#### **SUPPLEMENTAL MATERIALS**

Title: Scavenger receptor BI, not LDL receptor, mediates adrenal stress response

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Running title: HDL/SR-BI mediates adrenal stress response

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## **Major Resources Table**

In order to allow validation and replication of experiments, all essential research materials listed in the Methods should be included in the Major Resources Table below. Authors are encouraged to use public repositories for protocols, data, code, and other materials and provide persistent identifiers and/or links to repositories when available. Authors may add or delete rows as needed.

#### **Animals (in vivo studies)**

Species	Vendor or Source	Background Strain	Sex	Persistent ID / URL
SR-BI+/- (B6;129S- Scarb1tm1Kri/J)	The Jackson Laboratory	129S2/SvPas	Male and female	https://www.jax.org/strain/003379 Stock No:003379  SR-BI KO
ApoB100	Taconic Biosciences	C57BL/6NTac	Male and female	https://www.taconic.com/transgenic- mouse-model/apob100 Model # 1004

## **Genetically Modified Animals**

onousuny mountou / minuto						
SR-BI <sup>-/-</sup>	Species	Vendor or	Background	Other	Persistent ID / URL	
		Source	Strain	Information		
Parent -	SR-BI+/-	The	129S2/SvPas		https://www.jax.org/strain/003379	
Male	(B6;129S-	Jackson			Stock No:003379  SR-BI KO	
	Scarb1tm1Kri/J)	Laboratory			·	
Parent -	SR-BI+/-	The	129S2/SvPas		https://www.jax.org/strain/003379	
Female	(B6;129S-	Jackson			Stock No:003379  SR-BI KO	
	Scarb1tm1Kri/J)	Laboratory				

SRBI <sup>-/-</sup>	Species	Vendor or	Background	Other	Persistent ID / URL
ApoBtg		Source	Strain	Information	
Parent -	SRBI-/-	The	129S2/SvPas		https://www.jax.org/strain/003379
Male		Jackson			
		Laboratory			
Parent -	ApoB100	Taconic	C57BL/6NTac		https://www.taconic.com/transgenic-
Female		Biosciences			mouse-model/apob100
					Model # 1004

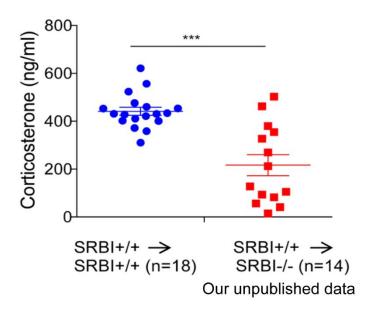
# LDL\_ratio

```
# fraction numbers
x <- c(6:31)
# control FPLC
\texttt{y1} \leftarrow \texttt{c(1.3615, 1.5565, 1.218, 0.8905, 0.6445, 0.3585, 0.0645, 0.171, 0.984, 0.8025, 1.492, 3.9, 9.955, 15.414, 19.73, 19.73, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.74, 19.
# LDL high mice
y2 <- c(0.3185,0.612,0.3895,0.5325,0.6225,0.572,1.948,3.257,6.176,7.9435,8.616,8.0365,8.9925,10.9915,13
## Loading required package: xts
## Loading required package: zoo
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
                  as.Date, as.Date.numeric
## Registered S3 method overwritten by 'xts':
## method from
## as.zoo.xts zoo
## Loading required package: TTR
## Registered S3 method overwritten by 'quantmod':
           method
## as.zoo.data.frame zoo
## Version 0.4-0 included new data defaults. See ?getSymbols.
## Loading required package: ALS
## Loading required package: nnls
## Loading required package: Iso
## Iso 0.0-18
## Loading required package: ptw
                   rt
                                          sd FWHM height
                                                                                                   area
## [1,] 3 1.273983 3 1.2180 3.889562
## [2,] 10 7.219235 17 0.8025 14.521991
## [3,] 16 2.123305 5 18.8785 100.477703
                                           sd FWHM height
                 rt
                                                                                                    area
## [1,] 3 10.616523 25 0.3895 10.36525
## [2,] 6 10.616523 25 0.5720 15.22188
## [3,] 12 4.671270 11 8.0365 94.10048
## [4,] 17 3.821948
                                                         9 14.1130 135.20541
LDL_ratio <- fit2[3, 5]/fit1[2, 5]
HDL_ratio <- fit2[4, 5]/fit1[3, 5]
LDL_ratio
```

### Caculation of LDL AUC

```
# fraction numbers
x < -c(6:31)
# SRBI+/+ApoBwt
y1 \leftarrow c(0,0.206,0.251,0.096,0,0.206,0.206,0.008,0,0,0,0,0.008,0.273,0.538,1.178,2.304,1.928,0.825,0.58,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.008,0.00
# SRBI+/+ApoBtq
y2 <- c(0,0,0,0,0.21,0.33,0.21,0,0.067,0.833,2.269,3.179,2.581,1.862,1.168,0.833,1.36,2.485,2.126,0.857
# SRBI-/-ApoBwt
y3 \leftarrow c(1.473, 1.553, 1.315, 0.788, 0.735, 0.498, 0, 0, 0, 0, 0.419, 2.502, 9.173, 15.606, 20.432, 18.823, 15.606, 11.09
# SRBI-/-ApoBtg
y4 <- c(0.366,1.263,0.946,1.078,1.052,1.157,2.027,2.634,5.534,7.986,9.357,8.223,8.434,9.519,12.68,13.89
## Loading required package: xts
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
                  as.Date, as.Date.numeric
## Loading required package: TTR
## Registered S3 method overwritten by 'quantmod':
##
         method
         as.zoo.data.frame zoo
## Version 0.4-0 included new data defaults. See ?getSymbols.
## Loading required package: ALS
## Loading required package: nnls
## Loading required package: Iso
## Iso 0.0-18
## Loading required package: ptw
##
                                              sd FWHM height
                rt
                                                                                                area
## [1,] NA
                                            NA NA NA
## [2,] 19 0.8493218 2 1.928 4.104585
                                              sd FWHM height
## [1,] 13 0.8493218 2 2.581 5.494779
## [2,] 19 0.8493218 2 2.126 4.526114
fit3
##
                    rt
                                            sd FWHM height
                                                                                                area
```

# Suppl Fig. 1



Suppl Fig 1. Adult recipient SRBI+/+ or SR-BI-/- mice were bilaterally adrenalectomized and adrenal gland from 9-day old SRBI+/+ mice was transplanted to the recipient mice. 6 weeks later, the mice were treated with cecal ligation and puncture (CLP) and the plasma corticosterone concentrations were quantified 4h post CLP.