Scavenger receptor class B type I-mediated uptake of serum cholesterol is essential for optimal adrenal glucocorticoid production

Menno Hoekstra, 1 Dan Ye, Reeni B. Hildebrand, Ying Zhao, Bart Lammers, Miranda Stitzinger, Johan Kuiper, Theo J. C. Van Berkel, and Miranda Van Eck

Division of Biopharmaceutics, Leiden/Amsterdam Center for Drug Research, Gorlaeus Laboratories, 2300RA Leiden, The Netherlands

Abstract  Impaired scavenger receptor class B type I (SR-BI)-mediated uptake of HDL-cholesterol esters (HDL-CE) induces adrenal insufficiency in mice. Humans contain an alternative route of HDL-CE clearance, namely through the transfer by cholesteryl ester transfer protein (CETP) to apolipoprotein B lipoproteins for subsequent uptake via the LDL receptor. In this study, we determined whether CETP can compensate for loss of adrenal SR-BI. Transgenic expression of human CETP (CETP Tg) in SR-BI knockout (KO) mice increased adrenal HDL-CE clearance from 33–58% of the control value. SR-BI KO/CETP Tg and SR-BI KO mice displayed adrenal hypertrophy due to equally high plasma adrenocorticotropic hormone levels. Adrenal cholesterol levels and plasma corticosterone levels were 38–52% decreased in SR-BI KO mice with and without CETP expression. SR-BI KO/CETP Tg mice also failed to increase their corticosterone level after lipo polysaccharide challenge, leading to an identical 4-fold increased tumor necrosis factor-α response compared with controls. These data indicate that uptake of CE via other routes than SR-BI is not sufficient to generate the cholesterol pool needed for optimal adrenal steroidogenesis. In conclusion, we have shown that CETP-mediated transfer of HDL-CE is not able to reverse adrenal insufficiency in SR-BI knockout mice. Thus, SR-BI-mediated uptake of serum cholesterol is essential for optimal adrenal function.—Hoekstra, M., D. Ye, R. B. Hildebrand, Y. Zhao, B. Lammers, M. Stitzinger, J. Kuiper, T. J. C. Van Berkel, and M. Van Eck. Scavenger receptor class B type I-mediated uptake of serum cholesterol is essential for optimal adrenal glucocorticoid production. J. Lipid Res. 2009. 50: 1039–1046.

Supplementary key words  adrenals • cholesteryl ester transfer protein • high density lipoprotein • low density lipoprotein • cholesteryl ester • corticosterone • inflammation

HDL is a small, dense protein-lipid complex that consists of a hydrophilic phospholipid monolayer and a hydrophobic core filled with cholesteryl esters. The HDL-mediated transport of cholesterol from peripheral tissues back to the liver for subsequent excretion from the body, often referred to as reverse cholesterol transport [reviewed in Van Eck et al. (1) and Lewis and Rader (2)], is considered to be an important physiological process to maintain total body cholesterol homeostasis. Scavenger receptor class B type I (SR-BI) is a multiligand cell surface receptor predominantly expressed in the liver and steroidogenic tissues (i.e., testis, ovary, and adrenals) (3). In mice, SR-BI is a prominent factor in the reverse cholesterol transport process, as it is the sole molecule involved in the selective uptake of cholesteryl esters from HDL by the liver (4). In addition to its established function in liver cholesterol metabolism, we (5) and others (6) have recently shown that SR-BI also plays a major role in adrenal cholesterol metabolism and steroid hormone production in mice. As a result, SR-BI deficiency in mice is associated with a depletion of adrenal cholesterol stores, resulting in a diminished steroid hormone (i.e., glucocorticoid) production in response to physiological stress. Importantly, the metabolism of HDL differs significantly between humans and mice as humans, in contrast to rodents, naturally express cholesteryl ester transfer protein (CETP). CETP is synthesized by macrophage-rich tissues, such as the spleen and liver, after which it is secreted into the blood circulation (7–10). In plasma, CETP is able to transfer cholesteryl esters from HDL to apolipoprotein B (ApoB)-containing particles, such as VLDLs and LDLs. LDL-associated cholesteryl esters can subsequently be

Abbreviations: ACTH, adrenocorticotrophic hormone; ApoB, apolipoprotein B; CETP, cholesteryl ester transfer protein; Ct, threshold cycle number; HPRT, hypoxanthine guanine phosphoribosyl transferase; KO, knockout; SR-BI, scavenger receptor class B type I; TNFa, tumor necrosis factor-α; WT, wild-type.

1 To whom correspondence should be addressed.

e-mail: hoekstra@lacdr.leidenuniv.nl

This research was supported by Top Institute Pharma (TIPharma project T2-110: M.H. and T.J.C.V.B.), by Grants 2001T41 (M.V.E.), 2006B107 (B.L.), and 2008T070 (M.H.) from the Netherlands Heart Foundation, and by VDI Grant 917.66.301 from the Netherlands Organization for Scientific Research (M.V.E.). M.V.E. is an Established Investigator of the Netherlands Heart Foundation (Grant 2007T056).

Manuscript received 1 August 2008 and in revised form 18 December 2008 and in re-revised form 26 January 2009.

Published, JLR Papers in Press, January 28, 2009.
DOI 10.1194/jlr.M800410-JLR200

Copyright © 2009 by the American Society for Biochemistry and Molecular Biology, Inc.

This article is available online at http://www.jlr.org
cleared from the blood circulation via whole particle uptake by the LDL receptor (ApoB receptor) (11, 12). In addition to SR-BI, the CETP→LDL→LDL receptor pathway can thus serve as an alternative route for the delivery of HDL-cholesteryl esters to cells in humans. However, the relative contribution of the SR-BI and the CETP→LDL→LDL receptor routes for the uptake of HDL-cholesteryl esters in humans is still unclear because polymorphisms in the human SR-BI gene leading to functional SR-BI deficiency so far have not been detected.

In vitro studies using adrenocortical cells have suggested that LDL receptor-mediated uptake of LDL is coupled to steroid hormone synthesis (13). In addition, Kita et al. (14) and Fong et al. (15) have shown that the adrenals express a functional LDL receptor in vivo. It is thus likely that the CETP→LDL→LDL receptor route is relevant for adrenal cholesterol homeostasis in the human situation. To gain insight in the possible role for the CETP→LDL→LDL receptor route in adrenal cholesterol metabolism, we have determined whether transgenic expression of human CETP can rescue the adrenal glucocorticoid insufficiency in SR-BI knockout (KO) mice.

## MATERIALS AND METHODS

### Animals

SR-BI KO mice were kindly provided by Dr. M. Krieger (Massachusetts Institute of Technology, Boston, MA) (16). CETP transgenic mice (CETP Tg; strain 5203) expressing human CETP under the control of its own promoter and other major regulatory elements were obtained from The Jackson Laboratory (Bar Harbor, ME) (17). SR-BI KO mice were bred with CETP Tg mice to generate heterozygous SR-BI KO mice expressing CETP on one allele. These mice were subsequently cross-bred to generate SR-BI KO mice expressing human CETP on one allele (SR-BI KO/CETP Tg) or control SR-BI KO mice and wild-type (WT) nontransgenic littermates. The presence of the targeted and wild-type SR-BI alleles as well as the CETP transgene was assessed by PCR amplification of DNA extracted from tail biopsies (primers: 5′-GATGGGACATGGGACACGAAGCCATTCT-3′ and 5′-TCTGTCTCCGTCCTCCCTCAGTCTGA-3′ for SR-BI and 5′-CTAGGCCACAGAATTGAAAGATCT-3′, 5′-GATGGTGGAATTCTAGCATCATCC-3′, and 5′-GAATGTCTCAGAGTGGGACATGGGACACGAAGCCATTCT-3′ for CETP). Mice were fed a sterilized regular chow powder diet (RM3; Special Diet Services, Witham, UK). Animal experiments were performed at the Gorlaeus Laboratories of the Leiden/Amsterdam Center for Drug Research in accordance with the national laws. All experimental protocols were approved by the Ethics Committee for Animal Experiments of the Leiden University.

### Isolation and labeling of HDL

Human HDL was isolated from blood of healthy subjects by differential ultracentrifugation as described by Redgrave, Roberts, and West (20) and dialyzed against PBS with 1 mM EDTA. HDL (1.063 < d < 1.21) was labeled with [3H]cholesteryl ether (CEt) via exchange from donor particles as reported previously (21).

### Adrenal uptake of [3H]-Cholesteryl ether HDL

A dose of 200 μg apolipoprotein (±1.2 × 10⁶ dpm) of [3H]CEt-HDL (total volume 100 μl) was injected into the tail vein. At 5 min after injection, a blood sample was drawn to verify the injected dose. For analysis of adrenal cholesteryl ether uptake, 24 h after tracer injection adrenals were excised, weighed, solubilized, and counted for 3H-radioactivity in a Packard liquid scintillation unit. A correction was made for the radioactivity in the blood present in the adrenals at the time of sampling (135.2 μl/g tissue).

### Lipoprotein distribution analysis

The distribution of cholesterol or [3H]-label over the different lipoproteins in plasma was analyzed by fractionation of pooled plasma using a Superose 6 column (3.2 × 300 mm, Smart-system; Pharmacia). Total cholesterol content of the effluent was determined using enzymatic colorimetric assays (Roche Diagnostics). Counting for [3H]-radioactivity was performed in a Packard liquid scintillation unit.

### Adrenal histology and immunohistochemical analysis for CETP

Formalin-fixed cryosections (8–10 μm) were prepared on a Leica CM3050-S cryostat. Cryosections were routinely stained with hematoxylin and Oil red O for nuclei and neutral lipids, respectively. For immunohistochemical staining of CETP, cryostat sections were incubated for 5 min with prewarmed (37°C) 0.025% trypsin at room temperature, blocked with 1% BSA in TBS, and

### Analysis of gene expression by real-time quantitative PCR

Quantitative gene expression analysis on snap-frozen organs was performed as described (18). In short, total RNA was isolated according to Chomczynski and Sacchi (19) and reverse transcribed using RevertAid™ reverse transcriptase. Gene expression analysis was performed using real-time SYBR Green technology (Eurogentec) with the primers displayed in Table 1, which were validated for identical efficiencies (slope = −3.3 for a plot of threshold cycle number (Ct) versus log ng cDNA). Hypoxanthine guanine phosphoribosyl transferase (HPRT), GAPDH, and acidic ribosomal phosphoprotein P0 (36B4) were used as the standard housekeeping genes. Relative gene expression numbers were calculated by subtracting the Ct of the target gene from the average Ct of HPRT, GAPDH, and 36B4 (Ct housekeeping) and raising 2 to the power of this difference. Genes that exhibited a Ct value of ≥35 were considered not detectable. The average Ct of three housekeeping genes was used to exclude that changes in the relative expression were caused by variations in the expression of the separate housekeeping genes.

### Quantitative gene expression analysis on snap-frozen organs

Analysis of gene expression by real-time quantitative PCR was performed as described (18). In short, total RNA was isolated according to Chomczynski and Sacchi (19) and reverse transcribed using RevertAid™ reverse transcriptase. Gene expression analysis was performed using real-time SYBR Green technology (Eurogentec) with the primers displayed in Table 1, which were validated for identical efficiencies (slope = −3.3 for a plot of threshold cycle number (Ct) versus log ng cDNA). Hypoxanthine guanine phosphoribosyl transferase (HPRT), GAPDH, and acidic ribosomal phosphoprotein P0 (36B4) were used as the standard housekeeping genes. Relative gene expression numbers were calculated by subtracting the Ct of the target gene from the average Ct of HPRT, GAPDH, and 36B4 (Ct housekeeping) and raising 2 to the power of this difference. Genes that exhibited a Ct value of ≥35 were considered not detectable. The average Ct of three housekeeping genes was used to exclude that changes in the relative expression were caused by variations in the expression of the separate housekeeping genes.

### Lipoprotein distribution analysis

The distribution of cholesterol or [3H]-label over the different lipoproteins in plasma was analyzed by fractionation of pooled plasma using a Superose 6 column (3.2 × 300 mm, Smart-system; Pharmacia). Total cholesterol content of the effluent was determined using enzymatic colorimetric assays (Roche Diagnostics). Counting for [3H]-radioactivity was performed in a Packard liquid scintillation unit.

### Adrenal histology and immunohistochemical analysis for CETP

Formalin-fixed cryosections (8–10 μm) were prepared on a Leica CM3050-S cryostat. Cryosections were routinely stained with hematoxylin and Oil red O for nuclei and neutral lipids, respectively. For immunohistochemical staining of CETP, cryostat sections were incubated for 5 min with prewarmed (37°C) 0.025% trypsin at room temperature, blocked with 1% BSA in TBS, and

### Analysis of gene expression by real-time quantitative PCR

Quantitative gene expression analysis on snap-frozen organs was performed as described (18). In short, total RNA was isolated according to Chomczynski and Sacchi (19) and reverse transcribed using RevertAid™ reverse transcriptase. Gene expression analysis was performed using real-time SYBR Green technology (Eurogentec) with the primers displayed in Table 1, which were validated for identical efficiencies (slope = −3.3 for a plot of threshold cycle number (Ct) versus log ng cDNA). Hypoxanthine guanine phosphoribosyl transferase (HPRT), GAPDH, and acidic ribosomal phosphoprotein P0 (36B4) were used as the standard housekeeping genes. Relative gene expression numbers were calculated by subtracting the Ct of the target gene from the average Ct of HPRT, GAPDH, and 36B4 (Ct housekeeping) and raising 2 to the power of this difference. Genes that exhibited a Ct value of ≥35 were considered not detectable. The average Ct of three housekeeping genes was used to exclude that changes in the relative expression were caused by variations in the expression of the separate housekeeping genes.

### Lipoprotein distribution analysis

The distribution of cholesterol or [3H]-label over the different lipoproteins in plasma was analyzed by fractionation of pooled plasma using a Superose 6 column (3.2 × 300 mm, Smart-system; Pharmacia). Total cholesterol content of the effluent was determined using enzymatic colorimetric assays (Roche Diagnostics). Counting for [3H]-radioactivity was performed in a Packard liquid scintillation unit.

### Adrenal histology and immunohistochemical analysis for CETP

Formalin-fixed cryosections (8–10 μm) were prepared on a Leica CM3050-S cryostat. Cryosections were routinely stained with hematoxylin and Oil red O for nuclei and neutral lipids, respectively. For immunohistochemical staining of CETP, cryostat sections were incubated for 5 min with prewarmed (37°C) 0.025% trypsin at room temperature, blocked with 1% BSA in TBS, and

### Analysis of gene expression by real-time quantitative PCR

Quantitative gene expression analysis on snap-frozen organs was performed as described (18). In short, total RNA was isolated according to Chomczynski and Sacchi (19) and reverse transcribed using RevertAid™ reverse transcriptase. Gene expression analysis was performed using real-time SYBR Green technology (Eurogentec) with the primers displayed in Table 1, which were validated for identical efficiencies (slope = −3.3 for a plot of threshold cycle number (Ct) versus log ng cDNA). Hypoxanthine guanine phosphoribosyl transferase (HPRT), GAPDH, and acidic ribosomal phosphoprotein P0 (36B4) were used as the standard housekeeping genes. Relative gene expression numbers were calculated by subtracting the Ct of the target gene from the average Ct of HPRT, GAPDH, and 36B4 (Ct housekeeping) and raising 2 to the power of this difference. Genes that exhibited a Ct value of ≥35 were considered not detectable. The average Ct of three housekeeping genes was used to exclude that changes in the relative expression were caused by variations in the expression of the separate housekeeping genes.

### Lipoprotein distribution analysis

The distribution of cholesterol or [3H]-label over the different lipoproteins in plasma was analyzed by fractionation of pooled plasma using a Superose 6 column (3.2 × 300 mm, Smart-system; Pharmacia). Total cholesterol content of the effluent was determined using enzymatic colorimetric assays (Roche Diagnostics). Counting for [3H]-radioactivity was performed in a Packard liquid scintillation unit.

### Adrenal histology and immunohistochemical analysis for CETP

Formalin-fixed cryosections (8–10 μm) were prepared on a Leica CM3050-S cryostat. Cryosections were routinely stained with hematoxylin and Oil red O for nuclei and neutral lipids, respectively. For immunohistochemical staining of CETP, cryostat sections were incubated for 5 min with prewarmed (37°C) 0.025% trypsin at room temperature, blocked with 1% BSA in TBS, and
incubated with a primary CETP antibody TP-2 (Ottawa Heart Institute, Ontario, Canada) and a secondary antibody conjugated to peroxidase (Jackson ImmunoResearch Labs, Suffolk, UK). Images were obtained with a Leica image analysis system, consisting of a Leica DMRE microscope coupled to a camera and Leica Qwin Imaging software (Cambridge, UK).

**Plasma hormone analysis**

Blood was drawn via the tail vein between 0900 and 1200 h for subsequent hormone analyses. Corticosterone and adrenocorticotropic hormone (ACTH) levels in plasma were determined using the CORTICOSTERONE and ACTH Double Antibody 125I Radioimmunoassay (RIA) kits from MP Biomedicals (Irvine, CA) according to the protocols from the supplier.

**Tumor necrosis factor-α response upon lipopolysaccharide challenge**

Mice were intravenously injected at 0900 h with 50 μg/kg LPS from Salmonella minnesota R595 (List Biological Laboratories, Hornby, Canada) into the tail vein. Blood samples were collected after 30, 60, 90, 120, and 180 min, and plasma tumor necrosis factor-α (TNF-α) protein levels were determined by ELISA (OptEIA kit; BD Biosciences Pharmingen, San Diego, CA).

**Data analysis**

Data were presented as means ± SEM. Statistical analyses were performed using one- and two-way ANOVA using Graphpad Prism Software (Graphpad Software, San Diego, CA). The level of statistical significance was set at P < 0.05.

**RESULTS**

In this study, we determined whether the CETP→LDL→LDL receptor route may be physiologically relevant for adrenal cholesterol metabolism and steroid hormone production. For this purpose, we cross-bred mice carrying the human CETP transgene linked to its natural flanking sequences (CETP Tg) with SR-BI KO mice to generate “human-like” SR-BI KO mice carrying the CETP transgene (SR-BI KO/CETP Tg mice). SR-BI KO/CETP Tg mice, like control SR-BI KO mice, lack a functional SR-BI gene, as evident from the undetectable mRNA expression of SR-BI in liver, spleen, adrenals, and white adipose tissue (Table 2). However, in contrast with control SR-BI KO mice, SR-BI KO/CETP Tg mice express relatively high levels of human CETP mRNA in macrophage-rich tissues, such as the liver and spleen, similarly as found in the human situation (8).

To investigate whether CETP expression in SR-BI KO mice induces transfer of cholesteryl esters from HDL to ApoB-containing lipoproteins for subsequent uptake by the LDL receptor in the adrenals, we determined the effect of transgenic expression of human CETP on plasma total cholesterol levels and the adrenal HDL-cholesteryl ester uptake in SR-BI KO mice. As shown in Fig. 1A, transgenic expression of human CETP induced an almost complete normalization of plasma total cholesterol levels in SR-BI KO mice, indicating that the CETP-mediated transfer of cholesteryl esters from HDL to LDL/LDL fraction can also act as an effective alternative pathway to remove cholesteryl esters from the blood circulation in SR-BI KO mice. In parallel to the observed change in plasma cholesterol levels, the uptake of [3H]CEt-HDL by the adrenals was also partially restored in SR-BI KO/CETP Tg mice compared with SR-BI KO mice [20 ± 1% vs. 11 ± 1% of the injected dose/g tissue (P < 0.05), respectively, compared with 34 ± 3% for wild-type mice; Fig. 1B]. CETP-mediated transfer of cholesteryl esters from HDL to ApoB-containing lipoproteins thus does significantly increase the adrenals uptake of HDL-cholesteryl esters in SR-BI KO mice probably via the action of the LDL receptor. Evaluation of the [3H]CEt distribution over the different lipoprotein fractions at 8 h after [3H]CEt-HDL injection showed that 4.2-fold more of the [3H]label resided in the nonHDL fraction per μg cholesterol in SR-BI KO/CETP Tg mice compared with SR-BI KO mice, while a 30% lower amount of [3H]label per μg cholesterol was detected in the HDL fraction (Fig. 1C), resulting in an overall 6-fold higher nonHDL/HDL label distribution ratio in SR-BI KO/CETP Tg mice compared with SR-BI KO mice. This suggests that human CETP, as anticipated, efficiently transports cholesteryl esters from HDL to ApoB-containing (nonHDL) lipoproteins in SR-BI KO mice.

Compared with wild-type mice, adrenals from SR-BI KO mice are darker due to an almost complete depletion of lipid in the adrenal cortex as a result of the impaired adrenal uptake of HDL-cholesteryl esters (5). Strikingly, macroscopically the adrenals of SR-BI KO/CETP Tg mice and SR-BI KO mice are essentially identical and are significantly different from those of wild-type mice. As can be appreciated from the photographs depicted in Fig. 2A, both the adrenals of SR-BI KO/CETP Tg and SR-BI KO mice are visually darker (red color instead of white color) and have a larger volume compared with those of wild-type mice. In agreement, an overall higher weight of the adrenals was observed (5.2 ± 0.4 mg and 4.8 ± 0.3 mg for SR-BI KO/CETP and SR-BI KO mice vs. 3.4 ± 0.2 mg for WT mice; P < 0.05 for both; Fig. 2B). As the external phenotype of the adrenals in SR-BI KO mice was not affected by the expression of CETP, we determined the extent of lipid loading in the adrenal cortex of the three different types of mice (Fig. 3). In accordance with our previous findings (5), Oil red O neutral lipid

---

**Table 2. Relative mRNA expression levels of SR-BI and CETP in different organs isolated from WT mice, SR-BI KO mice, and SR-BI KO mice expressing human CETP (SR-BI KO/CETP Tg)**

<table>
<thead>
<tr>
<th></th>
<th>Liver</th>
<th>Spleen</th>
<th>Adrenal</th>
<th>White Adipose Tissue</th>
</tr>
</thead>
<tbody>
<tr>
<td>SR-BI</td>
<td>0.12 ± 0.02</td>
<td>n.d.</td>
<td>0.018 ± 0.001</td>
<td>n.d.</td>
</tr>
<tr>
<td>SR-BI KO</td>
<td>n.d.</td>
<td>n.d.</td>
<td>0.42 ± 0.12</td>
<td>n.d.</td>
</tr>
<tr>
<td>SR-BI KO/CETP Tg</td>
<td>n.d.</td>
<td>0.14 ± 0.13</td>
<td>n.d.</td>
<td>0.012 ± 0.002</td>
</tr>
</tbody>
</table>

Values are relative to the housekeeping gene expression and represent means ± SEM of three mice per group. n.d., not detectable.

SR-BI is essential for optimal adrenal function 1041
staining in the adrenal cortex was almost completely absent in SR-BI KO mice compared with wild-type mice (Fig. 3A). Furthermore, no remarkable change in Oil red O staining was observed upon expression of CETP in SR-BI KO mice (Fig. 3A) as only limited Oil red O staining could be seen specifically in the zona glomerulosa (the mineralocorticoid producing cells) but not the zona fasciculata (the glucocorticoid producing cells) or zona reticularis (production of androgens) of the adrenal cortex of both SR-BI KO/CETP and control SR-BI KO mice. In contrast, all three zones of the cortex stained positive for neutral lipid in the adrenals of WT mice (Fig. 3A). Lipid analyses indicated that the adrenal cholesteryl ester content was not significantly different between SR-BI KO (7.3 ± 1.1 µg/mg protein) and SR-BI KO/CETP Tg mice (8.5 ± 0.6 µg/mg protein) but was significantly lower (P < 0.001 for both) compared with wild-type controls (16.5 ± 1.4 µg/mg protein). Immunohistochemical staining for CETP showed that the CETP protein is specifically expressed in the zona glomerulosa within the cortex of adrenals from SR-BI KO/CETP Tg mice (Fig. 3B). Combined, these findings suggest that the transfer of HDL-cholesteryl esters to ApoB-containing lipoproteins leading to subsequent uptake via the LDL receptor does not significantly contribute to cholesteryl ester storage in the adrenal cortex, while the adrenal expression of human CETP is insufficient to normalize cholesteryl ester storage.

Importantly, adrenocortical cells within the zona fasciculata need cholesterol to produce glucocorticoid hormones, such as cortisol in humans and corticosterone in rodents. As the zona fasciculata of SR-BI KO mice with or without transgenic expression of CETP is equally depleted of lipid, we determined the effect of CETP expression on plasma corticosterone levels. Under fed (nonstressed) conditions, no significant change in the plasma corticosterone levels was observed between the three different genotypes (Fig. 4). In contrast, plasma corticosterone levels in SR-BI KO and SR-BI KO/CETP Tg mice were significantly lower (P < 0.01 for both) compared with those of WT mice under conditions of physiological stress induced by overnight fasting (Fig. 4). Measurements of ACTH, a potent activator of adrenal cortex growth and steroidogenesis (22), indicated that the impaired...
The corticosterone response to fasting was not due to a decrease in ACTH levels as SR-BI KO and SR-BI KO/CETP Tg mice displayed a 2- to 3-fold increase ($P < 0.05$) in plasma ACTH levels compared with WT controls under fasting conditions (Fig. 5A). In addition, the mRNA expression of the LDL receptor, which is rapidly induced by ACTH in adrenocortical cells both in vitro (13, 23) and in vivo (24), was stimulated >3-fold ($P < 0.01$) in the adrenals of SR-BI KO mice and SR-BI KO/CETP Tg mice, indicative for optimal ACTH signaling in the adrenals of these animals (Fig. 5B). As CETP expression does not normalize the corticosterone response to fasting at a background of relatively high adrenal LDL receptor expression in SR-BI KO mice, it is suggested that SR-BI KO/CETP Tg mice similarly to control SR-BI KO mice also suffer from adrenal glucocorticoid insufficiency.

Recent studies by Cai et al. (6) have indicated that SR-BI KO mice are more susceptible to LPS-induced septic shock and death as a result of the adrenal glucocorticoid insufficiency. We also observed a significantly ($P < 0.001$) enhanced response of the pro-inflammatory cytokine TNF-α in SR-BI KO mice compared with wild-type mice upon injection of a sublethal dose of LPS (50 μg/kg) (Fig. 6A). Interestingly, the peak value of the plasma TNF-α level at 60 min after LPS injection (6.25 ± 0.52 ng/ml for SR-BI KO/CETP Tg and 6.23 ± 0.03 ng/ml for SR-BI KO mice) and the total TNF-α response in time (area-under-curve 553 vs. 573 ng/ml min, respectively) were identical between SR-BI KO/CETP Tg and control SR-BI KO mice upon the challenge with LPS (Fig. 6A). In addition, both SR-BI KO and SR-BI KO/CETP Tg mice failed to induce plasma corticosterone levels in response to LPS (Fig. 6B), which further establishes that transgenic expression of CETP is not able to overcome the adrenal glucocorticoid insufficiency in SR-BI KO mice.

**DISCUSSION**

SR-BI deficiency in mice leads to glucocorticoid insufficiency as a result of an impaired adrenal uptake of cholesteryl esters from HDL (4, 5). Within the adrenals, a constant supply of cholesterol is required to serve as precursor for the synthesis of mineralocorticoids and glucocorticoids (25). The cholesterol needed for optimal steroid synthesis in the adrenals can be acquired from 1) intracellular de novo synthesis of free cholesterol by the enzyme HMG-CoA reductase, 2) intracellular catabolism of stored cholesteryl esters.
esters to free cholesterol by neutral cholesteryl ester hydrolase, or 3) receptor-mediated uptake and subsequent intracellular catabolism of cholesteryl esters from circulating VLDL, LDL, and HDL particles [reviewed in Kraemer (25)]. In addition to the “HDL receptor” SR-BI, adrenals express the LDL receptor, which is involved in the whole particle endocytosis of the ApoB-containing lipoproteins VLDL and LDL (12). Importantly, the expression of SR-BI and the LDL receptor and steroidogenesis are coordinately regulated by activators of both glucocorticoid (dibutyryl cAMP and ACTH) and mineralocorticoid (PMA) synthesis in adrenocortical cells both in vitro and in vivo (24, 26–31), leading to the generally accepted suggestion that the LDL receptor may also contribute to adrenal steroid hormone synthesis in vivo. However, it has never been experimentally defined to what extent the LDL receptor indeed allows optimal adrenal hormone production, i.e., corticosterone production. There are no patients with a deficiency of the human ortholog of SR-BI called CLA-1 (CD36 and lysosomal integral membrane protein-II analogous-1) (31). In the human situation, an improper functioning of CLA-1 is expected to be compensated for by the presence of CETP, a protein that redistributes cholesteryl esters from HDL to ApoB-containing lipoproteins, including LDL, leading to an alternative uptake route of cholesteryl esters by the adrenals. In our experiments, we confirm that expression of human CETP in SR-BI KO mice indeed leads to the transfer of cholesteryl esters from the relatively large HDL particles to ApoB-containing lipoproteins accessible for LDL receptor-mediated internalization (32). As already anticipated, the presence of CETP in SR-BI KO mice resulted in an increased delivery of cholesteryl esters from HDL via ApoB-containing lipoproteins to the adrenals. The enhanced delivery of cholesteryl esters upon CETP expression, however, was not associated with a reversal of the changes in adrenal morphology or plasma corticosterone and ACTH levels associated with SR-BI deficiency. Furthermore, no difference was observed in the LPS-induced TNF-α response between SR-BI KO mice with and without CETP expression. These data show for the first time that the uptake of cholesteryl esters to free cholesterol by neutral cholesteryl ester hydrolase, or 3) receptor-mediated uptake and subsequent intracellular catabolism of cholesteryl esters from circulating VLDL, LDL, and HDL particles [reviewed in Kraemer (25)]. In addition to the “HDL receptor” SR-BI, adrenals express the LDL receptor, which is involved in the whole particle endocytosis of the ApoB-containing lipoproteins VLDL and LDL (12). Importantly, the expression of SR-BI and the LDL receptor and steroidogenesis are coordinately regulated by activators of both glucocorticoid (dibutyryl cAMP and ACTH) and mineralocorticoid (PMA) synthesis in adrenocortical cells both in vitro and in vivo (24, 26–31), leading to the generally accepted suggestion that the LDL receptor may also contribute to adrenal steroid hormone synthesis in vivo. However, it has never been experimentally defined to what extent the LDL receptor indeed allows optimal adrenal hormone production, i.e., corticosterone production. There are no patients with a deficiency of the human ortholog of SR-BI called CLA-1 (CD36 and lysosomal integral membrane protein-II analogous-1) (31). In
esters via other routes than SR-BI (i.e., via the LDL receptor) is not sufficiently effective to generate the cholesterol pool needed for optimal adrenal steroid hormone production. In accordance with this, recent studies by Kraemer et al. (33) using LDL receptor KO mice have indicated that endocytic adrenal cholesterol uptake via the LDL receptor is not necessary for ACTH- and dibutyryl cAMP-induced maximal corticosterone production in vivo. Furthermore, inhibition of SR-BI function in murine adrenocortical cells in vitro induced a marked 70–80% decrease in selective cholesteryl ester uptake and corticosterone secretion (34). We now firmly establish that SR-BI-mediated uptake of cholesteryl esters is the primary route for the delivery of HDL-cholesterol to the steroidogenic pathway in vivo and that LDL receptor-mediated uptake cannot compensate for the absence of the SR-BI-mediated uptake route.

In conclusion, we have shown that under human-like conditions, that is, in the presence of human CETP, the transfer of cholesteryl esters from HDL to ApoB-containing lipoproteins by CETP is not able to reverse the adrenal insufficiency in SR-BI KO mice. Our findings suggest that the adrenal uptake of HDL-cholesteryl esters via SR-BI probably also contributes significantly to adrenal steroidogenesis in humans. However, it should be acknowledged that the relative acceptor (i.e., LDL) availability for CETP-mediated transfer of cholesteryl esters from HDL is much higher in the human situation (high LDL and HDL levels) than in our current SR-BI KO mouse model (high HDL and very low LDL levels). An additional key point is that the higher LDL levels in humans provide more substrate for the LDL receptor-mediated uptake of LDL cholesterol by the adrenals. This is also why the results with LDL-poor mice should be interpreted cautiously when extrapolating to humans. It will therefore be important to study whether carriers of polymorphisms in the human SR-BI (CLA-1) gene that are associated with changes in HDL-cholesterol levels (35–38) are indeed more susceptible to adrenal insufficiency, stress-related diseases (i.e., anxiety and depression), and/or inflammatory diseases (i.e., sepsis, rheumatoid arthritis, and atherosclerosis).}

REFERENCES


