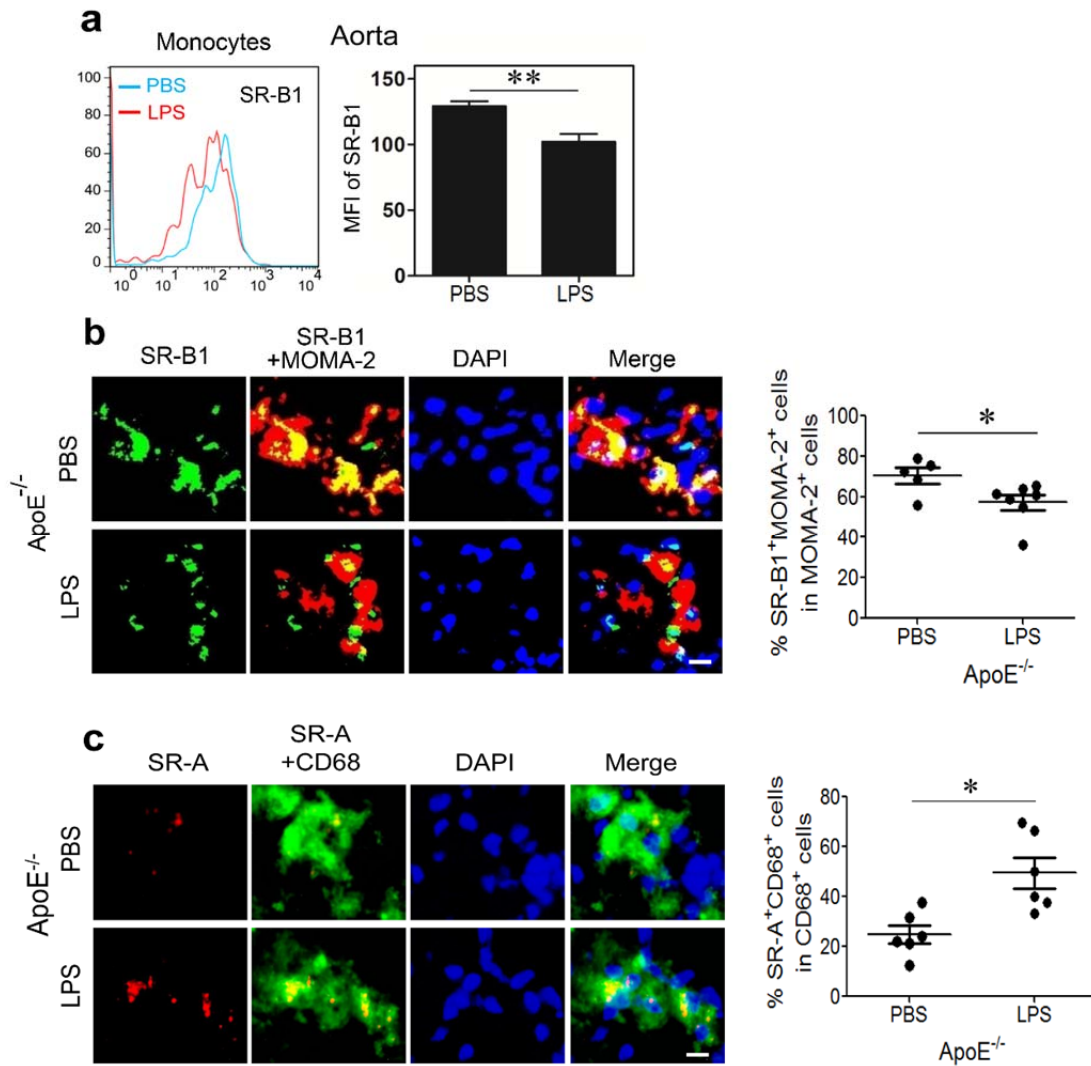


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**Supplementary Fig 1. Subclinical endotoxemia reduces plaque smooth muscle actin (SMA) levels.** ApoE<sup>-/-</sup> mice were pre-conditioned with PBS or super-low dose LPS for 4 weeks together with high fat diet, followed by high fat diet feeding only for an additional 4 weeks. **(a)** Representative images of SMA<sup>+</sup> staining within atherosclerotic plaques of aortic root areas. Scale bar: 300  $\mu$ m. **(b)** Quantification of SMA<sup>+</sup> positive staining areas per mm<sup>2</sup> in the lesion area of aortic root. Data are shown for aortic plaque areas from PBS (n = 7) and super-low dose LPS conditioned (n = 7) mice. Data represent two similar experiments. Error bars show means  $\pm$  s.e.m.; \* P < 0.05; student t-test.

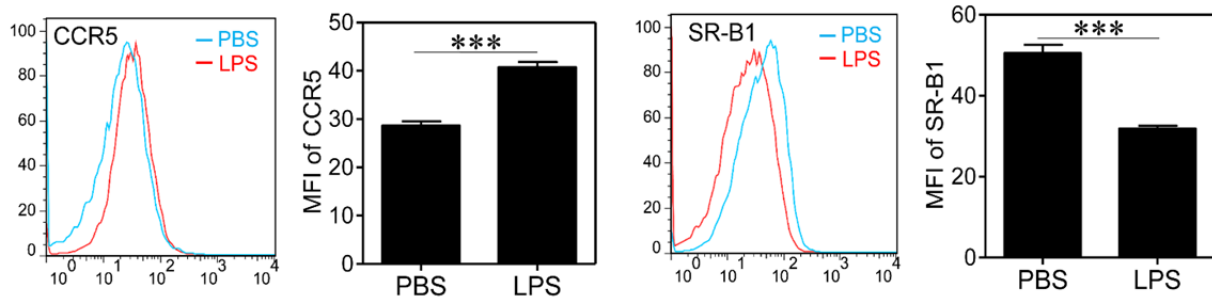
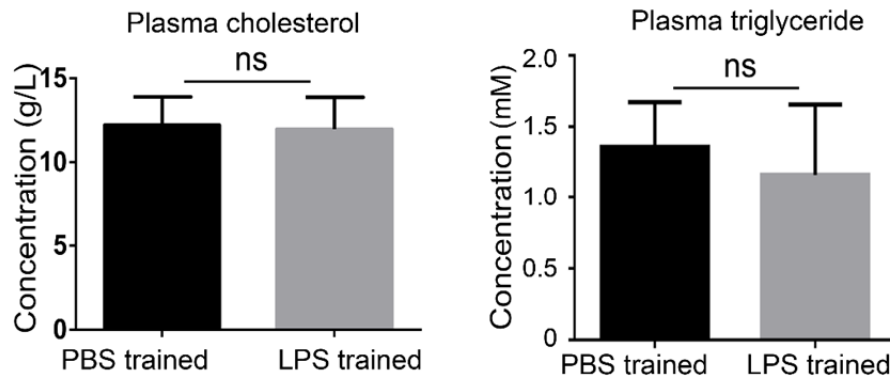
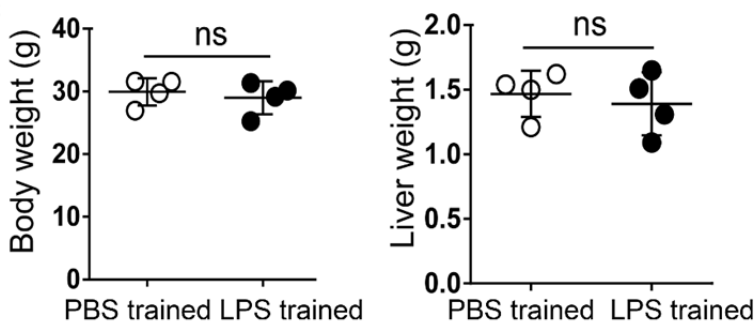
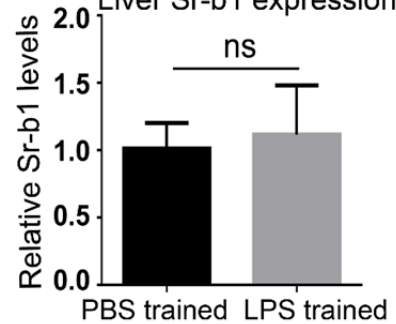


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17 **Supplementary Fig 2. Subclinical endotoxemia increases the expression of SR-A and**  
 18 **reduces the expression of SR-B1.** *ApoE*<sup>-/-</sup> mice were pre-conditioned with PBS or super-low  
 19 dose LPS for 4 weeks together with high fat diet, followed by high fat diet feeding only for an  
 20 additional 4 weeks. (a) Single cell suspensions were prepared from aorta, and the expression  
 21 levels of SR-B1 within monocytes were also analyzed. Data are shown for aortic monocytes from  
 22 PBS (n = 6) and super-low dose LPS conditioned (n = 6) mice. (b) Representative images of SR-  
 23 B1<sup>+</sup>MOMA-2<sup>+</sup> macrophages in the atherosclerotic plaques of aortic root areas. Scale bar: 100  $\mu$ m.  
 24 Quantification of SR-B1<sup>+</sup>MOMA-2<sup>+</sup> area as a percentage of the total atherosclerotic plaque area  
 25 within the aortic root. (c) Representative images of atherosclerotic plaques within aortic root  
 26 areas after staining with anti-SR-A and anti-CD68 antibodies. Scale bar: 100  $\mu$ m. The percentage  
 27 of SRA<sup>+</sup> cells within CD68<sup>+</sup> macrophages was quantified. Data represent two similar  
 28 experiments. Error bars show means  $\pm$  s.e.m.; \* P < 0.05; \*\* P < 0.01; student t-test.

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**a****b****c****d** Liver Sr-b1 expression

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34 **Supplementary Fig 3. Adoptive transfer of monocytes programmed by super-low dose LPS**35 **does not alter plasma levels of lipids or overall body and liver weights.** BM cells from36 *ApoE*<sup>-/-</sup> mice were cultured with M-CSF (10 ng/ml) in the presence of either PBS or LPS (0.1

37 ng/ml) for 5 days. Surface expression levels of CCR5 and SR-B1 on inflammatory monocytes

38 were determined by flow cytometry. (b-c) PBS or LPS programmed (trained) BM cells were then

39 adoptively transferred through *i.v.* injection to HFD-fed *ApoE*<sup>-/-</sup> mice ( $3 \times 10^6$  cells/mouse) once

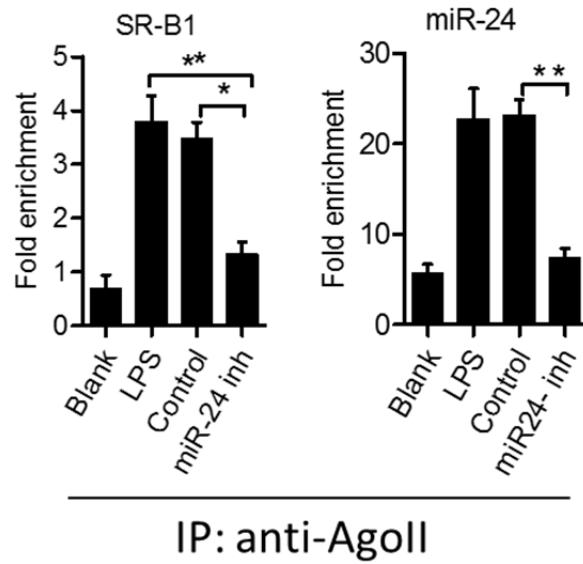
40 a week for 4 weeks. (b) Plasma levels of total cholesterol and triglyceride were measured and

41 quantified. (c) Total body weight and liver weight were quantified. (d) Liver levels of Sr-b1

42 expression were measured by real-time RT-PCR. Data represent two similar experiments. Error

43 bars represent means  $\pm$  s.e.m.; ns: no significance, \*\*\*, P < 0.001, student t-test.

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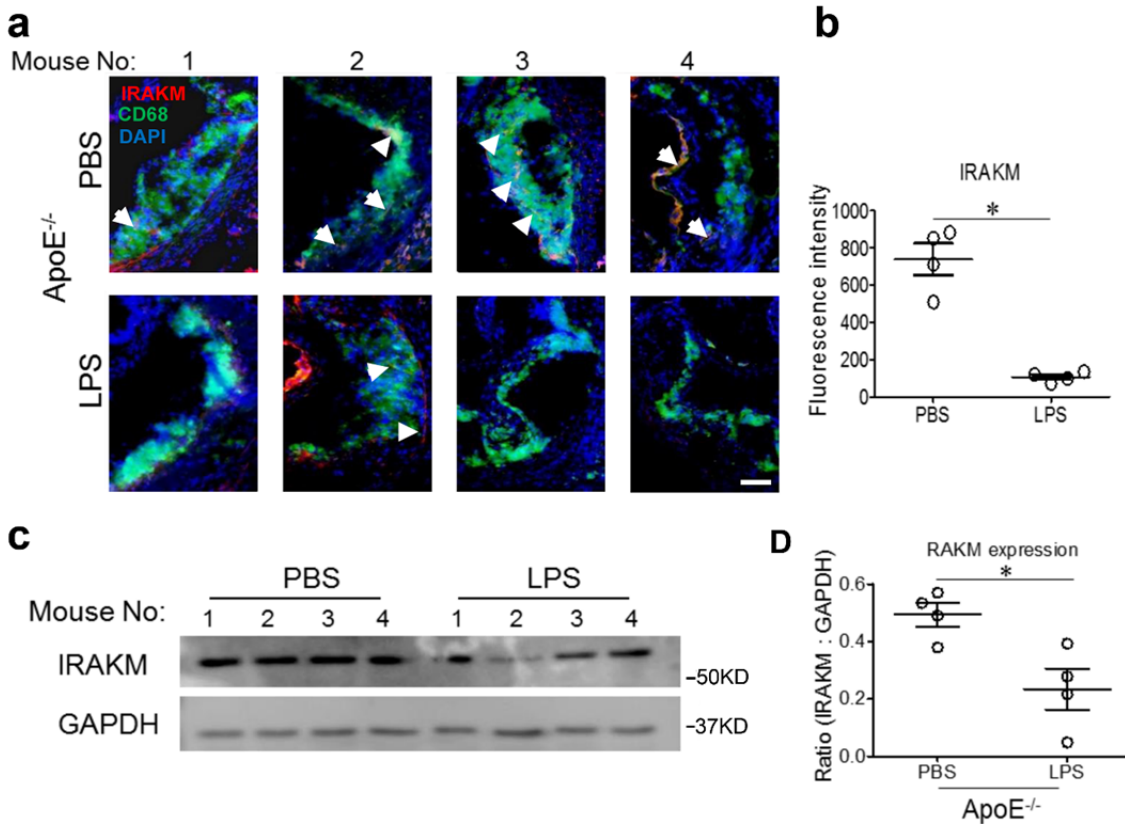
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47 **Supplementary Fig 4. miR-24 forms a close complex with Sr-b1 mRNA as measured by the**  
 48 **RIP assay.** Total RNAs were harvested from LPS treated WT bone marrow macrophages ( $30 \times$   
 49  $10^6$ ) and used for co-immunoprecipitation with an anti-AgoII antibody, in the presence of either  
 50 miR-24 mimics or miR-24 antagomoir . The relative enrichment of Sr-b1 messenger RNAs were  
 51 analyzed by real-time RT-PCR. Results are presentative of three experiments and are expressed  
 52 as fold enrichment relative to AgoII-immunoprecipitation control samples.

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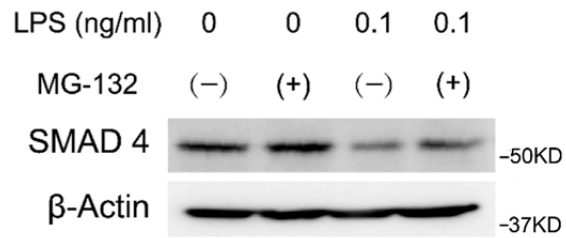
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60 **Supplementary Fig 5. Reduced IRAK-M expression in mice with subclinical endotoxemia.**

61 ApoE<sup>-/-</sup> mice were pre-conditioned with PBS or super-low dose LPS for 4 weeks together with  
62 high fat diet, followed by high fat diet feeding only for an additional 4 weeks. (a) Representative  
63 images of atherosclerotic plaques within aortic root areas after staining with anti-IRAK-M and  
64 anti-CD68 antibodies. Scale bar: 100  $\mu$ m. (b) The fluorescent intensity of IRAK-M was  
65 quantified. Data are shown from PBS (n = 4) and super-low dose LPS conditioned (n = 4) mice.  
66 (c) Western blot analyses of IRAK-M expression in the splenic cells. (D) Relative expression  
67 levels of IRAK-M were quantified. Data are shown from PBS (n = 4) and super-low dose LPS  
68 conditioned (n = 4) mice. Results are presentative of two experiments. Error bars show means  $\pm$   
69 s.e.m.; \*, P < 0.05; student t-test.

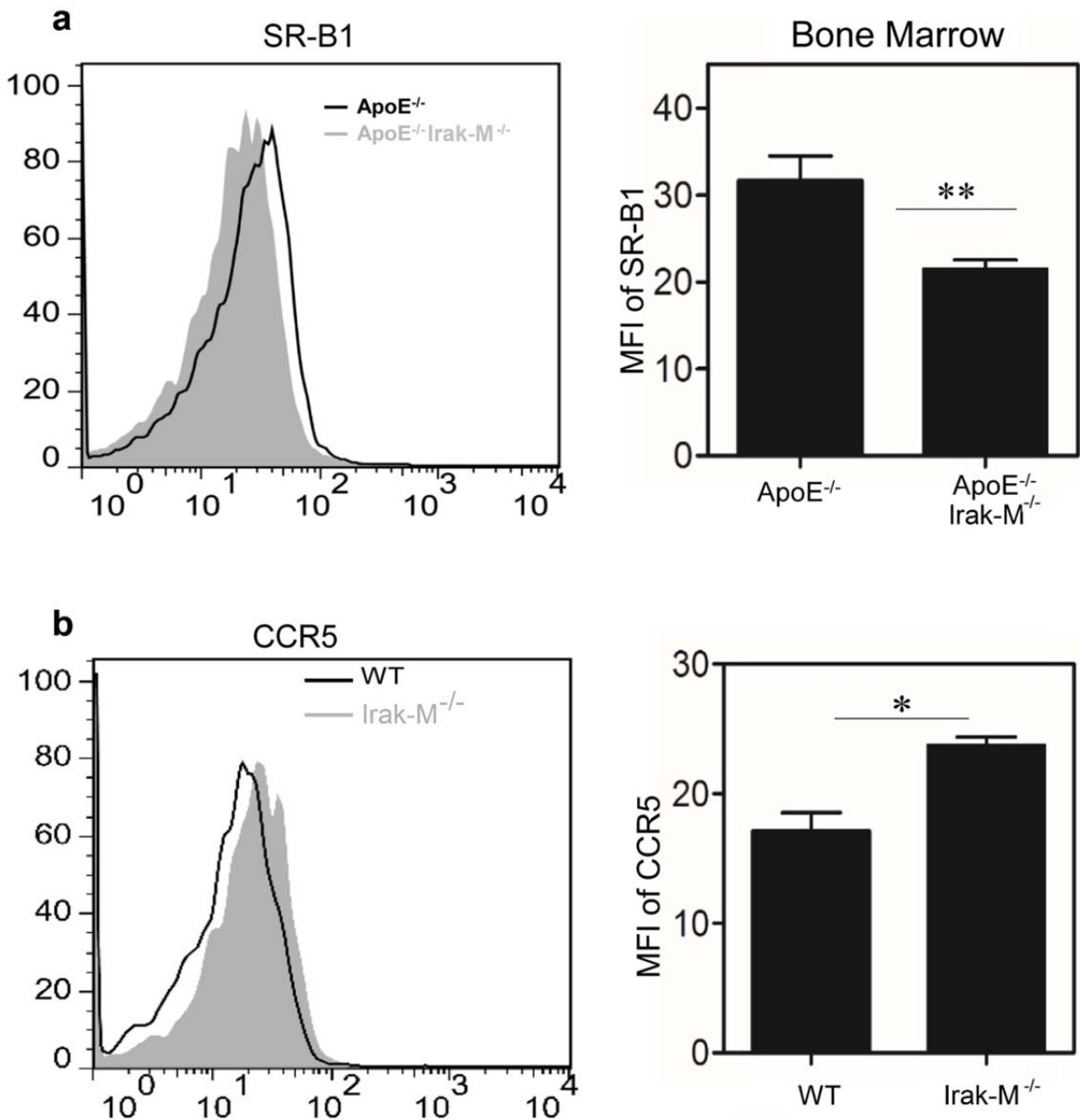
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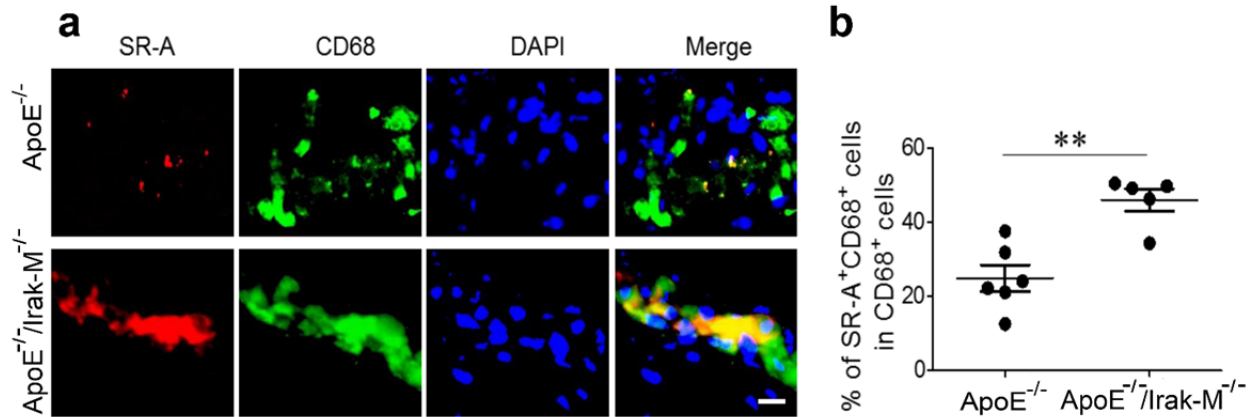
**Supplementary Fig 6. The reduction of Smad4 in monocytes by super-low dose LPS is not caused by protein degradation.** WT BMM were cultured in the presence of absence of 100 pg/ml LPS, together with or without MG-132 (25 nM) for 5 days. The cellular levels of Smad4 were detected by Western Blot. Data represents two similar experiments.



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**Supplementary Fig 7. Disruption of IRAK-M suppresses SR-B1 and up-regulates CCR5.** (a)  $ApoE^{-/-}$  and  $ApoE^{-/-}Irak-M^{-/-}$  mice were fed with HFD for 8 weeks. The expression levels of SR-B1 within BM monocytes were compared between  $ApoE^{-/-}$  (n = 7) and  $ApoE^{-/-}Irak-M^{-/-}$  (n = 6) mice. Error bars show means  $\pm$  s.e.m.; \*\*, P < 0.01; student t-test. (b) BM cells from WT C57 BL/6 mice or  $Irak-M^{-/-}$  mice were cultured with M-CSF (10 ng/ml) in the presence of super-low dose LPS (0.1 ng/ml). On day 5, the expression levels of CCR5 within CD11b<sup>+</sup>Ly6C<sup>++</sup> monocytes were examined. Quantified data are shown from WT and  $Irak-M^{-/-}$  cells (n = 3). Results are representative of three experiments. Error bars show means  $\pm$  s.e.m.; \*, P < 0.05; student t-test.

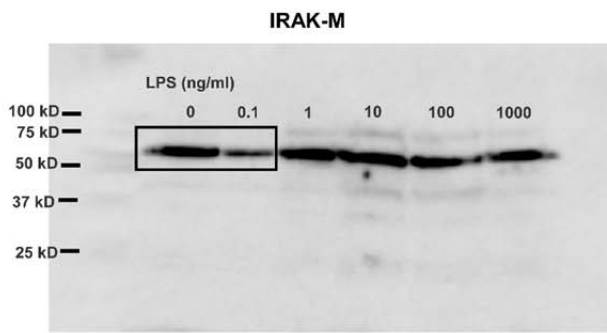
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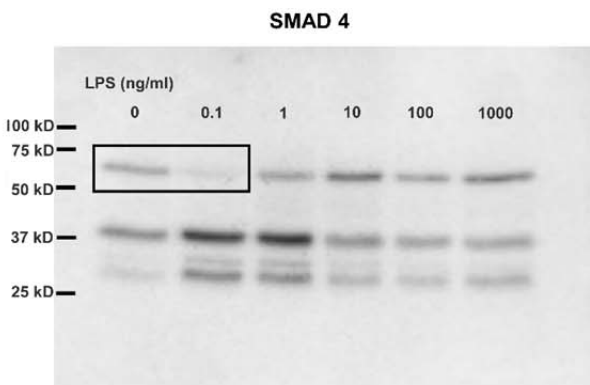
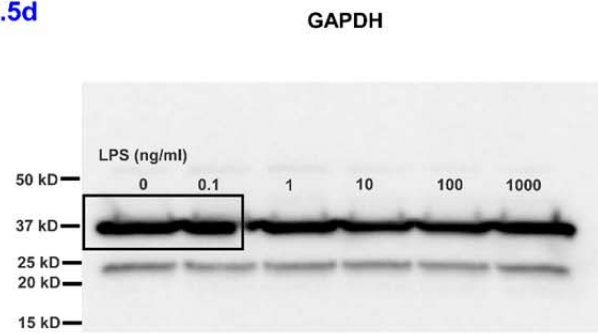
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**Supplementary Fig 8. Disruption of IRAK-M increases SR-A expression of macrophages within lesion area.** *ApoE*<sup>-/-</sup> and *ApoE*<sup>-/-</sup>/*Irak-M*<sup>-/-</sup> mice were fed with HFD for 8 weeks. (a) Representative images of atherosclerotic plaques within aortic root areas after staining with anti-SRA and anti-CD68 antibodies. Scale bar: 100 μm. (b) The percentage of SRA<sup>+</sup> cells within CD68<sup>+</sup> macrophages was quantified. Data are shown from *ApoE*<sup>-/-</sup> (n = 6) and *ApoE*<sup>-/-</sup>/*Irak-M*<sup>-/-</sup> (n = 5) mice. Results are representative of two similar experiments. Error bars show means ± s.e.m.; \*\* P < 0.01; student t-test.

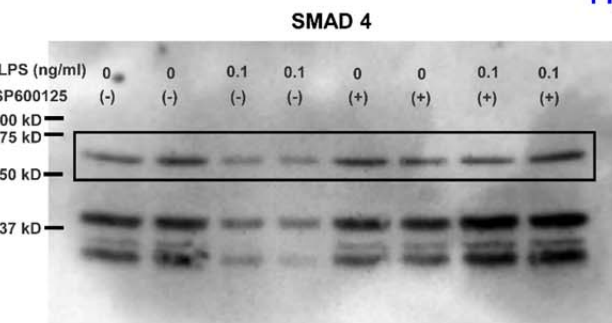
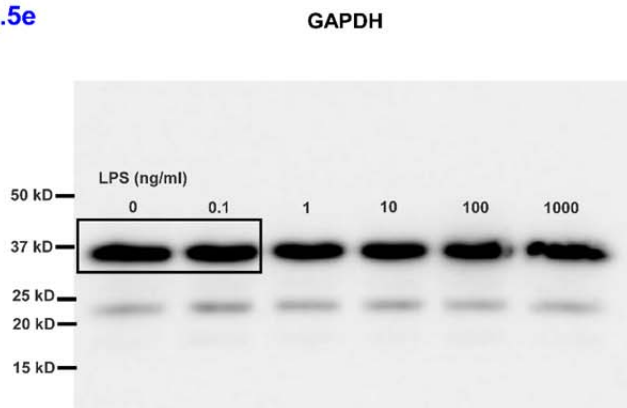




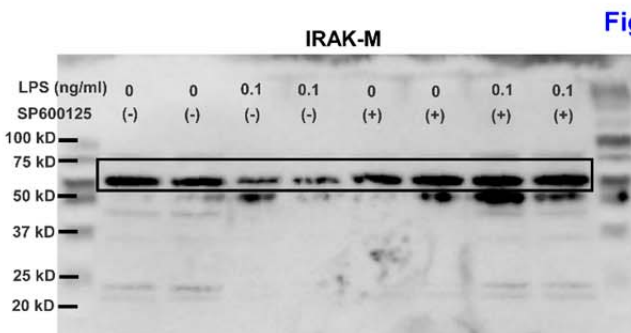
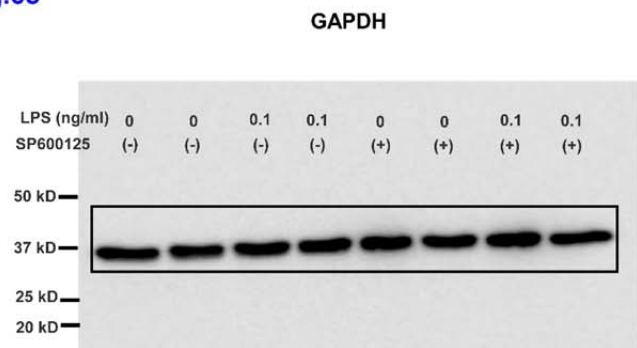
**Fig.5d**



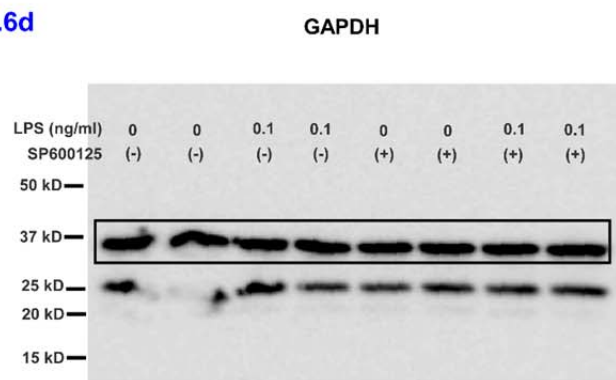
**Fig.5e**



**Fig.6c**

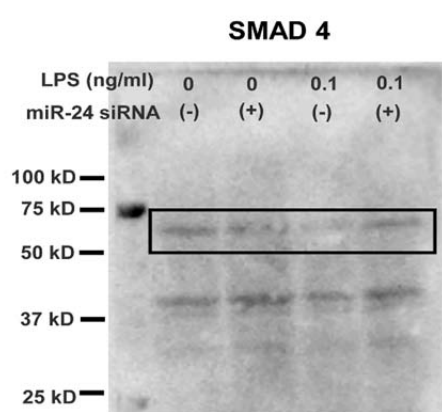


**Fig.6d**

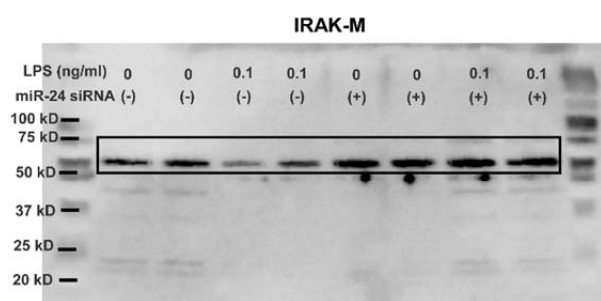
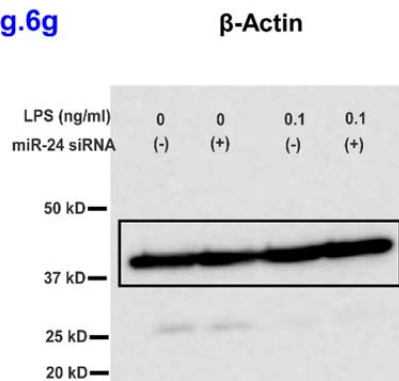


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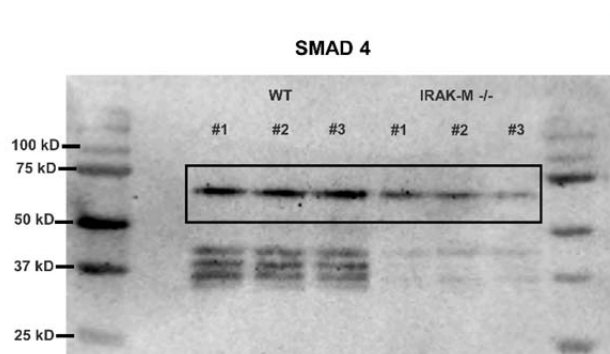
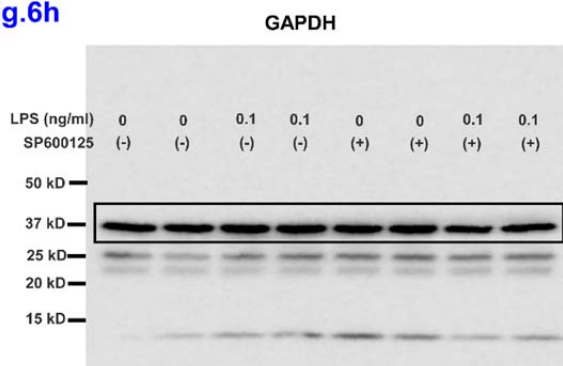
110 **Supplementary Fig 9. Original gels shown in the main manuscript.** Blots correspond to those  
111 shown in Figure 5d, 5e, and Figure 6c, and 6d within the main manuscript.



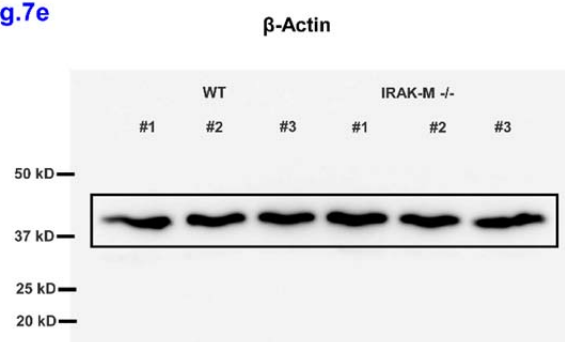
**Fig.6g**



**Fig.6h**



**Fig.7e**



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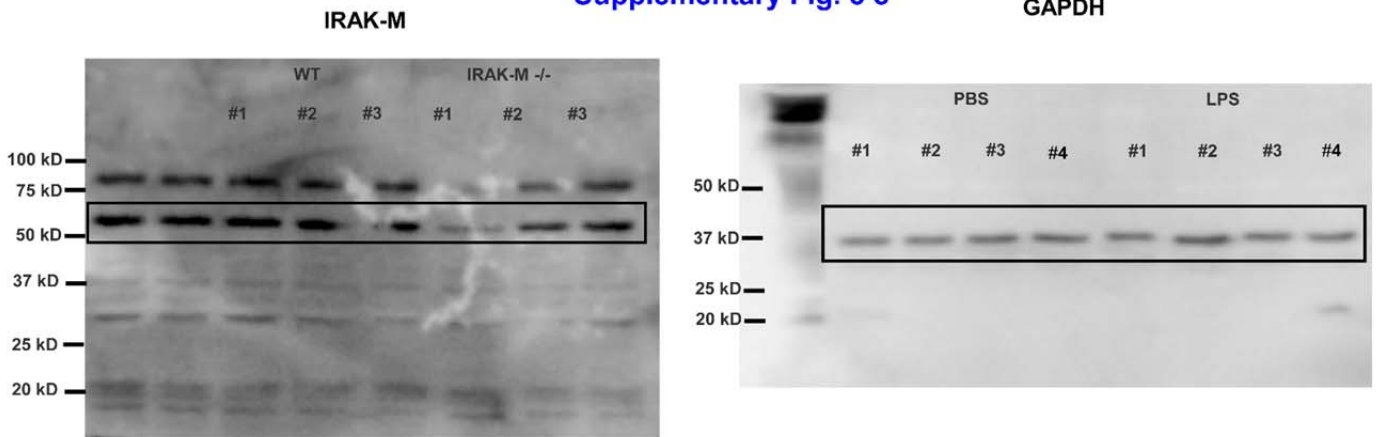
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114 **Supplementary Fig 10. Uncropped original gels shown in the main manuscript.** Blots  
115 correspond to those shown in Figure 6g, 6h, and Figure 7f within the main manuscript.

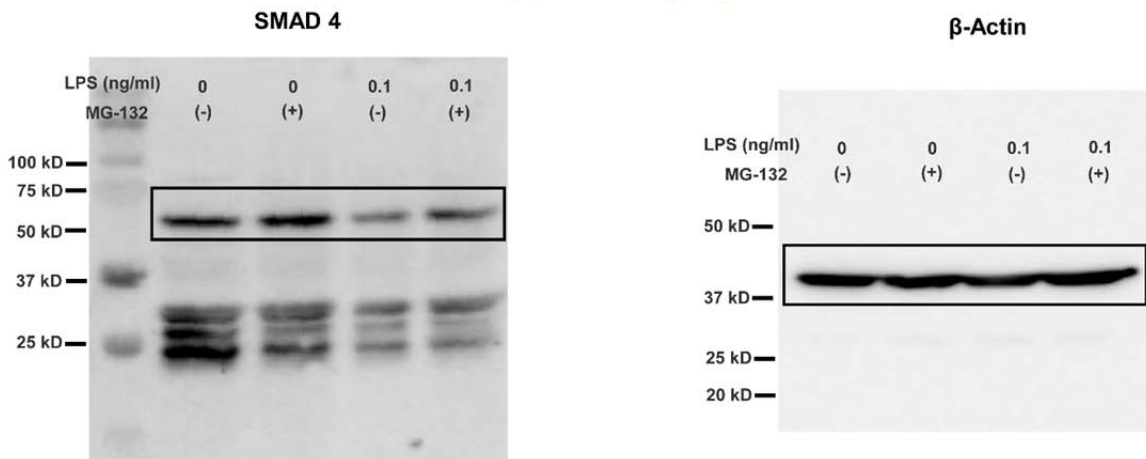
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Supplementary Fig. 5 c



Supplementary Fig. 6



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120 **Supplementary Fig 11. Uncropped original gels shown in the Supplementary Fig 5c and**  
121 **Supplementary Fig 6**

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**Supplementary Table 1. miRs selectively induced by subclinical dose LPS**

id	PBS	LPS	Fold Change	log2Fold Change
<b><i>mmu-miR-24-3p</i></b>	5163.2669	57785.0508	11.19156766	3.4843402
mmu-miR-3068-5p	19.061420	183.523741	9.628020362	3.2672391
mmu-miR-146a-5p	6615.9765	55275.4873	8.35484932	3.0626138
mmu-miR-1965	0.7773613	6.05221449	7.78558712	2.9608058
mmu-miR-1953	0.9549579	5.23425630	5.481137857	2.4544754
mmu-miR-450b-3p	3.4718013	17.9982757	5.184131734	2.3741023
mmu-miR-449a-5p	2.1102558	9.17771225	4.349099244	2.1207166
mmu-miR-152-3p	359.01571	964.823028	2.687411723	1.4262173
mmu-miR-6236	11.600608	30.2133288	2.604460776	1.3809847
mmu-miR-29b-2-5p	16.226452	41.4310964	2.553305899	1.3523663
mmu-miR-200a-3p	22.1733757	55.1966714	2.489321973	1.31575284
mmu-miR-6240	218.527628	471.464116	2.157457709	1.10933228
mmu-miR-2137	37.9225004	75.8060019	1.998971613	0.99925798
mmu-miR-155-5p	186.703613	337.736411	1.808944158	0.85514787
mmu-miR-188-5p	329.895138	594.127773	1.800959471	0.84876571
<b><i>mmu-miR-29a-5p</i></b>	112.545596	201.605426	1.79132221	0.84102486
mmu-miR-5115	251.576054	445.134329	1.769382744	0.82324615
mmu-miR-194-5p	132.146298	230.562523	1.744752032	0.80302201

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**Supplementary Table 2. Arbitrary Model Parameter Values**

<b>Parameter</b>	<b>Description</b>	<b>Value</b>
$\gamma_X$	Rate of <i>X</i> reaching its steady state	1
$\gamma_Y$	Rate of <i>Y</i> reaching its steady state	1
$\gamma_{IRAKM}$	Rate of <i>IRAK-M</i> reaching its steady state	1
$\omega_X$	Basal inhibition of <i>X</i>	-1.4
$\omega_{X,X}$	<i>X</i> auto-activation	6.4
$\omega_{Y,X}$	<i>X</i> inhibition by <i>Y</i>	-2
$\omega_{LPS,X}$	<i>X</i> activation by LPS	1
$\omega_{IRAKM,X}$	<i>X</i> inhibition by <i>IRAK-M</i>	-1
$\omega_Y$	Basal activation of <i>Y</i>	0.2
$\omega_{Y,Y}$	<i>Y</i> auto-activation	6.4
$\omega_{X,Y}$	<i>Y</i> inhibition by <i>X</i>	-2
$\omega_{IRAKM}$	Basal activation of <i>IRAK-M</i>	0.2
$\omega_{X,IRAKM}$	<i>IRAKM</i> inhibition by <i>X</i>	-2
<i>LPS</i>	Level of LPS	0 ~ 1

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