

SUPPLEMENTAL MATERIAL

Targeted deletion of hepatocyte Abca1 increases plasma HDL reverse cholesterol transport via the LDL receptor

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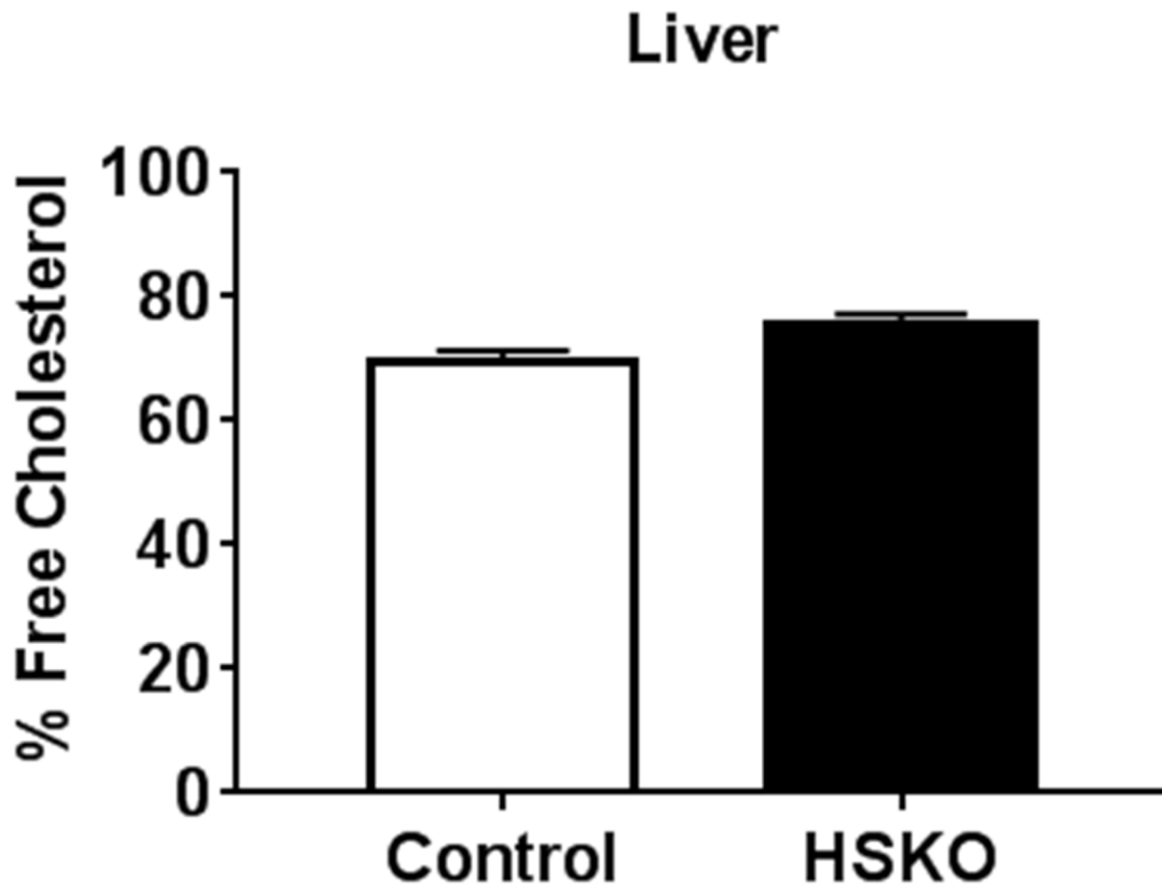
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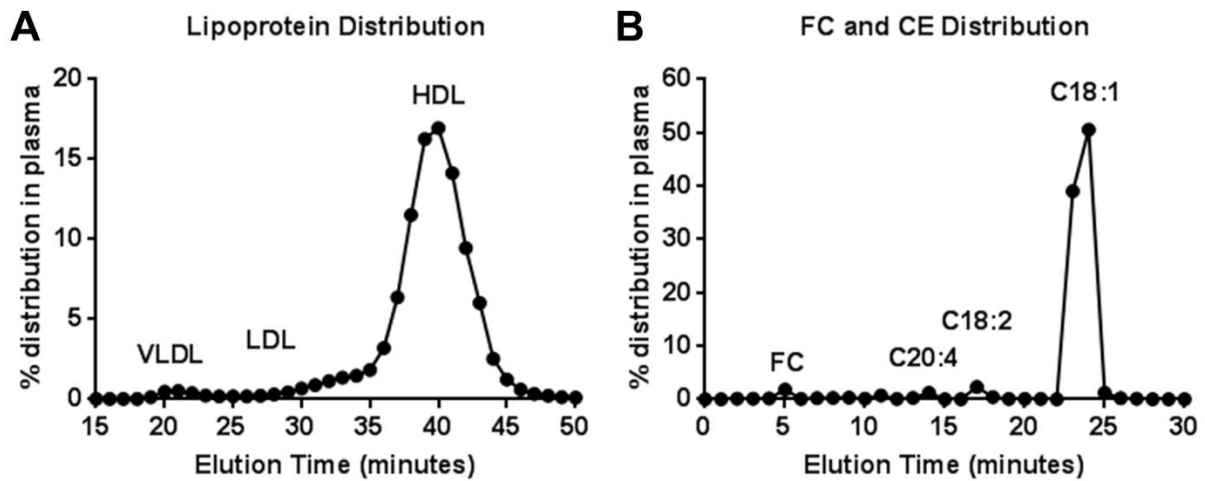
Short title: Liver Abca1 and HDL reverse cholesterol transport

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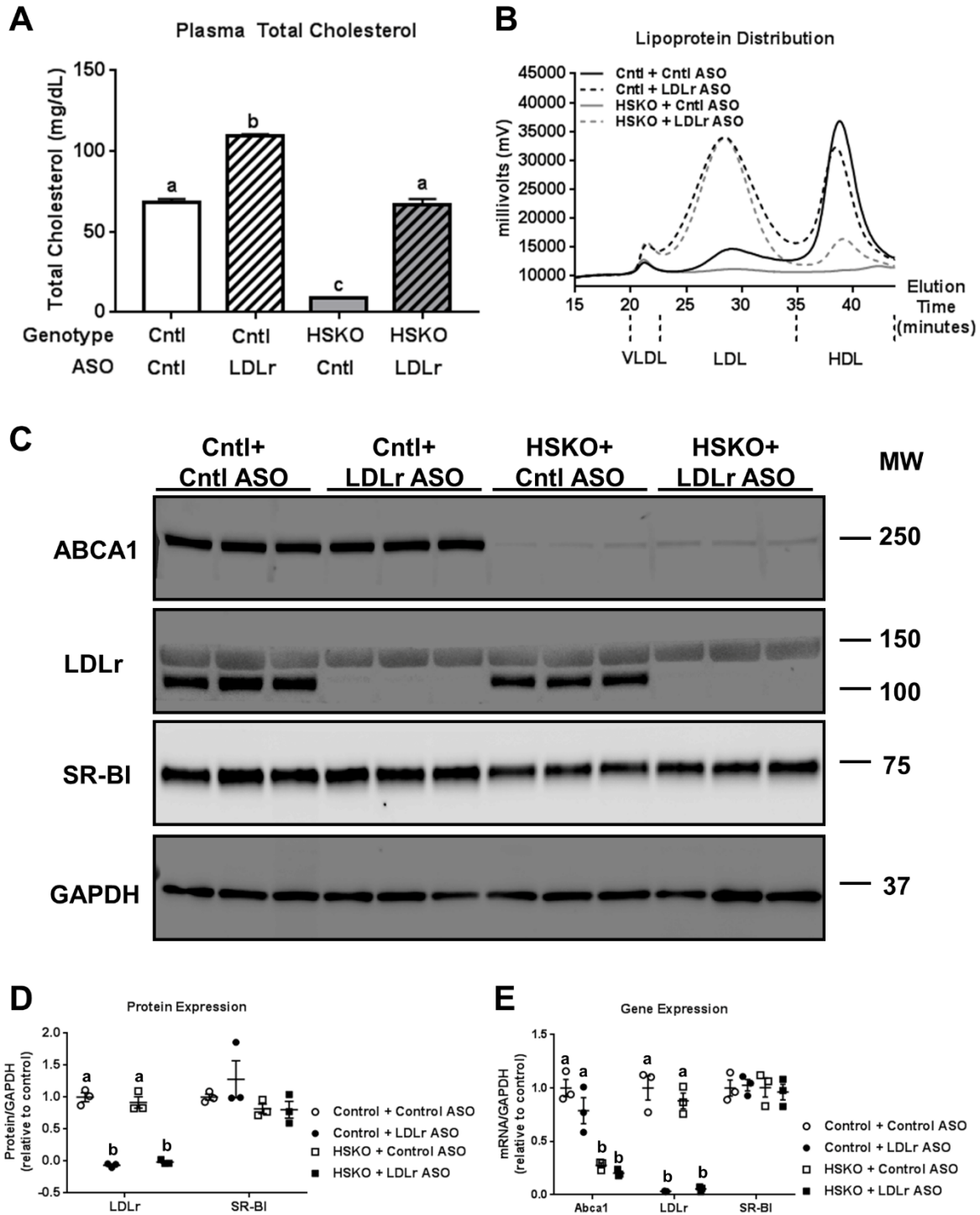
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Supplemental Figure I. Hydrolysis of ^3H -CO human HDL tracer to ^3H -FC in liver. Percentage ^3H -FC in liver 24 hours following ^3H -CO human HDL injection into control or HSKO mice was determined after lipid extraction of liver, thin-layer chromatographic separation of FC and CE, and quantification of radiolabel by liquid scintillation counting. Approximately 70% of the plasma HDL-derived radioactivity was FC in both genotypes of mice, indicating significant hepatic hydrolysis of ^3H -CO to ^3H -FC. Data are mean \pm SEM; n=5/genotype.

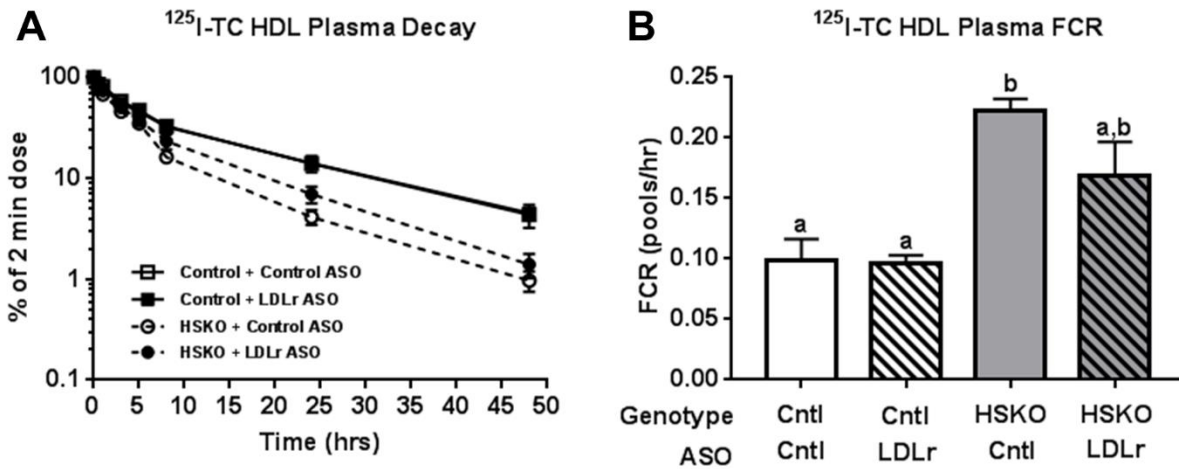


Supplemental Figure II. $^3\text{H-CO}$ ($^3\text{H-C18:1}$) HDL stability in plasma. $^3\text{H-CO}$ radiolabeled human HDL was incubated with plasma at 37°C for 2 min up to 24 hours. Plasma was then fractionated by FPLC to determine ^3H lipoprotein distribution (**A**). The plasma samples were then lipid extracted and $^3\text{H-FC}$ and $^3\text{H CE}$ fatty acyl distribution was analyzed by reverse phase HPLC (**B**). Only data from the 24-hour incubation are shown. Nearly all (95%) of the $^3\text{H-CO}$ HDL tracer eluted in the HDL region of the FPLC column. After a 24-hour incubation of the $^3\text{H-CO}$ HDL tracer with plasma at 37°C , 94% of the radiolabel eluted in HDL region, suggesting minimal transfer of radiolabel to other plasma lipoproteins (A). In the 2-min incubation, 96.5% of the HDL radiolabel eluted in the CO peak compared with 89.4% in the 24-hour time point, indicating minimal hydrolysis of the $^3\text{H-CO}$ to $^3\text{H-FC}$ over the 24-hour plasma incubation.

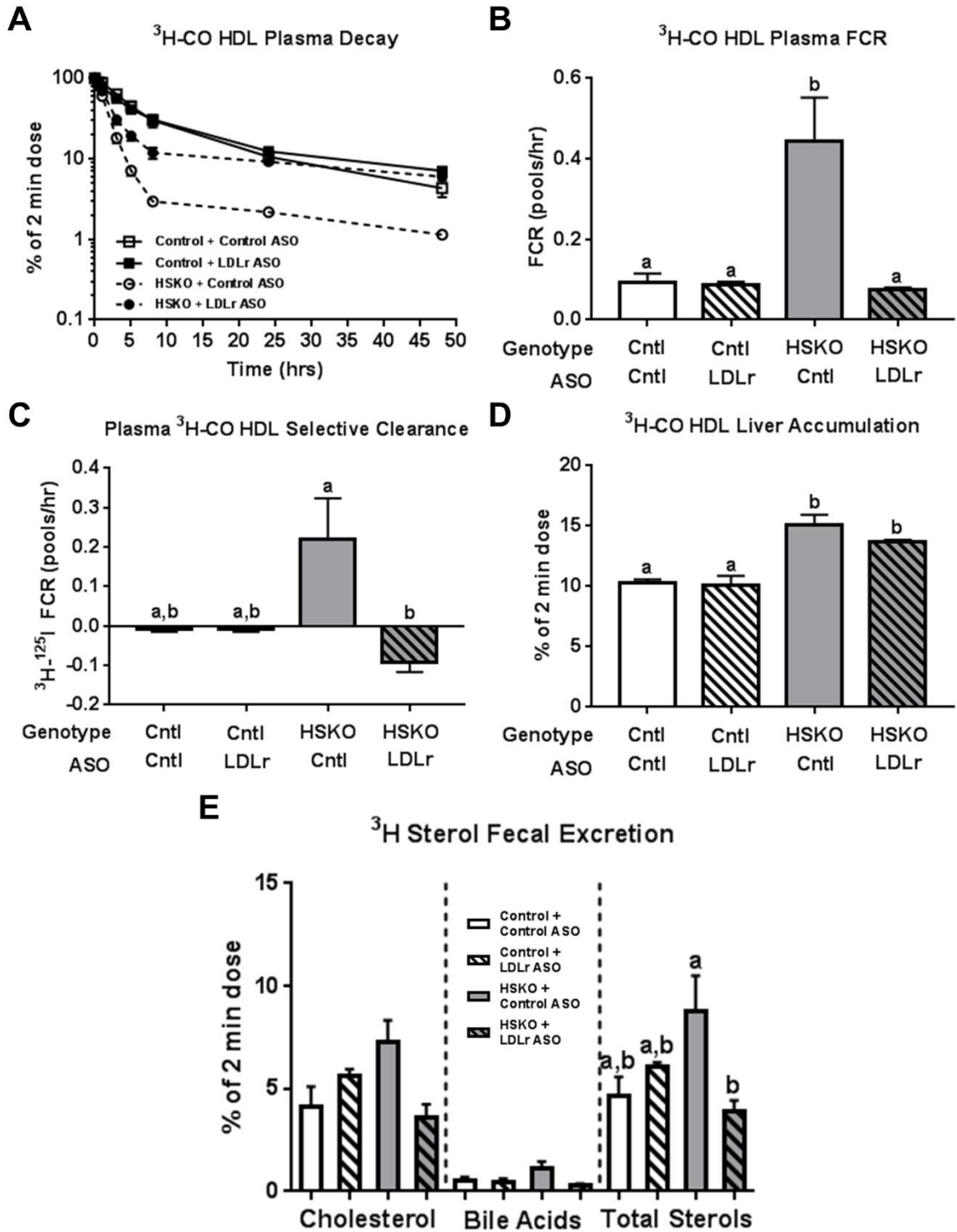


Supplemental Figure III. LDLr ASO treatment increases plasma LDL-C and HDL-C in HSKO mice. Efficiency of LDLr silencing was monitored by quantifying total plasma cholesterol (**A**) and

plasma lipoprotein cholesterol distribution by FPLC (mV signal from online cholesterol analyzer) **(B)** following 4 weeks of ASO treatment. Whole liver lysates were immunoblotted for Abca1, LDLr, and SR-BI **(C)** and blots were quantified **(D)** by calculating fold change in protein/GAPDH ratio relative to control mice treated with control ASO. Real-time PCR was performed to analyze gene expression for Abca1, LDLr, and SR-BI and the mRNA/GAPDH ratio relative to control mice treated with control ASO was plotted **(E)**. Data are mean \pm SEM; n=3 for each group. Groups with different letters are statistically different. Protein and gene expression data are from recipient mice used in mouse HDL tracer studies (Supplemental Figures IV and V).



Supplemental Figure IV. Effect of LDLr ASO treatment on *in vivo* catabolism of ¹²⁵I-TC labeled mouse HDL. ¹²⁵I-TC labeled HDL was injected intravenously in control and HSKO treated with either a control or LDLr targeting ASO. Periodic blood samples were taken over 48 hours to analyze plasma decay (**A**) and FCR (**B**). Data are mean \pm SEM. Groups with different superscripts are statistically different ($p < 0.05$), $n = 3$ for each group. Control turnover curves in panel A are identical and SEM in nearly all points falls within the symbol.



Supplemental Figure V. Effect of LDLr ASO treatment on *in vivo* catabolism of $^3\text{H-CO}$ radiolabeled mouse HDL. $^3\text{H-CO}$ radiolabeled HDL was injected intravenously in control and HSKO mice treated with either a control or LDLr targeting ASO for 4 weeks. Periodic blood

samples were taken over 48 hours to analyze plasma decay (**A**), plasma FCR (**B**), and plasma HDL ³H-CO selective clearance, using data from Supplemental Figure IVB for ¹²⁵I-TC FCR (**C**). At the termination of the study, tissues were harvested to quantify liver accumulation (**D**) and fecal excretion (**E**) of the tracer. Data are mean ± SEM. Groups with different superscripts are statistically different (p<0.05), n=3 for each group.