are presented as mean ± SEM and representative blots are shown. *p<0.001 vs. SS; **p<0.001 vs. HSD1 siRNA; #p<0.05 vs. SS

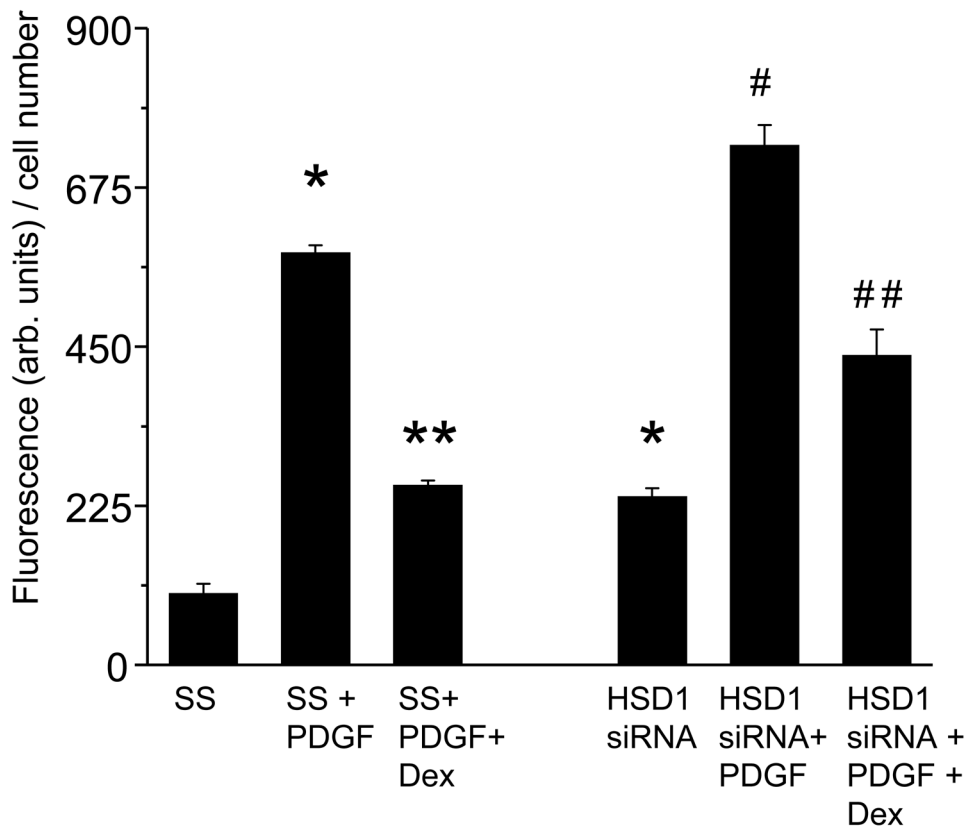


Fig. 5. 11β -HSD1 mediates the inhibitory effects of dexamethasone on PDGF-stimulated HCSMC

Human coronary artery smooth muscle cells (HCSMC) were transfected with an siRNA to decrease 11β -HSD1 expression, synchronized by serum starvation (0.2% serum) for 48 h, and stimulated with 5% fetal calf serum and platelet-derived growth factor-BB (PDGF) (10 ng/ml) in the presence or absence of dexamethasone (10^{-7} mol/L). Cell proliferation was determined by measuring BrdU incorporation (n=4). Data are presented as mean \pm SEM and results are corrected for cell number. * $p < 0.001$ vs. SS; ** $p < 0.001$ vs. SS + PDGF; # $p < 0.001$ vs. HSD1 siRNA; ## $p < 0.001$ vs. HSD1 siRNA + PDGF

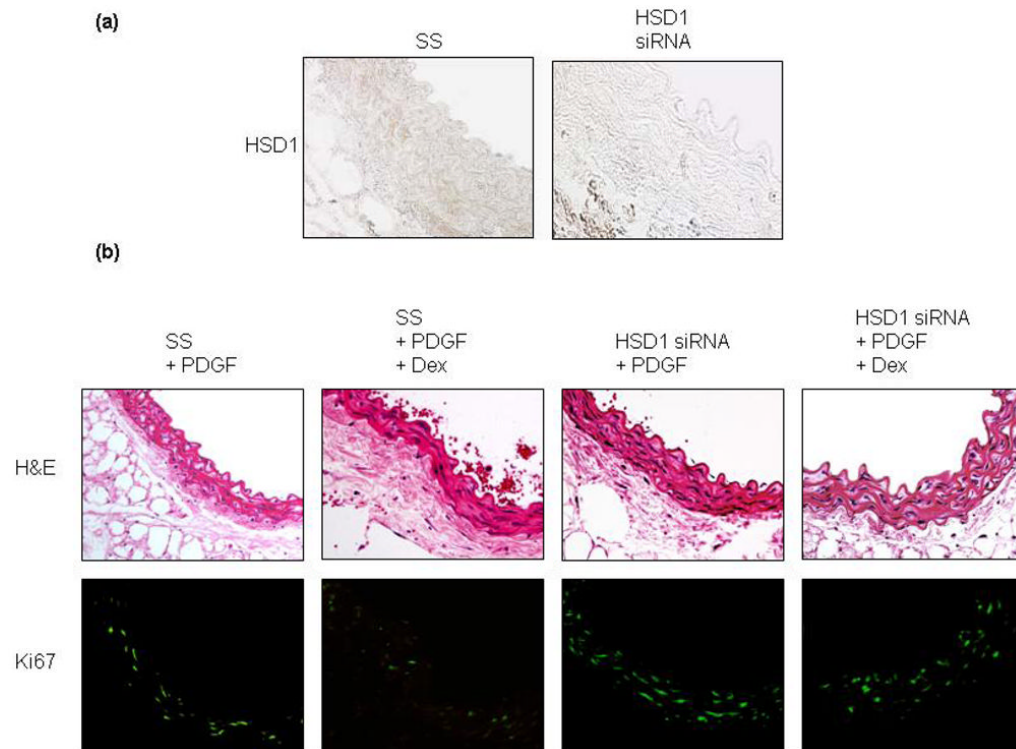


Fig. 6. 11 β -HSD1 mediates the inhibitory effects of dexamethasone on proliferation in PDGF-stimulated murine aortas

Thoracic aortas from C57Bl/6 male mice (n=12) were isolated, sectioned into 5 mm rings, and transfected with 80 nmol/L Stealth RNAiTM to 11 β -HSD1 mRNA or scrambled control (SS) added to smooth muscle basal medium. (a) After 48 h, 11 β -HSD1 expression was examined by immunohistochemistry of paraffin embedded sections. (b) The influence of 11 β -HSD1 on platelet-derived growth factor-BB (PDGF) (10 ng/ml)-stimulated proliferation, in the presence or absence of dexamethasone (10^{-7} mol/L), was examined by immunofluorescence labeling of Ki67. Representative hematoxylin & eosin stained sections are provided for comparison (*top*) and corresponding fluorescent sections are shown (*bottom*). Images are magnified 200X.

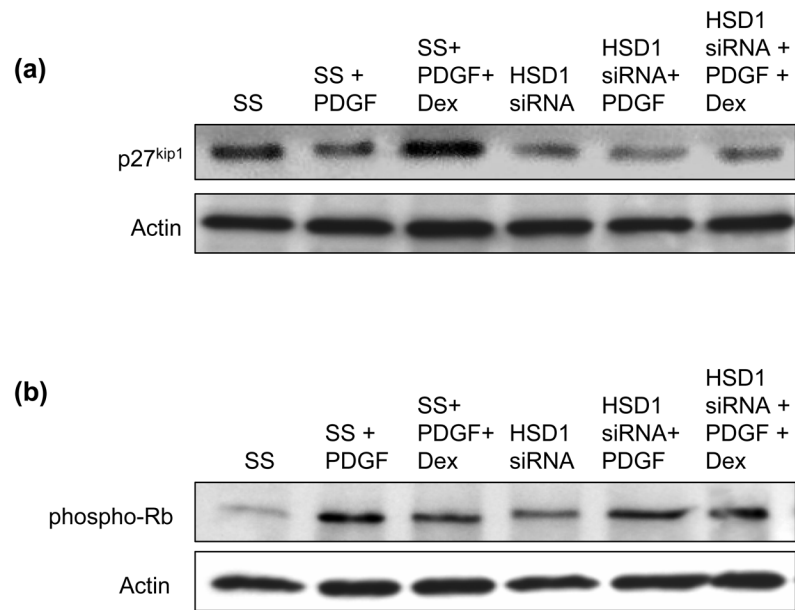


Fig. 7. 11 β -HSD1 influences the expression of cell cycle regulators in response to PDGF and dexamethasone

Human coronary artery smooth muscle cells (HCSMC) were transfected with an siRNA to decrease 11 β -HSD1 expression, synchronized by serum starvation (0.2% serum) for 48 h, and stimulated with 5% fetal calf serum and platelet-derived growth factor-BB (PDGF) (10 ng/ml) in the presence or absence of dexamethasone (10^{-7} mol/L). Expression of (a) p27^{kip1} (n=3) and (b) phosphorylation of retinoblastoma protein (phospho-Rb) at Ser608 (n=3) were determined by Western immunoblotting. Representative blots are shown