# **Supplemental Figure 1**



### Supplemental Figure 1: Bioengineering 3D human arteries.

(a) 3D bioengineered human vessels were subjected to abluminal A $\beta$ 42 treatment with or without luminal HDL treatment. Histological characterization of the vessels shows (b) a monolayer of endothelial cells on the luminal aspect visualized with CD31 and (c) multiple layers of smooth muscle cells throughout the vessel visualized with smooth muscle actin (SMA). CD31: cluster of differentiation 31, SMA: smooth muscle actin, DAPI: 4',6-di-amidino-2-phenylindole.

## **Supplemental Figure 2**



# Supplemental Figure 2: *HDL isolation and depletion of apoB-containing lipoproteins from human plasma.*

HDL or apoB-depleted plasma was prepared from young, healthy donor plasma by (**a**) sequential density gradient ultracentrifugation (HDL) using KBr, (**b**) polyethylene glycol (PEG-P) precipitation of non-HDL lipoproteins, (**c**) a combination of the two methods (PEG-UC), or (**d**) a single UC spin (UC-P). KBr: potassium bromide, LDL: low-density lipoprotein, VLDL: very low-density lipoprotein, HDL: high-density lipoprotein.

## **Supplemental Figure 3**



Total denatured protein

MW (kDa) LDL HDL PEG-P Plasma 250 150 100 75 50 37 25 20 15

#### Supplemental Figure 3: Composition and cholesterol efflux function of HDL and apoB-depleted plasma.

The levels of (a) HDL-C, (b) apoA-I and (c) apoE, (d) phospholipid, and (e) cholesterol ester levels were measured in unfractionated plasma, HDL isolated from plasma by ultracentrifugation, and apoB-depleted plasma produced by PEG precipitation and normalized to HDL-C concentration. (f) Cholesterol efflux capacity was measured using RAW 264.7 macrophages loaded with [3]H-cholesterol. (g) Total protein was observed in HDL using SDS-PAGE denaturing gels stained with Coomassie Brilliant Blue. Scatter plots represent independent experiments with mean ±95% confidence interval. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, or exact p-values by one-way analysis of variance (ANOVA) with Tukey's multiple comparisons test. MW: molecular weight, d: diameter, kDa: kilodalton, LDL: low-density lipoprotein, HDL: high-density lipoproteins isolated by sequential density gradient ultracentrifugation, PEG-P: apoB-depleted plasma by polyethylene glycol precipitation, A.U.: arbitrary units, SDS-PAGE: sodium dodecyl sulphate polyacrylamide gel electrophoresis.

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