

## Aldosterone Production in Human Adrenocortical Cells Is Stimulated by High-Density Lipoprotein 2 (HDL2) through Increased Expression of Aldosterone Synthase (CYP11B2)

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Adrenal aldosterone production is regulated by physiological agonists at the level of early and late rate-limiting steps. Numerous studies have focused on the role of lipoproteins including high-density lipoprotein (HDL) as cholesterol providers in this process; however, recent research suggests that HDL can also act as a signaling molecule. Herein, we used the human H295R adrenocortical cell model to study the effects of HDL on adrenal aldosterone production and *CYP11B2* expression. HDL, especially HDL2, stimulated aldosterone synthesis by increasing expression of *CYP11B2*. HDL treatment increased *CYP11B2* mRNA in both a concentration- and time-dependent manner, with a maximal 19-fold increase (24 h, 250  $\mu$ g/ml of HDL). Effects of HDL on *CYP11B2* were not additive with natural agonists including angiotensin II or  $K^+$ . HDL effects were likely mediated by a calcium signaling cascade, because a calcium channel blocker and a calmodulin kinase inhibitor abolished the *CYP11B2*-stimulating effects. Of the two subfractions of HDL, HDL2 was more potent than HDL3 in stimulating aldosterone and *CYP11B2*. Further studies are needed to identify the active components of HDL, which regulate aldosterone production. (*Endocrinology* 152: 751–763, 2011)

The adrenal cortex is the only source of the mineralocorticoid aldosterone. Aldosterone secretion is normally under the tight control of the renin/angiotensin II (AngII)/aldosterone system (RAAS). Considerable evidence supports the role of two key regulatory steps in the control of adrenal cell aldosterone synthesis. The first has been termed the early rate-limiting step. AngII and  $K^+$  control this step through regulation of both protein expression and activities of the steroidogenic acute regulatory (StAR) protein, which facilitates cholesterol transport from the outer to inner mitochondrial membrane where conversion to pregnenolone occurs (1, 2). The second limiting reaction has been termed the late rate-limiting step, which represents the conversion of deoxycorticoste-

rone to aldosterone by the mitochondrial enzyme aldosterone synthase (CYP11B2) (3, 4).

CYP11B2 is expressed almost solely within the zona glomerulosa of the adrenal cortex (5, 6). As the final enzyme needed for aldosterone synthesis, CYP11B2 is regulated by multiple physiological agonists, including AngII and  $K^+$ . Both agonists increase intracellular calcium levels and activate calmodulin and calmodulin kinases (7, 8), which leads to increased CYP11B2 transcription (9–13). The expression of CYP11B2 influences directly the capacity of zona glomerulosa cells to produce aldosterone, and its adrenal expression varies *in vivo* depending on dietary sodium through regulation of the RAAS.

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Abbreviations: AngII, angiotensin II; FPLC, fast protein liquid chromatography; HDL, high-density lipoprotein; 12-HETE, 12-hydroxyeicosatetraenoic acid; LDL, low-density lipoprotein; qPCR, quantitative real-time RT-PCR; RAAS, renin/AngII/aldosterone system; siRNA, small interfering RNA; S1P, sphingosine 1 phosphate; SR-BI, scavenger receptor, class B type I; StAR, steroidogenic acute regulatory.

Circulating lipoproteins are important sources of cholesterol for adrenal steroidogenesis. Earlier studies have shown that in both fetal and adult human adrenals, low-density lipoprotein (LDL) increases ligand-stimulated steroid production by providing needed cholesterol for steroid biosynthesis (14–22). However, the function of high-density lipoprotein (HDL) in human adrenal steroidogenesis and particularly aldosterone synthesis remains poorly defined. Recent reports in vascular model systems suggest that, in addition to the traditional role in cholesterol transport, HDL can also act as a signaling molecule by activating the ERK1/2 pathway (23–26). The observation that HDL receptors [scavenger receptor, class B type I (SR-BI)] are highly expressed in both normal adrenal glands and aldosterone-producing adenomas (27, 28) supports a potential role for HDL in adrenal function and particularly aldosterone biosynthesis. Capponi and co-workers (29) have reported that HDL can stimulate aldosterone production in both bovine glomerulosa and H295R cells, but the mechanism was assumed to be through provision of cholesterol as a substrate for steroid synthesis.

In the current study, we demonstrate for the first time that HDL, especially HDL2, is able to stimulate adrenal cell aldosterone synthesis and steroidogenic enzyme expression. This indicates a potential new physiological mechanism for the regulation of aldosterone production by diseases or drugs that raise serum HDL2 levels.

## Materials and Methods

### Cell culture and reagents

The human adrenocortical cell line H295R was routinely cultured in DME/Ham F12 medium (Life Technologies, Inc., Carlsbad, CA) supplemented with 10% Cosmic calf serum (Hyclone, Logan, UT) and antibiotics including 1% penicillin/streptomycin (Life Technologies) and 0.1% gentamicin (Sigma-Aldrich, St. Louis, MO). One day before the experiment, cells were changed to low-serum experimental medium overnight (DME/F12 medium supplemented with 0.1% Cosmic calf serum and antibiotics). On the next morning, cells were treated with the indicated reagents in the low-serum experimental medium for the indicated time. For inhibitor studies, cells were preincubated with inhibitors for 30 min before any ligands were added. BLT-1 was purchased from Chembridge Corp. (San Diego, CA) and stored in dimethylsulfoxide. Other reagents, unless otherwise noted, were ordered from Sigma-Aldrich.

### Lipoproteins

Native human HDLs were purchased commercially from Calbiochem (Darmstadt, Germany) (three lots tested), AbD Serotec (Raleigh, NC) (one lot tested), and Millipore (Billerica, MA) (two lots tested) or prepared by our collaborators using previously described methods (30). All the figures represent results obtained using more than one source.

HDL2 and HDL3 (three sets) were provided and prepared as previously described (30). Briefly, HDL2 (1.066 g/ml  $\leq$  d  $\leq$  1.125 g/ml) and HDL3 (1.125 g/ml  $\leq$  d  $\leq$  1.210 g/ml) were isolated by sequential ultracentrifugation. Before use, HDL was dialyzed extensively against 0.9% NaCl, 10 mM HEPES (pH 7.4) and sterilized by filtration through a 0.45- $\mu$ m Millipore filter. Concentrations used for HDL represent the protein content in the lipoprotein particle.

### RNA isolation and quantitative real-time RT-PCR (qPCR)

Total RNA was extracted from cells using RNeasy mini kit (QIAGEN, Valencia, CA) following the manufacturer recommendations. Purity and integrity of the RNA was checked spectroscopically using a NanoDrop instrument (NanoDrop Technologies, Wilmington, DE). Total RNA (2  $\mu$ g) treated with deoxyribonuclease I (Ambion Inc., Austin, TX) was reverse transcribed using the High Capacity cDNA Archive Kit (Applied Biosystems, Foster City, CA) following the manufacturer recommendations and stored at  $-80$  C for further application.

The primer and probe sets for human *StAR*, side-chain cleavage (*CYP11A1*),  $3\beta$ -hydroxysteroid dehydrogenase-2 (*HSD3B2*), 21-hydroxylase (*CYP21*), *CYP11B1*, and *CYP11B2* were designed using Primer Express 3.0 (Applied Biosystems) and purchased from Integrated DNA Technologies (Coralville, IA) as published previously (12). *SR-BI* and *LDLR* gene expression assays were obtained from Applied Biosystems.

qPCR were performed in the ABI Prism 7500 sequence detection system (Applied Biosystems) in a total reaction mixture volume of 20  $\mu$ l following the reaction parameters recommended by the manufacturer. The TaqMan 2 $\times$  Master Mix (Applied Biosystems), 900 nM of each primer, 250 nM of probe, and 5  $\mu$ l of each first-strand cDNA sample were combined in each reaction vessel for gene detection. Negative controls contained water instead of first-strand cDNA. Relative quantification of mRNA levels between different tissues were determined using the comparative cycle threshold value as described previously (31), and *18S* rRNA was used as an internal control. The quantification of the *18S* rRNA in each sample was performed using a TaqMan ribosomal RNA reagent kit (Applied Biosystems) following the manufacturer recommendations.

### Protein extraction and protein assay

Cells were lysed in 100  $\mu$ l 1 $\times$  mammalian protein extraction reagent (Pierce Chemical Co., Rockford, IL). The protein content of samples was then determined by the bicinchoninic acid protein assay using the micro bicinchoninic acid protocol (Pierce).

### Steroid assay

Cell medium was collected after treatment and stored at  $-20$  C. The aldosterone content of the medium was analyzed using an RIA kit from Siemens Healthcare Diagnostics (Los Angeles, CA), and radioactivity was read by multicrystal  $\gamma$ -counter (Berthold Technologies, Bad Wildbad, Germany). Assays were conducted following the manufacturer's recommendation except that standard curves were prepared in the experimental cell culture medium. The results were normalized to protein amount and shown as the fold change over basal condition.

### Small interfering RNA (siRNA) silencing

Cell electroporation was performed using an AMAXA nucleofector kit following the manufacturer's recommendation (Lonza Group Ltd., Basel, Switzerland). Briefly, H295R cells were cultured until 80% confluence in normal growth medium before use. Cells were trypsinized and pelleted at a speed of 1000 rpm for 5 min. Supernatants were discarded, and cell pellets were resuspended in supplemented buffer R (Lonza) at a ratio of  $3 \times 10^6$  cells/100  $\mu$ l. For siRNA silencing experiments, Mission predesigned siRNA was ordered from Sigma-Aldrich and dissolved in nuclease-free water at a stock concentration of 50  $\mu$ M, and 2  $\mu$ l of siRNA was used per  $1 \times 10^6$  cells and electroporation was performed using program T-20 in the AMAXA system. Cells were allowed to recover in growth medium for 10 min before plating, and treatments were performed 48 h later.

### Western analysis

Immunoblotting was performed using XCell SureLock system (Invitrogen) following the manufacturer's recommendation. Briefly, samples were lysed with lysis buffer (2% sodium dodecyl sulfate, 62.5  $\mu$ M Tris, 0.04% bromophenol blue, 0.5 M dithiothreitol) and heated at 95 C for 5 min. Proteins were loaded in

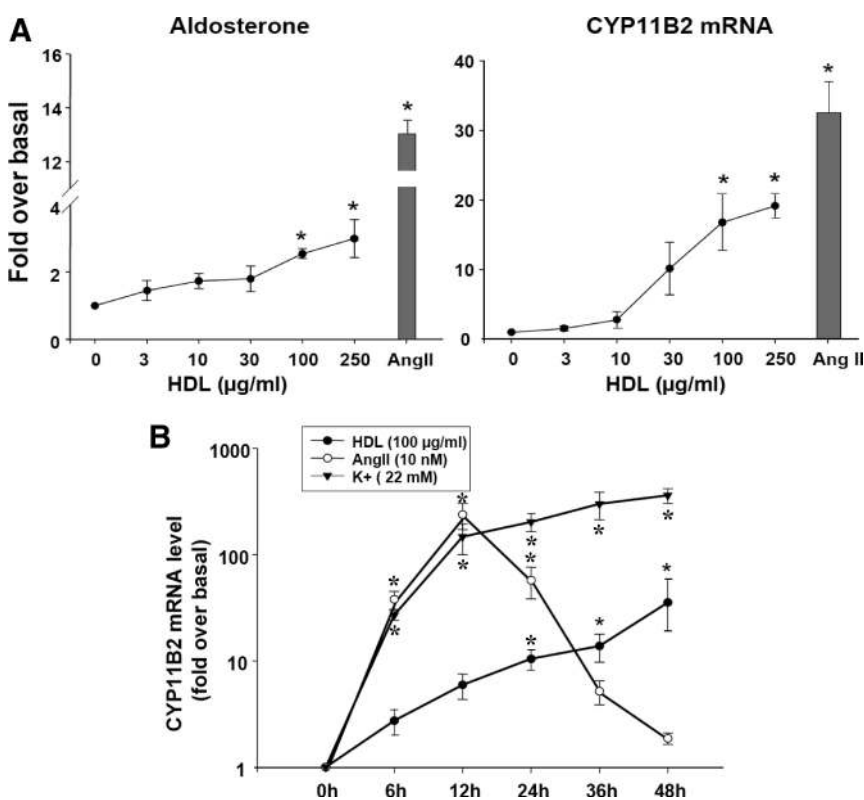
equal amounts to 10% Bis-Tris gel and allowed to separate for 1.5 h before transferring to polyvinylidene difluoride membranes. After blocking with 5% BSA for 1 h, the membrane was incubated with either SR-BI antibody (rabbit antihuman, 1:5000 in BSA; Novus Biologicals, Littleton, CO) or Nurr-1 antibody (rabbit antihuman, 1:1000; Santa Cruz Biotechnology Inc., Santa Cruz, CA), and second antibody (goat anti-rabbit, 1:7000; Invitrogen).

### Transfection and luciferase assay

The 5' flanking DNA from the human *CYP11B2* (–1521/+2) was inserted upstream of the firefly luciferase gene in the reporter vector pGL3 Basic (Promega, Madison, WI). Transient transfection assays were performed using Transfast (Promega) reagents in a ratio of 4  $\mu$ l/ $\mu$ g plasmid DNA. Luciferase activity was tested using the Luciferase assay system (Promega), and values are expressed as arbitrary light units normalized to the  $\beta$ -galactosidase activity of each sample and compared with basal condition for the fold change.

### Intracellular calcium assay

Calcium concentration was measured using Fluo-4 NW calcium assay kit (Invitrogen) following the manufacturer's recommendation. Briefly, H295R cells were cultured to 80–90% confluence in 96-well culture plates and preloaded with fluo-4 dye for 30 min at 37 C and measured at the same temperature with FLUOstar OPTIMA system (BMG LABTECH, Offenburger, Germany) using an excitation filter of 494 nm and emission of 516 nm. Basal levels were measured for 20 sec before injection, followed by 2 sec shaking and 100 sec of measurement. Inhibitors (losartan and BAPTA-AM) were added to loading dye buffer and incubated for 30 min before experiments.



**FIG. 1.** Effects of HDL on aldosterone production and *CYP11B2* expression in human H295R adrenal cells. **A**, HDL concentration-dependent effects on aldosterone production (*left*) and aldosterone synthase expression (*right*). Cells were treated with the indicated concentrations of HDL for 24 h. Media content of aldosterone was measured by immunoassay, and qPCR was used to quantify cellular *CYP11B2* mRNA. **B**, HDL time-dependent effects on *CYP11B2* expression. Cells were treated with HDL (100  $\mu$ g/ml), AngII (10 nM), or K<sup>+</sup> (22 mM) for the indicated times, and the expression of *CYP11B2* mRNA was quantified by qPCR. qPCR data were normalized to 18S rRNA. Data are shown as the fold change compared with basal. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using one-way ANOVA followed by a Dunnett test. \*,  $P < 0.05$ .

### Gel filtration chromatography [fast protein liquid chromatography (FPLC)]

FPLC was performed as previously described (32). Briefly, the lipoprotein samples were loaded (1 ml) onto a Superdex 200 column (60  $\times$  1.6 cm) using an Akta FPLC system and eluted with 10 mM Tris buffer at a flow rate of 1 ml/min (120 ml total column volume). After each run, the Superdex 200 column was washed with one column volume each of water and elution buffer.

### Statistical analysis

Results are provided as means  $\pm$  SEM. Individual experiments were repeated at least three times. Statistical analysis was performed using one-way ANOVA (including one-way ANOVA on rank) and paired *t* tests with a confidence interval of 95% using GraphPad Prism version 3.0 (GraphPad Software, Inc., San Diego, CA).

## Results

### HDL stimulates aldosterone production and CYP11B2 mRNA expression

Incubation of H295R cells with native human HDL for 24 h increased both aldosterone production and *CYP11B2* mRNA expression in a concentration-dependent manner with maximal responses detected at a concentration of 250  $\mu\text{g/ml}$  HDL (3-fold for aldosterone and 19-fold for *CYP11B2*) (Fig. 1A). Compared with AngII (10 nM), HDL stimulated aldosterone production only moderately (3-fold *vs.* 13-fold). This may be explained by the difference in StAR regulation by the two reagents. StAR protein Western analysis indicated that HDL did not significantly change basal StAR levels, whereas AngII stimulated its expression by 6-fold after a 6-h treatment (data not shown).

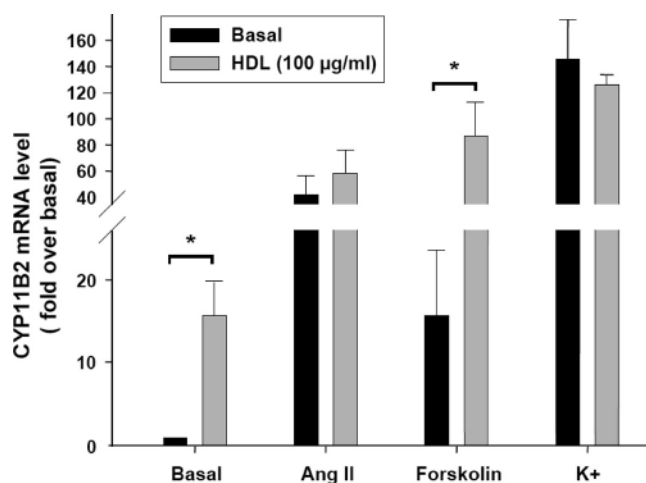
The time-dependent effects of HDL on *CYP11B2* mRNA expression were also examined in H295R (Fig. 1B). Although AngII caused a significant time-dependent increase in *CYP11B2* mRNA that peaked at 12 h, this increase was short-lived, and *CYP11B2* transcript levels almost returned to basal by 48 h. On the other hand,  $\text{K}^+$ -stimulated *CYP11B2* expression plateaued after 12 h. Like  $\text{K}^+$  treatment, HDL (100  $\mu\text{g/ml}$ ) increased the expression of *CYP11B2* transcript continuously for up to 2 d, suggesting that HDL is not working through the AngII signaling pathway. This was supported by the ability of the type 1 AngII receptor antagonist losartan to block AngII but not HDL stimulation of *CYP11B2* transcript levels (Supplemental Fig. 1, published on The Endocrine Society's Journals Online web site at <http://endo.endojournals.org>).

### Effects of HDL alone and with aldosterone-stimulating agonists on CYP11B2 expression

Cells were treated with HDL (100  $\mu\text{g/ml}$ ) alone or together with AngII (10 nM), forskolin (10  $\mu\text{M}$ ), or  $\text{K}^+$  (22 mM) for 24 h. Although HDL alone stimulated *CYP11B2* transcript levels by 15-fold, this effect was not additive with either AngII or  $\text{K}^+$  in H295R cells. However, when combined with the cAMP pathway agonist forskolin, HDL enhanced forskolin stimulation of *CYP11B2* from 15-fold to more than 80-fold (Fig. 2A). This result strongly suggests that HDL used a signaling pathway other than cAMP, but a pathway that was shared by AngII and  $\text{K}^+$ .

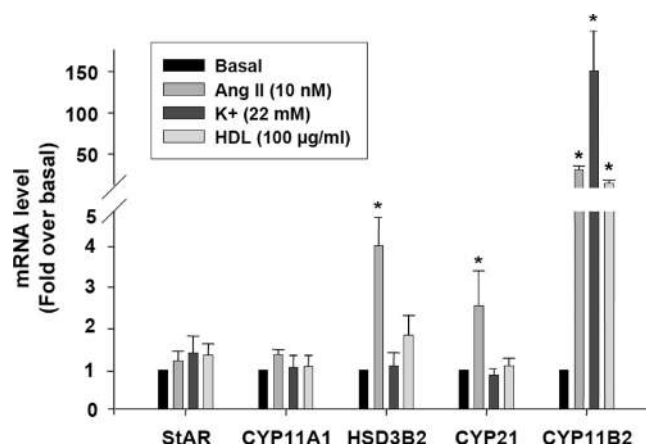
### HDL effects on steroidogenic enzyme expression

To further characterize the effects of HDL on aldosterone stimulation, the mRNA encoding the enzymes/proteins involved in aldosterone biosynthesis were examined and compared with AngII and  $\text{K}^+$ . As reported previously, AngII significantly increased expression of *HSD3B2*,

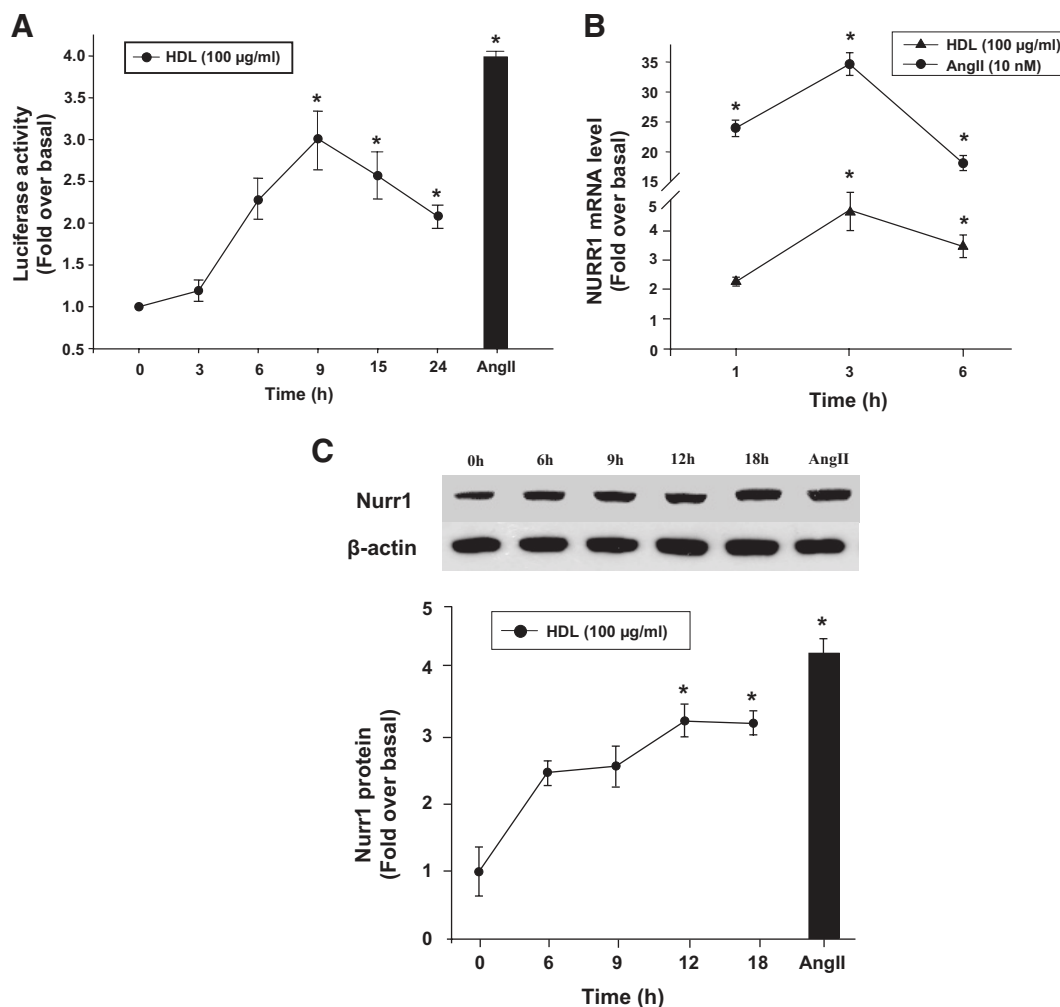


**FIG. 2.** Effects of HDL alone and with aldosterone-stimulating agonists on *CYP11B2* expression. H295R adrenal cells were treated with AngII (10 nM), forskolin (10  $\mu\text{M}$ ), or  $\text{K}^+$  (22 mM) with or without HDL for 24 h. RNA was isolated and used for qPCR measurement of *CYP11B2* with data normalized to *18S* rRNA and shown as the fold change compared with basal. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using one-way ANOVA followed by a Dunnett test. \*,  $P < 0.05$ .

*CYP21*, and *CYP11B2* (33–36). However, *CYP11B2* was the only transcript that was significantly up-regulated by HDL treatment (Fig. 3). This is similar to the effects of  $\text{K}^+$  on steroidogenic enzyme transcript expression. Transcripts related to cortisol and dehydroepiandrosterone production were also examined, and there was no difference in *CYP17* expression level, whereas *CYP11B1* was moderately increased (4.3-fold with 100  $\mu\text{g/ml}$  HDL treatment, data not shown). The expression of this transcript can also be increased by Ang II and  $\text{K}^+$  (37, 38).



**FIG. 3.** HDL effects on steroidogenic enzyme transcript levels. H295R cells were treated with AngII (10 nM),  $\text{K}^+$  (22 mM) or HDL (100  $\mu\text{g/ml}$ ) for 24 h. RNA was isolated and reverse transcribed for qPCR quantification. Data were normalized to *18S* rRNA and shown as the fold change compared with basal. Statistical significances were determined using one-way ANOVA followed by a Dunnett test. \*,  $P < 0.05$ .



**FIG. 4.** HDL effects on *CYP11B2* promoter activity and NURR1 expression. **A**, HDL effects on *CYP11B2* promoter activity. H295R cells were transfected with the *CYP11B2* promoter construct (1  $\mu$ g/ml) and allowed to recover overnight. Luciferase activity was tested after treatment with or without HDL treatment (100  $\mu$ g/ml) for the indicated time. Data were normalized to cotransfected  $\beta$ -galactosidase vector and are shown as the fold induction over the untreated group. As a positive control, cells were treated with AngII (10 nM) for 9 h. Results represent the means  $\pm$  SEM from at least three independent experiments each performed in triplicate. Statistical differences were determined using one-way ANOVA followed by Dunnett test. \*,  $P < 0.05$ . **B** and **C**, HDL stimulates both transcript (**B**) and protein (**C**) expression of NURR1. H295R cells were treated with AngII (10 nM) or HDL (100  $\mu$ g/ml) for the indicated times (1–18 h). qPCR was used to quantify *NURR1* mRNA level, and Western analysis was used for NURR1 protein quantification. qPCR results were normalized to 18S rRNA, and Western data were normalized to  $\beta$ -actin protein levels. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using one-way ANOVA followed by a Dunnett test. \*,  $P < 0.05$ .

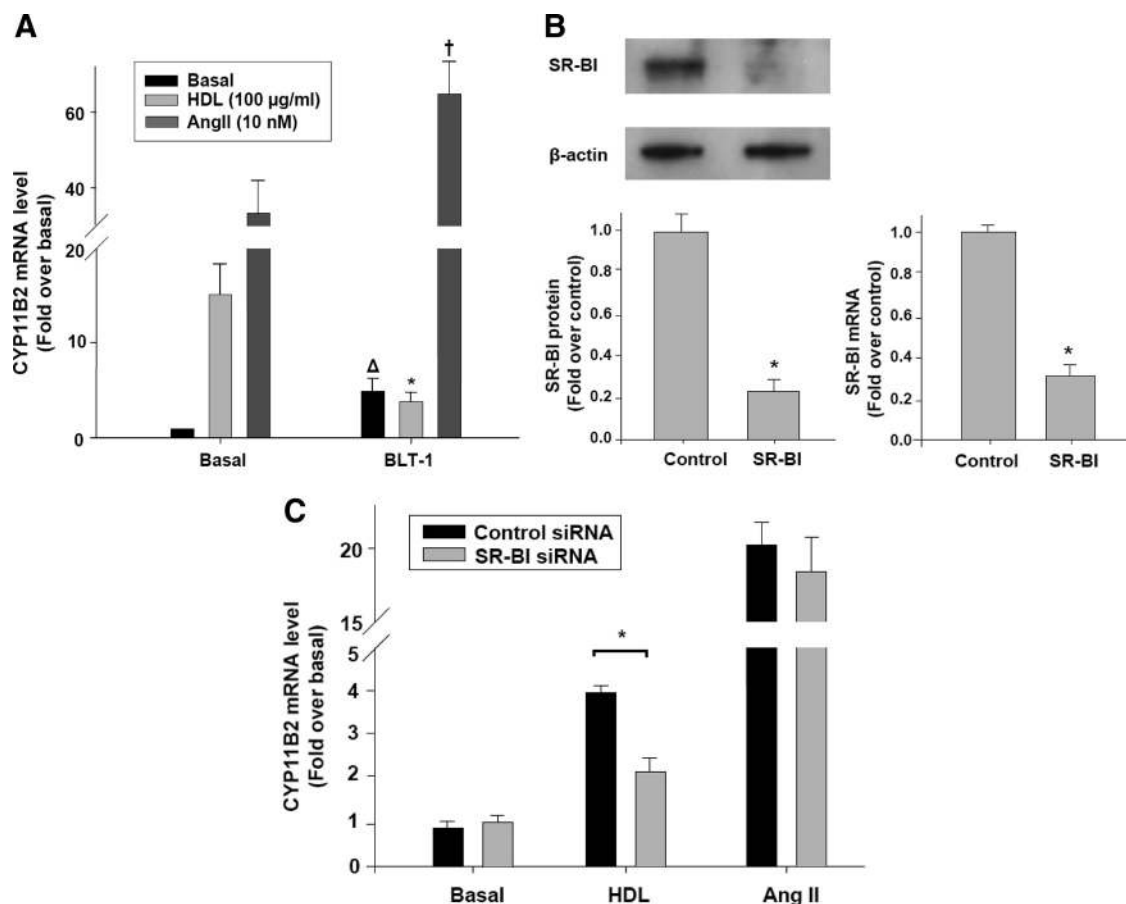
### HDL stimulates *CYP11B2* promoter activity and NURR1 expression

H295R cells were transfected with a vector possessing the 5' flanking *CYP11B2* promoter region and treated with HDL or AngII for 3–24 h. HDL significantly stimulated promoter activity in a time-dependent manner, starting from 6 h and peaking around 9 h, with a maximal stimulation of 2.8-fold (Fig. 4A). We have previously demonstrated a role for the transcription factor NURR1 in the transcription of *CYP11B2* (11, 39). HDL caused a time-dependent increase in *NURR1* transcript (4.5-fold at 3 h) (Fig. 4B) and protein (3-fold at 12 h) (Fig. 4C), suggesting that the HDL-triggered signaling cascade activates the expression of NURR1 and thus stimulates *CYP11B2* transcription.

### HDL stimulation is mediated by SR-BI

The adrenal cortex of several species expresses high levels of the HDL receptor SR-BI (40–44). The human adrenal is no exception, and transcripts and protein expression have been documented (27, 28, 45). To examine the role of SR-BI in the HDL-mediated signaling pathway, we used a SR-BI-specific inhibitor, BLT-1. As shown in Fig. 5A, although BLT-1 itself increased the basal level of *CYP11B2* mRNA, preincubating with BLT-1 blocked the HDL stimulatory effects. The inhibition appeared to be specific, because BLT-1 did not decrease AngII activation of cellular *CYP11B2* expression.

The involvement of SR-BI was further confirmed by siRNA knockdown of SR-BI expression. Transient deliv-



**FIG. 5.** The role of SR-BI receptor in HDL regulation of *CYP11B2*. **A**, BLT-1 effects on HDL stimulation of *CYP11B2*. H295R cells were preincubated with BLT-1 (3  $\mu$ M) for 20 min, followed by basal, HDL (100  $\mu$ g/ml), or AngII (10 nM) treatment for 24 h. RNA were isolated and reverse transcribed for qPCR quantification. qPCR data were normalized to 18S rRNA and shown as the fold change compared with untreated basal cells. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistics were calculated using one-way ANOVA followed by a Dunnett test.  $\Delta$ ,  $P < 0.05$ , compared with basal; \*,  $P < 0.05$ , compared with HDL treatment; †,  $P < 0.05$ , compared with AngII treatment. **B** and **C**, SR-BI knockdown effects on HDL stimulation of *CYP11B2* mRNA. H295R cells were allowed to grow to 80–90% confluence. Electroporation was performed as described in *Materials and Methods*, and cells were plated for 2 d before treatment with HDL (100  $\mu$ g/ml) for 24 h. RNA was isolated, and qPCR was performed for *CYP11B2* mRNA quantification. Data were normalized to 18S rRNA and shown as the fold change compared with control siRNA-transfected cells (**B**). Western blotting was performed as described in *Materials and Methods*. Western data were normalized to  $\beta$ -actin protein level and shown as the fold change compared with control siRNA-transfected cells (**C**). Data represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using either Student's paired  $t$  test (**B**) or one-way ANOVA followed by a Dunnett test (**C**). \*,  $P < 0.05$ .

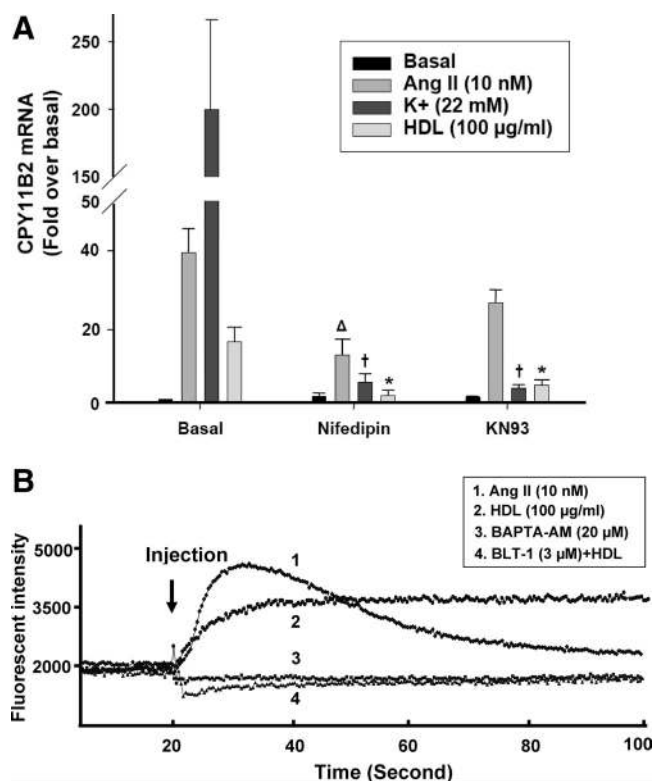
ery of siRNA for SR-BI decreased the endogenous expression of *SR-BI* mRNA to less than 30% and the protein level to 25% when compared with cells transfected with scrambled siRNA (Fig. 5B). After 24 h treatment with HDL, the knockdown group exhibited decreased HDL-stimulating effects on *CYP11B2* by 50%, although AngII effects were not influenced (Fig. 5C). However, we did notice a decrease in overall fold stimulation in response to both ligands as a result of the AMAXA electroporation.

### HDL stimulates adrenal cell *CYP11B2* expression through calcium signaling pathways

The importance of calcium signaling for HDL-stimulated *CYP11B2* expression was investigated with the help of the calcium channel blocker nifedipine and calmodulin kinase inhibitor KN93 (Fig. 6A). Preincubation with 10  $\mu$ M nifedipine completely blocked HDL effects on

*CYP11B2* and 5  $\mu$ M KN93 inhibited 80% of HDL activity. The inhibition of both HDL and  $K^+$  activity by nifedipine and KN93 suggests that like  $K^+$ , HDL signals mainly through calcium. In agreement with previous results, both nifedipine and KN93 only partially inhibit AngII-stimulating effects, indicating the involvement of another signaling cascade (likely protein kinase C) in AngII effects on *CYP11B2* expression.

This result was confirmed by examination of intracellular calcium concentrations in H295R cells (Fig. 6B). As published previously (46–48), AngII caused a significant but transient increase in intracellular calcium concentrations. HDL triggered a lower-magnitude peak  $Ca^{2+}$  increase, but this increase was maintained for more than 100 sec. Both a calcium chelator (BAPTA-AM) and an SR-BI inhibitor (BLT-1) blocked this change in calcium concentration, suggesting that HDL stimulated calcium influx



**FIG. 6.** Role of calcium signaling in HDL stimulation of *CYP11B2* expression. **A**, Calcium signaling antagonists inhibited HDL-induced stimulation of *CYP11B2*. H295R cells were preincubated with nifedipine (10  $\mu$ M) or KN93 (5  $\mu$ M) for 20 min followed by treatment with agonists (24 h). *CYP11B2* qPCR data were normalized to 18S ribosome RNA and shown as the fold change compared with untreated basal cells. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using Student's paired *t* test.  $\Delta$ ,  $P < 0.05$ , compared with AngII treatment;  $\dagger$ ,  $P < 0.05$ , compared with K<sup>+</sup> treatment; \*,  $P < 0.05$ , compared with HDL treatment. **B**, HDL increased the intracellular calcium concentration in H295R cells. Cells were allowed to grow to 80–90% confluence on a 96-well dish and incubated with fluorescent dye (100  $\mu$ l) for 40 min before use. Cells were preincubated with and without BAPTA-AM or BLT-1 for 30 min. Fluorescent intensity was recorded for the 20 sec before and 80 sec after agonist addition.

and this process was mediated by SR-BI. Preincubation with AT1 receptor block losartan completely abolished the calcium-stimulating effects of AngII but did not influence HDL function (data not shown).

### HDL2 is more potent than HDL3 in stimulating adrenal cells

When comparing HDL from different sources, we observed a range of adrenal cell *CYP11B2* responses. All three commercially obtained samples (from Calbiochem, AbD Serotec, and Millipore) showed similar significant effects on aldosterone and *CYP11B2* stimulation, whereas samples by our collaborators had only minor activity (Fig. 7A). To clarify the differences in HDL function, FPLC was performed to determine the HDL components of an HDL preparation with high stimulatory activity (sample 2, from

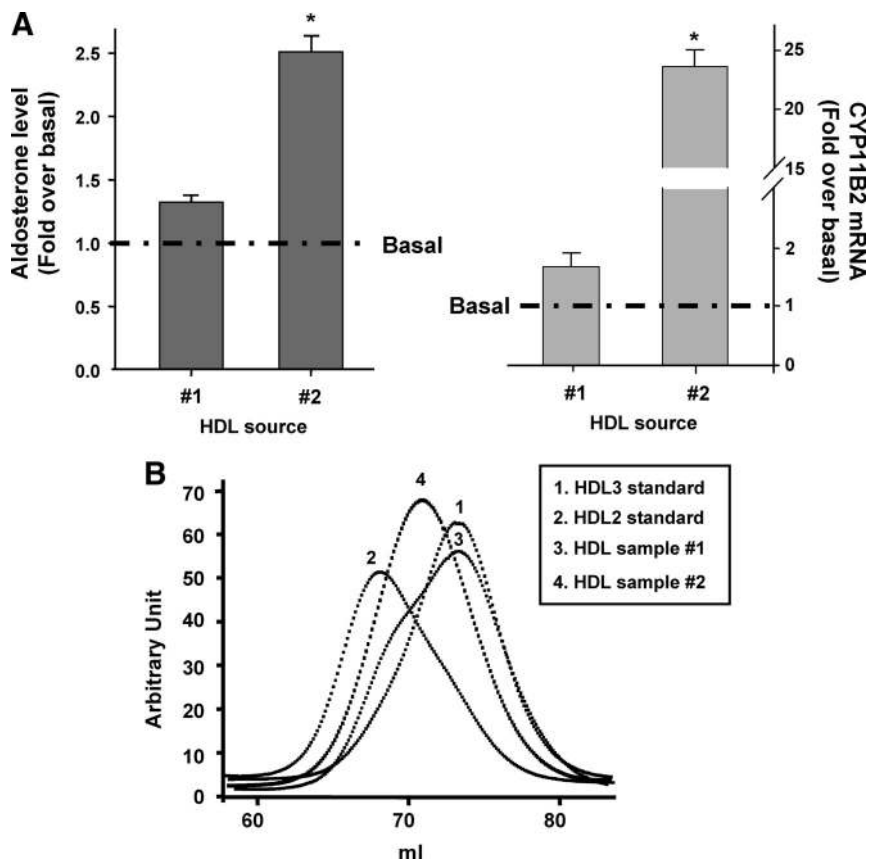
Calbiochem) and low stimulatory activity (sample 1, prepared within our group). As shown in Fig. 7B, HDL from our group was mainly composed of the HDL3 subfraction, whereas Calbiochem HDL had more of an HDL2 component. This result suggested that different HDL subfractions might influence the aldosterone/*CYP11B2*-stimulating effect. This hypothesis was tested by specifically isolating HDL2 and HDL3 subfractions, with the total HDL isolated from the same sample as control, and using them to treat H295R cells. HDL2 (100  $\mu$ g/ml) significantly increased *CYP11B2* transcript levels, whereas HDL3 (100  $\mu$ g/ml) did not influence *CYP11B2* expression (Fig. 8A).

To confirm the above finding, a similar set of experiments was performed comparing HDL2- and HDL3-stimulating effects on aldosterone production, *CYP11B2* transcripts, promoter activity, intracellular calcium concentration, and *NURR1* mRNA level (Fig. 8). HDL2 was more active than HDL3, and this idea suggested that HDL2 might be a more potent initiator of the calcium signaling pathway in human adrenocortical cells.

## Discussion

Like all steroid-producing glands, the adrenal synthesizes steroid hormones using cholesterol as a precursor. There are three primary cholesterol sources for steroidogenic tissues: *de novo* synthesis, LDL, and HDL. Different animal species have their own preferred source of cholesterol. For humans, LDL is the main cholesterol source for adrenal steroidogenesis (14–17), whereas rats and mice use HDL as the major source of cholesterol in steroid synthesis (49, 50). The varied role of lipoproteins in steroid production may relate in part to the apoprotein content in HDL that has been shown to be quite different between those species. Although direct incubation of HDL with rat adrenocortical cells did not increase steroid production *in vitro*, HDL has been shown to enhance steroidogenesis with ACTH (50). However, Leydig cells from rats can use HDL to regulate testosterone production via an apolipoprotein A1-dependent pathway (51). Furthermore, in mouse, HDL has been shown to stimulate StAR protein expression (52) and act as a potential growth factor for adrenocortical cells (53).

Bovine steroidogenic cells, on the other hand, appear to use both LDL and HDL for steroidogenesis (29, 54), and bovine glomerulosa cells respond to HDL with increases in aldosterone production in a mechanism that may involve a calcium signaling pathway (55). We also observed a significant stimulation of bovine glomerulosa cell aldosterone production by HDL in the current study (data not shown). Similar effects were reported by Capponi's group,



**FIG. 7.** Variation in HDL2 vs. HDL3 ratio in HDL from different sources and their effects on aldosterone production and *CYP11B2* expression. **A**, Variable effects of HDL on aldosterone production and *CYP11B2* expression. H295R cells were treated with the indicated concentration of HDL for 24 h. RIA was performed to determine the concentration of aldosterone produced by cells, and qPCR was performed for quantifying *CYP11B2* mRNA level. All data are shown as the fold change compared with basal. Results represent the means  $\pm$  SEM from at least three independent experiments. Statistical significances were determined using one-way ANOVA followed by a Dunnett test. \*,  $P < 0.05$ . **B**, Representative diagram of FPLC comparison of the HDL components from different sources. Purified HDL2 and HDL3 were used as controls. HDL sample 1 was isolated by our research laboratory, and sample 2 was purchased from Calbiochem.

and they further extended the stimulating effects of HDL to H295 human adrenocortical cells (29). However, the latter article made the assumption that the HDL effect was associated with its cholesterol-providing function. For the current study, we replicated the aldosterone-stimulating effects in H295R cells and proposed a novel mechanism that HDL may initiate a calcium signaling pathway leading to increased expression of the late rate-limiting step enzyme *CYP11B2*.

HDL is used by most cells mainly through binding to SR-BI. SR-BI is an 82-kDa membrane protein with 509 amino acids (56, 57). It contains one putative membrane-spanning hydrophobic domain near each end of the protein and a large extracellular region with a cysteine-rich carboxyl-terminal half and multiple potential *N*-glycosylation sites. SR-BI is highly glycosylated and can also be modified by fatty acylation (58). As opposed to the receptor-mediated endocytosis pathway used by LDL receptors

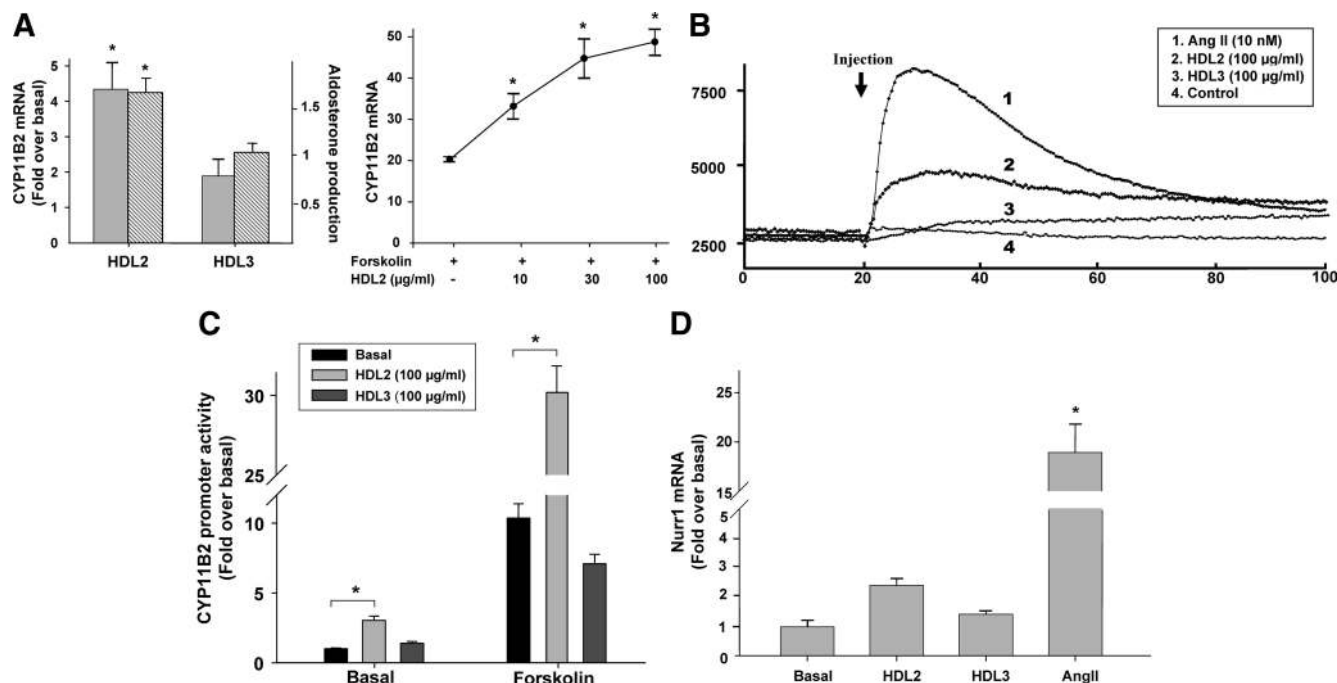
in lipid uptake, SR-BI uses a selective uptake mechanism, which efficiently transports lipid but not the carrier protein into cells (57, 59). This process can be divided into two independent steps: receptor-mediated surface binding and lipid transportation across the membrane. With the use of the specific inhibitor of SR-BI, BLT-1 (60), we were able to separate these two steps and show that lipid uptake rather than receptor binding is required to trigger aldosterone production and *CYP11B2* stimulation in adrenocortical cells.

As a lipoprotein family member, HDL can be further divided into two subfractions depending on particle size and density, namely HDL2 and HDL3. Although lower levels of both HDL subfractions have been shown to be markers of coronary heart disease (61), HDL3 exerts a greater preventive effect on  $\text{Cu}^{2+}$ -induced LDL oxidation (62). HDL2 and HDL3 differ in both lipoprotein and lipid components (63–65). A study conducted by Rothblat and colleagues (30) suggested that the phospholipid composition of HDL is a major determinant of the bidirectional flux and net movement of cellular free cholesterol mediated by SR-BI, raising the possibility that HDL2 and HDL3 have diverse effects on SR-BI-mediated cholesterol transport. The HDL2/HDL3 ratio can be shifted by both physiological

(exercise, menopause, and gender) (66–69) and pathological (familial hypercholesterolemia, obesity, alcohol consumption, smoking, and substance abuse) conditions (70–75). Our data suggest that the changes in HDL level (especially HDL2/HDL3 ratio) might modulate adrenal aldosterone production. Although aldosterone levels are primarily regulated by RAAS, most humans who are on a high-sodium diet have very low renin levels. Under these conditions, HDL2 may play a role in regulating aldosterone production. Furthermore, our study suggested that HDL2 has the potential to act with ACTH to increase aldosterone production. In addition, the high expression level of SR-BI in adrenal steroid-producing tumors (76) further emphasized the need to examine the role of HDL in the dysregulation of steroidogenesis seen in adrenal disease.

One mechanism for HDL activation of adrenal cell signaling would be through lipids or proteins associated with





**FIG. 8.** Potency of HDL2 vs. HDL3 in stimulating adrenal cells. **A**, HDL2 had stronger stimulatory effects on both aldosterone production and *CYP11B2* mRNA expression. H295R cells were treated with HDL2 or HDL3 (100 µg/ml) with or without forskolin (10 µM) for 24 h in 0.1% low-serum medium. RIA was performed to determine the concentration of aldosterone produced by cells, and qPCR was performed for quantifying *CYP11B2* mRNA level. All data are shown as the fold change compared with basal (solid bar, *CYP11B2* mRNA; dashed bar, aldosterone production). Results represent the means ± SEM from at least three independent experiments. Statistical differences were determined using one-way ANOVA followed by Dunnett test. \*,  $P < 0.05$ . **B**, HDL2 increased the intracellular calcium concentration in H295R cells. Cells were allowed to grow to 80–90% confluence on a 96-well dish and incubated with fluorescent dye (100 µM) for 40 min before use. Cells were exposed to the indicated agonists, and the plate was shaken for 2 sec to allow quick mixing. Fluorescent intensity was recorded for the 20 sec before and 80 sec after agonist addition. **C**, HDL2 was more potent than HDL3 in stimulating *CYP11B2* promoter activity and was additive with forskolin. H295R cells were transfected with 1 µg of *CYP11B2* promoter construct and allowed to recover overnight. Luciferase activity was tested after treatment with or without HDL2, HDL3 (100 µg/ml), or forskolin (10 µM) for 6 h. Data were normalized to cotransfected β-galactosidase vector and are shown as the fold induction over the untreated group. Results represent the means ± SEM from at least three independent experiments. Statistical differences were determined using ANOVA on rank. \*,  $P < 0.05$ . **D**, HDL2 stimulates the expression of *NURR1*. H295R cells were treated with AngII (10 nM) or HDL2 or HDL3 (100 µg/ml) for 3 h. qPCR was used to quantify cellular *NURR1* mRNA level, and data were normalized to 18S rRNA. Results represent the means ± SEM from at least three independent experiments. Statistical differences were determined using one-way ANOVA. \*,  $P < 0.05$ .

the HDL particle. Based on previous studies (77–82), potential candidates for the active component of HDL were examined using adrenocortical cells. Sphingosine 1 phosphate (S1P) has been shown, in multiple studies, to account for many of the antiatherogenic activities of HDL (77, 78, 83–86). Mounting evidence supports the hypothesis that HDL, by binding to SR-BI, increases endothelial nitric oxide synthase phosphorylation via the activation of both phosphatidylinositol 3-kinase and MAPK pathways (79, 87). Recent results also suggest the involvement of S1P receptors (especially type 1 and type 3) in this signaling process (24). However, treatment of H295R cells with S1P or its analog, FTY720, did not stimulate H295R cells and preincubation of a S1P receptor inhibitor (VPC 23019) failed to block HDL stimulation of *CYP11B2* (data not shown). In addition, HDL3 has twice the amount of S1P than the more steroidogenically active HDL2 (88), further suggesting that S1P is not the stimulatory agent for adrenal cells.

Another candidate is 12-hydroxyeicosatetraenoic acid (12-HETE), which has been proposed to have a role in mediating AngII-induced aldosterone secretion. Nadler and co-workers (81, 82, 89–91) reported that AngII treatment increases the expression of 12-lipoxygenase in human cultured glomerulosa cells and stimulates 12-HETE production in H295R cells. Elevated 12-lipoxygenase levels raised aldosterone secretion under both basal and AngII-stimulated conditions, accompanied by an increase in *CYP11B2* transcript levels (89). In addition, in both bovine and rat glomerulosa cell models, treatment of 12-HETE causes an influx of calcium (80, 92). However, because of the fact that no 12-HETE receptor has been identified, the mechanisms for 12-HETE stimulation of aldosterone production have not been determined. In the present study, we were not able to observe the effects of 12-HETE on H295R *CYP11B2* and aldosterone (data not shown), suggesting that this lipid is not the active component found in HDL.

Recently, an interesting finding from a phase III clinical trial was published, in which a potent inhibitor of cholesteryl ester transfer protein was studied as a potential treatment for cardiovascular disease. Noticeably, treatment caused a 20% decrease in LDL and a 61% increase in HDL in patients after torcetrapib/statin administration (93–95). The increase in circulating HDL was mainly due to elevated HDL2 concentrations (93, 96). However, treatment also caused an elevation in blood pressure and aldosterone levels in subjects given torcetrapib. Current evidence suggests that the effects of torcetrapib were related to a nonspecific action that was unrelated to the inhibition of cholesteryl ester transfer protein (97–99). Although there are no clinical data indicating that individuals with high HDL have more hypertension or higher aldosterone levels (or lower  $K^+$  levels), our *in vitro* observations using cultured adrenal cells suggest that further studies are now needed to determine whether HDL2 can directly influence adrenal gland production of aldosterone.

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