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## Supplemental Text

### Materials and Methods

#### CETP Assay

##### *Sample Preparation and Western blotting*

For lipoprotein fractionation, equal volumes of plasma samples were pooled from six MMPs and six rabbits fed normal chow. The lipoproteins were fractionated using fast protein liquid chromatography (FPLC) with a Superose 6 10/300 GL FPLC column (Amersham Biosciences, Piscataway, NJ). Diluted plasma and aliquots of the FPLC fractions were subjected to SDS-PAGE and transferred onto PVDF membranes. Immunoblotting was performed as previously described. Antibodies against cholesteryl ester (CE) transfer protein (CETP) (mouse monoclonal, designated as "antibody A," and rabbit polyclonal, designated as "antibody B") were purchased from EMD Millipore (Billerica, MA) and Abcam (Cambridge Science Park, Cambridge, UK), respectively. The proteins were visualized using a chemiluminescence method (ECL Plus Western Blotting Detection System; GE Healthcare Piscataway, NJ).

##### *Diagnosis of CETP Deficiency and Determination of the CETP Mass*

A 59-year-old man and 54-year-old woman with high HDL-C levels (172 and 191 mg/dL, respectively) were referred to the National Defense Medical College

Hospital. Blood samples were obtained, and the sera were isolated with centrifugation. Genomic DNA was prepared using a commercially available kit (Qiagen, Frederick, MA). The CETP mass was measured using a sandwich ELISA with JHC1 and JHC2, two monoclonal antibodies specific to human CETP, as previously described<sup>1</sup>. Common CETP mutations (an intron 14 splicing defect, Int14 +1 G>A; amino acid substitution from aspartic acid to glycine at position 442, D442G) were identified using an Invader<sup>®</sup> assay<sup>2</sup>.

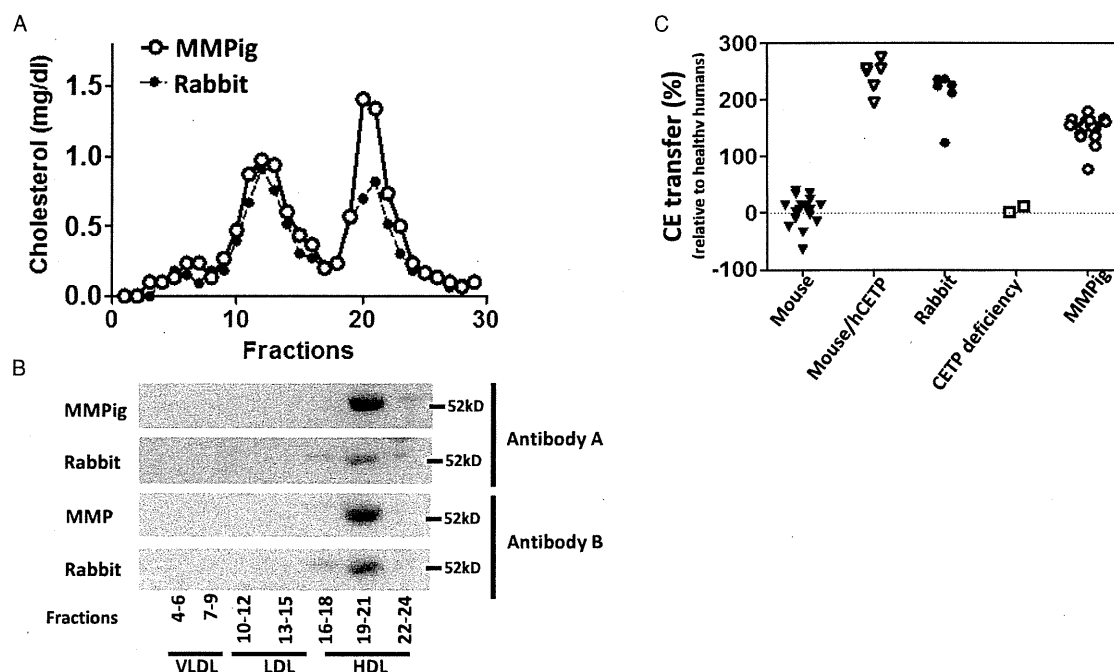
##### *Cloning and Generation of Recombinant Adenoviruses Encoding for Human CETP*

A recombinant adenovirus expressing human CETP (Ad-hCETP) was produced using the ViraPower Adenoviral Expression System (Invitrogen, Carlsbad, CA, USA), according to the manufacturer's instructions. Briefly, an entry clone of the Gateway system (Invitrogen) was generated by cloning the open reading frame into a pENTR/D-TOPO vector (Invitrogen) using first-strand complementary DNA derived from human liver RNA (Clontech, Pao Alto, CA) as a template. The specific primers were as follows: forward: 5'-CAC CAT GCT GGC TGC CAC AGT -3'; reverse: 5'-CTA GCT CAA GCT CTG GAG GAA A -3'. An expression clone for the adenoviral vector was then generated via an LR recombination reaction between the entry clone and a pAd/CMV/V5-DEST (Invitrogen), according to the manufacturer's protocol.

**Supplemental Table 1.** Genes investigated and primers used for PCR

Gene	Assay ID	Custom made primers	RefSeq	GenBank
LDLr	Ss03374441_u1		NM_001206354.1	AF065990.1
SR-BI	Ss03391104_m1		NM_213967.1	AF467889.2
HMGCR	Ss03390147_m1		NM_001122988.1	DQ432054.1
SREBP-2	Ss03376492_u1			AY493571.1
APOBEC-1		Forward: CCATGGTGTGACCATCCAGAT Reverse: AGTTGACAAAATTCCTCCAGCAGTA Probe: TGGGAGCCCCAGAGTA	XM_003126519.1	
NPC1L1		Forward: CCTGTTCCGGAGCGAGTCTCTA Reverse: GAAAGAGGAAATAGTCGAGCAGGTA Probe: CTGCCCAAGGACTC	XM_003134893.1	
GAPDH	Ss03375435_u1		NM_001206359.1	AY307771.1

The assay IDs are listed for the predesigned gene expression assay (ABI). The primers and probes for the APOBEC-1 and NPC1L1 analyses were created based on swine sequences.



**Supplemental Fig. 1.** MMPig sera activate CE transfer from HDL to LDL

(A) The cholesterol levels in the MMPig and rabbit lipoprotein fractions (pooled from six animals each) isolated using FPLC. (B) A Western blot analysis using two antibodies to detect the CETP expression in the MMPig and rabbit lipoprotein fractions, as described in the Methods. (C) The cholesteryl ester (CE) transfer activity in the sera obtained from C57BL/6 mice ( $n=18$ ), C57BL/6 mice injected with Ad-hCETP ( $n=6$ ), rabbits ( $n=6$ ), subjects with CETP deficiency ( $n=2$ ) and MMPigs ( $n=18$ ).

The recombinant adenoviral plasmid was purified and then transfected into 293A cells. The adenovirus was purified using the Adeno-X Virus Purification Kit (Clontech), after a sufficient cytopathic effect was observed in the cells. The multiplicity of infection (MOI) was defined as the ratio of the total number of plaque-forming units to the total number of cells infected.

#### *Injection of the Adenoviral Vector and Blood Sampling*

C57BL/6 mice were obtained from Clea Japan (Tokyo, Japan) and handled according to the guidelines of the National Defense Medical College Institutional Animal Care and Use Committee. The mice were injected with Ad-hCETP ( $5 \times 10^8$  pfu) via the tail vein. Five days after injection, the mice were exsanguinated, and blood samples were obtained.

#### *Determination of the CETP Activity*

The serum CETP activity was determined according to the method of Cheung *et al.*<sup>3)</sup> with a minor modification. In brief, the  $d > 1.125$  fraction isolated via ultracentrifugation from pooled healthy human plasma ( $n=6$ , fasting) was incubated with

$^{14}\text{C}$ -cholesteryl oleate (Perkin-Elmer) at  $37^\circ\text{C}$  for 16 hours, after which the  $1.125 < d < 1.21$  fraction was isolated via ultracentrifugation. Following dialysis, the HDL fractions were incubated with human LDL isolated from pooled human plasma ( $n=6$ , fasting) in the presence or absence of serum obtained from MMPigs, rabbits, mice or humans. Ten hours after incubation, LDL was precipitated by adding heparin (500 U/mL) and manganese chloride ( $0.2 \mu\text{mol/L}$ ) to the samples. The radiotracer counts in the supernatants were determined using a scintillation counter. The percentage cholesteryl ester transfer was calculated by dividing the differences between the values obtained in the absence and presence of sera by the values obtained in the absence of sera. The data are expressed as the CE transfer relative to the mean value of sera obtained from eight healthy humans.

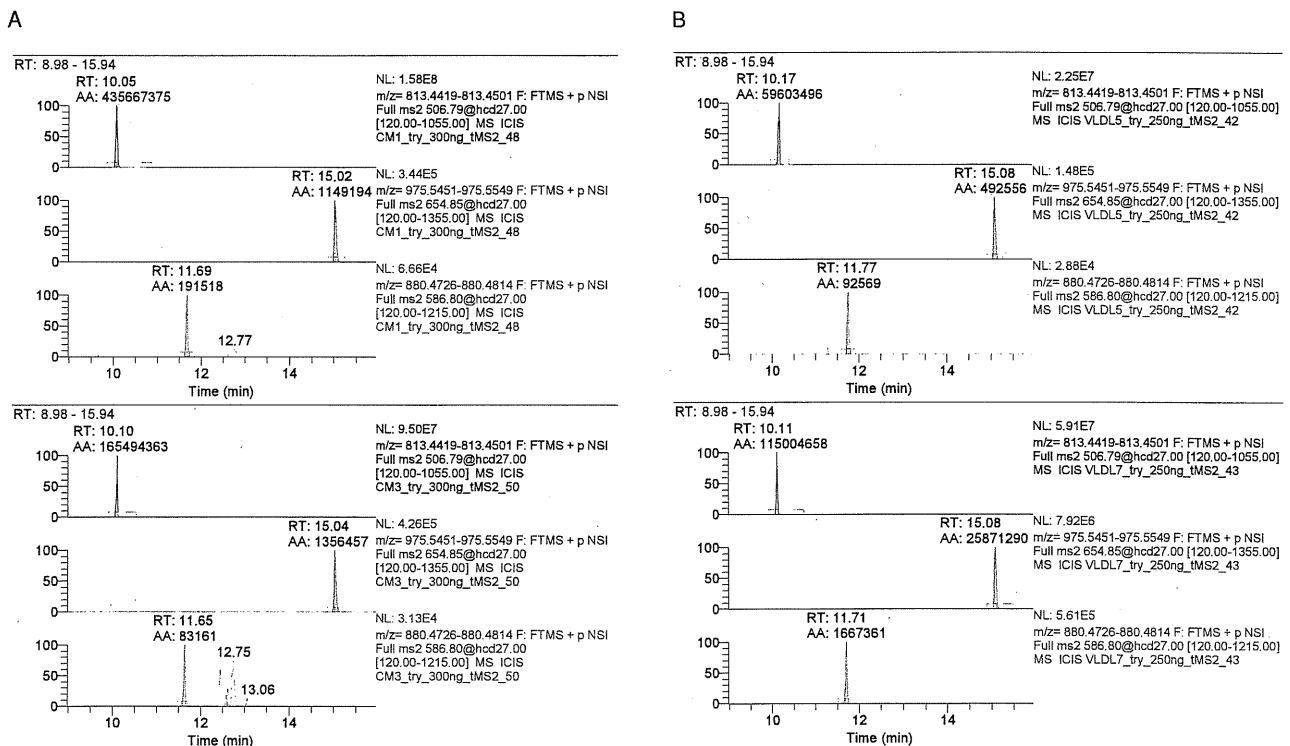
#### **MS/MS Analysis of Apolipoprotein A1 and B**

To detect apolipoprotein (apo)A1, B48 and B100 in the plasma, we utilized a HPLC/MS/MS-based method that can be applied across species. A multiple reaction monitoring (MRM) method was developed for apoA1 and apoB48/100<sup>4)</sup>. First, the chylomicron

**Supplemental Table 2.** MRM transitions for the targeted peptides

	m/z (precursor, z=2)	m/z (product)
Apolipoprotein A1		
AKPALEDLR	506.7929	813.446
Apolipoprotein B48/100		
GFPEPTLEALFGK	654.8453	975.55
YENYELTLK	586.7953	880.477

Thermo tube lens = 120 for all precursor ions. Z=2 for all precursor ions

**Supplemental Fig. 2.**

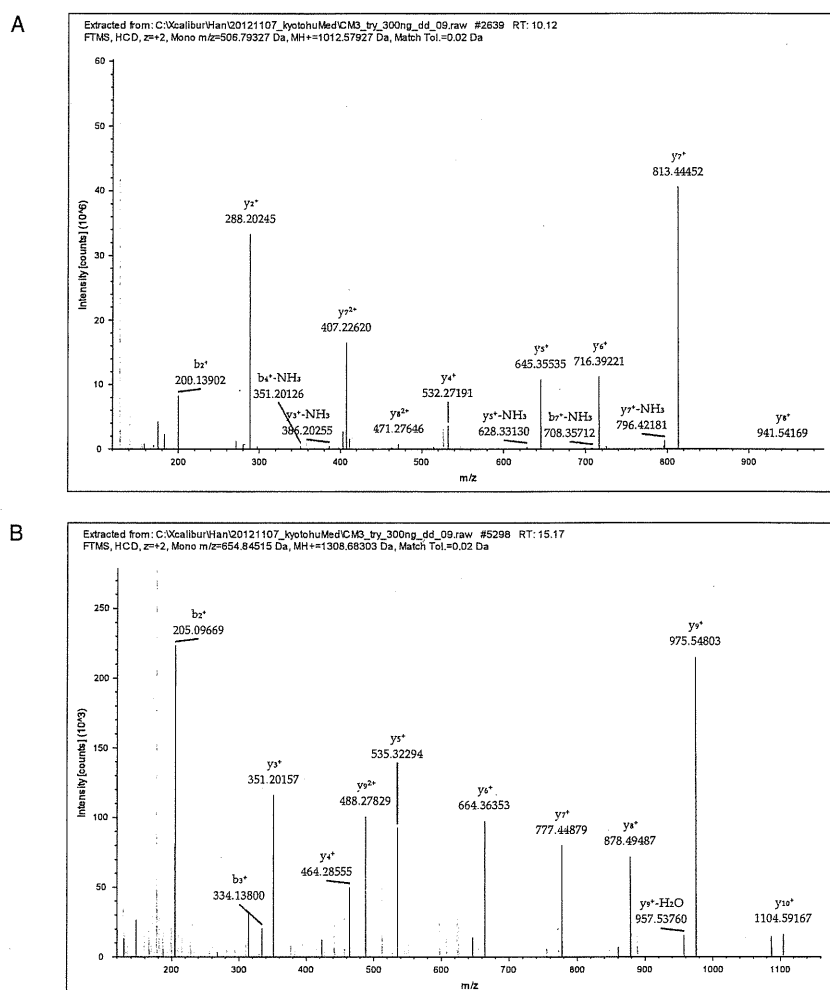
Extracted MRM chromatograms for ApoA1 (AKPALEDLR) and ApoB48/100 (GFPEPTLEALFGK, YENYELTLK) from a nanoHPLC-MS/MS analysis of digested human (A) and MMPig (B) plasma fractions, chylomicron (CM) and very-low-density lipoprotein (VLDL).

(CM) and very-low-density lipoprotein (VLDL) fractions were separated from human and MMPig plasma using the Himac Centrifuge (CP70MX, Himac Centrifuge, Hitachi Medical Systems), according to the manufacturer's protocol. The samples were alkylated and digested overnight with trypsin and analyzed on a nano HPLC system (Easy-nLC, ThermoScientific) coupled with a triple quadrupole mass spectrometer (Orbitrap Q Exactive, ThermoScientific). The trap column was the EASY column, C18, 0.1 × 2 cm, 5 μm (ThermoScientific), and the analytical column was the Tip column, ODS, 0.075 × 120 mm (Nikkyo Technos). The flow rate was 0.3 mL/min.

## Results

### CETP Assay

**Supplemental Fig. 1A** shows the cholesterol distribution in the lipoprotein subfractions isolated using FPLC. The lipoprotein profiles of the MMPs were similar to those of rabbits, except for the higher HDL cholesterol levels. We performed a Western blot analysis using the FPLC subfractions to investigate whether the CETP expression was evident in the MMP. Two antibodies against CETP enabled the visualization of bands of similar molecular size in the HDL fractions obtained from both MMPigs and rabbits, as previ-



**Supplemental Fig. 3.**

Sequencing of the tandem mass spectrum of (C) a peptide with 506.8 m/z obtained from the tryptic digest of MMPig CM proteins and (D) a peptide with 654.8 m/z obtained from the tryptic digest of MMPig CM proteins.

ously demonstrated in humans<sup>5</sup>) and hamsters<sup>6</sup>) (**Supplemental Fig. 1B**). We next assessed whether sera obtained from MMPigs activated CE transfer from HDL to LDL. We confirmed that the sera obtained from C57BL/6 wild-type mice do not possess a CE transfer activity and that the intravenous injection of Ad-hCETP results in a 2.4-fold increase in the activity compared with that observed in human sera (**Supplemental Fig. 1C**). To further validate the reliability of our CETP activity assay, we measured the CE transfer activity in the sera obtained from the subjects with CETP deficiency. The CETP mass in the sera of the 59-year-old man (HDL-C, 172 mg/dL) and 51-year-old woman (191 mg/dL) was <0.1 and 0.4  $\mu$ g/mL,

respectively (normal range, 1.04-3.55  $\mu$ g/mL). A genetic analysis revealed that the man carried the homozygous intronic mutation Int14 +1 G>A, while the woman carried the Int14 +1 G>A/D442G mutation as a compound heterozygote. We also confirmed the absence of the CE transfer activity in the sera of both subjects. Finally, we observed a robust CE transfer activity in the MMPig sera, 1.5-fold higher than that observed in the humans and 0.7-fold lower than that observed in the rabbits, supporting the possible expression of CETP.

### Identification of ApoA1- and ApoB48/100-Specific Peptides and MRM Transitions

**Supplemental Table 2** shows the identified peptide sequence along with the corresponding experimentally determined MRM transitions. **Supplemental Fig. 2A** and **2B** show MRM chromatograms extracted from both the human and pig plasma fractions, CM and VLDL. The amino acid sequences for apoA1 and B48/100 in the humans and pigs were obtained with Proteome Discoverer software program using a Swiss-Prot-TrEMBL database. The sequencing of the tandem mass spectrum of peptides with 506.8 and 654.8 m/z from the tryptic digest of porcine CM proteins was AKPALEDLR and GFPTLEALFGK, corresponding to peptides of porcine apoA1 (**Supplemental Fig. 3A**) and apoB48/100 (**Supplemental Fig. 3B**), respectively.

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