



**Universiteit
Leiden**
The Netherlands

Profiling of proteins and targeting of myeloid mechanisms in atherosclerosis

Delfos, L.

Citation

Delfos, L. (2026, June 30). *Profiling of proteins and targeting of myeloid mechanisms in atherosclerosis*. Retrieved from <https://hdl.handle.net/1887/4307144>

Version: Publisher's Version

License: [Licence agreement concerning inclusion of doctoral thesis in the Institutional Repository of the University of Leiden](#)

Downloaded from: <https://hdl.handle.net/1887/4307144>

Note: To cite this publication please use the final published version (if applicable).

Chapter 4

NLRP3 Inflammasome inhibition by the novel bispecific antibody InflammAb attenuates atherosclerosis in apolipoprotein E-deficient mice

Lucie Delfos, MSc¹, Marie A.C. Depuydt, PhD¹, Melody Chemaly, PhD², Sophie Coyle, MSc², Frank H. Schaftenaar, PhD¹, Peter J. van Santbrink, BSc¹, Pier P. Lindenberg, MSc¹, Mireia N.A. Bernabé Kleijn, BSc¹, Ciara Costello, MSc², Christine A. Power, PhD³, Rebecca Coll, PhD⁴, Aaron Peace, MD PhD⁵, Meredith Gregory-Ksander, PhD⁶, Amanda C. Foks, PhD¹, Johan Kuiper, PhD¹, Victoria McGilligan, PhD^{2#}, Ilze Bot, PhD^{1#}

¹Division of BioTherapeutics, Leiden Academic Centre for Drug Research, Leiden University, Leiden, The Netherlands.

²Personalised Medicine Centre, School of Medicine, Ulster University, Derry-Londonderry, Northern Ireland.

³Christine Power Consulting, Thoiry, Auvergne-Rhône-Alpes, France

⁴School of Medicine, Dentistry and Biomedical Sciences, Wellcome Wolfson Institute for Experimental Medicine

⁵Department of Cardiology, Western Health and Social Care Trust, Derry BT47 6SB, UK

⁶Department of Ophthalmology, Schepens Eye Research Institute, Massachusetts Eye & Ear Infirmary and Harvard Medical School, Boston, MA 02114, USA.

shared senior author

Published in JACC: Basic to Translational Science 2025;10(6):826-840.

Running title: InflammAb attenuates preclinical atherosclerosis

Abstract

Background: Cardiovascular disease remains the most common cause of mortality worldwide, which is attributable to the underlying chronic inflammatory condition atherosclerosis. The NLRP3 inflammasome contributes to this inflammatory process in atherosclerosis by producing IL-1 β . Components of the intracellular NLRP3 inflammasome have been shown to be expressed by macrophages in the atherosclerotic plaque and are a potential target for intervention.

Objectives: Here, we aimed to determine the efficacy of the novel bispecific antibody InflammAb, designed to target the interleukin-1 receptor type 1 and the NLRP3 inflammasome, in inhibiting atherosclerosis.

Methods: We established the efficacy of InflammAb to inhibit IL-1 β production upon NLRP3 inflammasome activation in mouse and human macrophages and in Western-type diet fed *ApoE*^{-/-} mice. Subsequently, we treated *ApoE*^{-/-} mice with developing collar-induced atherosclerosis and *ApoE*^{-/-} mice with established atherosclerotic plaques with InflammAb.

Results: InflammAb inhibited IL-1 β secretion from bone marrow derived macrophages and circulating IL-1 β levels *in vivo*, upon NLRP3 inflammasome activation. Furthermore, InflammAb treatment significantly inhibited atherosclerotic plaque development, which was accompanied by a reduction in relative macrophage and necrotic core content. Established atherosclerotic lesion size in the aortic root was not affected by InflammAb treatment, however, InflammAb significantly reduced relative macrophage and necrotic core content in these plaques.

Conclusions: In conclusion, inhibition of the NLRP3 inflammasome by the bispecific antibody InflammAb shows promising efficacy in inhibiting atherosclerotic plaque development and destabilization in mice.

Keywords: Atherosclerosis, NLRP3 Inflammasome, Interleukin-1 receptor type 1, bispecific antibody, Interleukin-1 β

Condensed Abstract

The NLRP3 inflammasome contributes to the inflammatory process in atherosclerosis by producing IL-1 β . Components of the intracellular NLRP3 inflammasome have been shown to be expressed by macrophages in the atherosclerotic plaque and are a potential therapeutic target. We aimed to determine the efficacy of the novel bispecific antibody InflammAb, designed to target the interleukin-1 receptor type 1 and the NLRP3 inflammasome, in inhibiting atherosclerosis. InflammAb effectively inhibited IL-1 β secretion from bone marrow derived

macrophages and reduced circulating IL-1 β levels in vivo. Furthermore, InflammAb treatment significantly inhibited atherosclerotic plaque development, accompanied by a reduction in relative macrophage and necrotic core content. InflammAb treatment did not affect the size of established atherosclerotic lesions, however, InflammAb significantly reduced relative macrophage and necrotic core content in these plaques. To conclude, inhibition of the NLRP3 inflammasome by the bispecific antibody InflammAb shows promising efficacy in inhibiting atherosclerotic plaque development and destabilization in *ApoE*^{-/-} mice.

Abbreviations

IL = Interleukin

NLRP3 = NOD (nucleotide oligomerization domain)-, LRR (leucine-rich repeat)-, and PYD (pyrin domain)-containing protein 3)

ASC = Apoptosis speck-like protein

ApoE^{-/-} = apolipoprotein E-deficient

LDL = low density lipoprotein

PAMPs = pathogen-associated molecular patterns

DAMPs = danger-associated molecular patterns

PRRs = pattern recognition receptors

IL-1R1 = IL-1 receptor type 1

CC = Cholesterol crystals

oxLDL = oxidized LDL

BMDMs = bone marrow derived macrophages

WTD = Western-type diet

Alum = Aluminium hydroxide

LPS = Lipopolysaccharides

TNF α = Tumor Necrosis Factor alpha

1. Introduction

Acute cardiovascular syndromes (ACS), such as myocardial infarction and stroke, remain a major cause of death worldwide. The pathology giving rise to these syndromes is atherosclerosis, which is characterized by the accumulation of lipids and inflammatory cells in the large and medium-sized arteries.^{1,2} Several immune cell subsets of both the myeloid and lymphoid lineage have been shown to contribute to the ongoing low-grade inflammation in atherosclerosis. Macrophages are a prominent immune cell type in both human³ and mouse atherosclerosis⁴ and are derived from monocytes that infiltrate the atherosclerotic lesion.⁵ A proportion of intraplaque macrophages may also derive from vascular proliferating macrophages.^{5,6} In addition, smooth muscle cells have been shown to transdifferentiate and present macrophage-like phenotype.⁷ In the atherosclerotic plaque, macrophages can produce pro- and anti-inflammatory cytokines, form foam cells by engulfing modified low density lipoprotein (LDL), and clear apoptotic cells via efferocytosis.^{8,9}

In recent years, a number of macrophage subsets have been identified, each with specific characteristics and inflammatory functions.¹⁰ Nowadays, single-cell RNA-sequencing technology has identified three major macrophage populations present in both human and mouse atherosclerotic vessels, namely the resident, inflammatory and foamy macrophages. The inflammatory macrophages largely express proinflammatory genes and are suggested to be derived from monocytes in the blood.^{4,11} Within the mouse inflammatory macrophages two subpopulations have been identified and classified as the inflammatory-*Nlrp3* and the $CCR2^{int}MHCII^{+}$ macrophages. The inflammatory-*Nlrp3* macrophages highly expressed *Nlrp3* and *Il1b* (Interleukin-1 β).¹¹

The NLRP3 inflammasome is a multiprotein complex located inside the cell and is composed of a sensor called NLRP3 (*NOD (nucleotide oligomerization domain)-, LRR (leucine-rich repeat)-, and PYD (pyrin domain)-containing protein 3*)), an adaptor the Apoptosis speck-like protein (ASC) and caspase-1 the effector. This inflammasome can be activated by a wide range of stimuli.^{12,13} Moreover, the NLRP3 inflammasome has been described to be an important driver of atherosclerotic inflammation.¹³ In human carotid artery plaques, NLRP3 mRNA was seen to be expressed at a higher level as compared to control obtained from transplant material and expression was higher in plaques of symptomatic versus asymptomatic patients.¹⁴ NLRP3 inflammasome components are predominantly expressed in macrophages and foam cells.^{14,15} Upon activation, the NLRP3 inflammasome contributes to the ongoing inflammatory response via activation of the inflammatory cytokines IL-1 β and IL-18. Activation of the NLRP3 inflammasome requires a priming signal, which can occur through binding of pathogen-associated molecular patterns (PAMPs) or danger-associated molecular patterns (DAMPs) signaling to pattern recognition receptors (PRRs). An alternative priming pathway is via IL-1 β itself through the IL-1 receptor type 1 (IL-1R1).^{12,16-18} Priming results in the transcriptional upregulation of the NLRP3 inflammasome components and the pro-forms of IL-1 β and IL-18.^{12,16} Cholesterol crystals (CC) and oxidized LDL (oxLDL), present in both initial and

advanced atherosclerosis, act as second-activation signals^{19,20}, leading to the assembly of the NLRP3 inflammasome. Then, caspase-1, also known as Interleukin-1 Converting Enzyme or ICE, is activated and cleaves pro-IL-1 β and pro-IL-18 into their active forms, IL-1 β and IL-18. Next, gasdermin D is cleaved and releases IL-1 β and IL-18 by pore formation.^{12,16}

In the previous Canakinumab Antiinflammatory Thrombosis Outcome Study (CANTOS), direct systemic targeting of IL-1 β with the monoclonal antibody canakinumab significantly lowered the rate of recurrence of cardiovascular events in patients with previous myocardial infarctions. However, this coincided with a higher occurrence of fatal infections and sepsis.²¹ Intervention in NLRP3 inflammasome activation using the small molecule MCC950 reduced atherosclerotic plaque development *in vivo*, which resulted from reduced macrophage content in the plaque.²² However, in Phase II clinical studies for rheumatoid arthritis with MCC950, a higher risk of drug-induced liver injury was found, which may be due to a high daily dose and high lipophilicity, both associated with hepatotoxicity risk.²³⁻²⁵ To circumvent the side-effects described in the studies above, alternative strategies to inhibit the NLRP3 inflammasome specifically in cells that contribute to the disease process are required. Therefore, we developed a novel bispecific antibody, InflammAb, designed to target both the IL-1R1 and the NLRP3 inflammasome. Via the IL-1R1, the antibody becomes internalized presumably by receptor-mediated endocytosis. Internalization of the bispecific antibody allows the antibody to reach the intracellular NLRP3 inflammasome target where it can exert its inhibitory activity. The construction of the bispecific antibody InflammAb is described elsewhere²⁶. In this study, we aimed to determine the ability of InflammAb to inhibit atherosclerosis. First, we established the efficacy of InflammAb *in vitro* and *in vivo*, after which we assessed the therapeutic anti-atherosclerotic potential of InflammAb during the development of atherosclerosis and in more advanced plaques in *ApoE*^{-/-} mice.

2. Materials and Methods

Generation of a bispecific antibody InflammAb, targeting the IL-1R1 and the NLRP3 inflammasome.

The construction of InflammAb is fully described in patent application WO2020053447A1²⁶. In brief, monoclonal antibodies targeting the IL-1R1 and the NLRP3 were generated at Fusion Antibodies (Northern Ireland, UK) using hybridoma technology after immunisation of the mice with the respective human target proteins. Following identification of functionally relevant antibodies from the hybridoma supernatants, the antibody heavy and light chain variable regions were cloned, sequenced and reformatted onto a mouse IgG2a framework. The bispecific antibody InflammAb, was generated by fusion of a scFV against NLRP3 to the C-terminal of the heavy chain of the mouse IgG2a mAb. The bispecific antibody was transiently expressed in Expi-CHO cells and purified using protein A affinity chromatography at Fusion Antibodies. Due to similar peptide sequences InflammAb also cross-reacts with the mouse proteins, which was confirmed in previous *in vitro* studies as part of the development process.

Human atherosclerotic plaque single-cell RNA sequencing (scRNA-seq) data

To determine which cell populations express both NLRP3 and IL1R1 in human atherosclerotic plaques, publicly available scRNA-seq datasets of the human atherosclerotic plaque (GSE155512, GSE159677, GSE131778, GSE253903)²⁷⁻³⁰ were downloaded, proximal adjacent samples were excluded from GSE159677 and the datasets were combined into a single object using Seurat packages (version 5.1.0)³¹ in R (version 4.4.1). Filtering was performed by removing doublets cells using scDblFinder (version 1.18.0)³², removing ambient RNA contamination by applying the decontX function of the celda R package (version 1.20.0; decontX: Decontamination of single cell genomics data. Version 1.2.0. Yin, Y, Yajima, M, Campbell, J; 2024. Accessed October 1, 2024. <https://www.bioconductor.org/packages/release/bioc/html/decontX.html>) and excluding cells with a mitochondrial gene percentage exceeding 10%, a total count of less than 800 or fewer than 500 unique genes detected. Layers were split based on the unique patient identifiers. The data were normalized, scaled and variable features were identified for each layer, using the SCTransform function of Seurat v5. Subsequently, integration was performed using the IntegrateLayers function of Seurat v5, where 'RPCAIntegration' was specified in 'method' argument and 'SCT' was specified in the 'normalization.method' argument. Double positive cells were identified using the WhichCells function of Seurat v5. The function's 'expression' argument was set to 'NLRP3 > 0.5 & IL1R1 > 0.5'.

Cell culture

Bone marrow was collected by flushing femurs and tibias from C57Bl/6J mice with ice-cold Phosphate buffered saline (PBS). Single-cell suspensions were obtained using a 70 µm cell strainer. Bone marrow cells were seeded at a density of 1×10^6 cells/mL and differentiated into bone marrow derived macrophages (BMDMs) by culturing for 7 days in RPMI with 10% FCS, 100 U/mL penicillin/streptomycin, 2 mM L-glutamine and 10 ng/mL m-CSF (Immunotools). To assess the efficacy of InflammAb *in vitro*, 0.1×10^6 BMDMs/well were plated in flat-bottom 96-well plates. After adhering overnight, cells were exposed to 50 ng/mL Lipopolysaccharides (LPS) (Salmonella minnesota R595, List Biological Laboratories Inc.) for 3 hours. After removal of the LPS, the cells were exposed to either 2.5 or 25 ng/mL InflammAb for 30 minutes. Subsequently, 50 µg/mL Aluminium hydroxide (Alum, Brenntag Biosector A/S) was added for 1 hour to activate the NLRP3 inflammasome, after which the supernatant was collected. IL-1 β and Tumor Necrosis Factor alpha (TNF α) concentrations were measured using mouse IL-1 β and TNF α ELISAs according to the manufacturer's protocol (BioLegend).

To assess the efficacy of InflammAb in a human cell line *in vitro*, THP-1 cells (ECACC) were seeded in a 96 well plate (Sarstedt, UK) at a density of 100,000 cells/well in complete RPMI containing 10% FBS, 1% penicillin/streptomycin and PMA (50 ng/mL). Cells were incubated for 48 hours in PMA to allow them to differentiate. After 48 hours, the medium was replaced with complete RPMI media without PMA. Cells were then left for another 24 hours before being primed with LPS (1 µg/mL) for 3 hours. InflammAb (50 µg/mL) or control antibodies were added for 24 hours. Cells were then activated with Nigericin (10 µM) (N7143, Sigma-Aldrich, UK) for 60 minutes before supernatants were removed and stored at -80°C until further analysis. THP-1 supernatants were diluted 1/25 prior to the IL-1 β ELISA (88-7261-77, Thermofisher, UK) or neat for a Caspase-Glo® 1 Inflammasome Assay (G9951 Promega, UK). To assess the binding of InflammAb to NLRP3, human recombinant NLRP3 protein (Cusabio, UK) was immobilised to a 96 well plate using coating buffer, before detection with HRP-tagged InflammAb.

Animal experiments

All experimental animal work was performed in compliance with the Dutch government guidelines, the Directive 2010/63/EU of the European Parliament and the ARRIVE guidelines. Study protocols were approved by the Ethics Committee for Animal Experiments and the Animal Welfare Body of Leiden University (project numbers 106002017887 and 10600202216361). C57BL/6J and atherosclerosis-prone apolipoprotein E-deficient (*ApoE*^{-/-}) mice were bred in the local animal facility and kept under standard laboratory conditions. Food and water were provided *ad libitum*.

To assess the presence of IL-1R1⁺NLRP3⁺ double positive cells in tissues of *ApoE*^{-/-} mice, 5 to 8 weeks old male *ApoE*^{-/-} mice were placed on a Western-type diet (WTD) containing 15% cocoa butter and 0.25% cholesterol for 12 weeks. At the end of these 12 weeks, the mice were anesthetized by subcutaneous administration of anesthetics (ketamine (100 mg/kg) and xylazine (10 mg/kg), after which sedation was monitored by toe pinch. Peritoneal cells were obtained by

flushing the peritoneal cavity (PC) with 10 mL ice-cold PBS. The mice were perfused with PBS through the left ventricle and organs, including vascular beds, were collected.

To study the efficacy of InflammAb *in vivo*, female *ApoE*^{-/-} mice were placed on a WTD containing 15% cocoa butter and 0.25% cholesterol for 2 weeks, during which the mice were treated with 100 µg InflammAb or isotype control antibody (*InVivo*MAb mouse IgG2a Isotype, BioXCell) in PBS, 3 times per week intraperitoneally (n=3-4 mice per group). At the end of these two weeks, an inflammatory challenge was performed by injecting low dose LPS (50 µg/kg i.v.), a dose which we have previously shown to be effective in inducing a potent circulating IL-1β response²². Plasma was collected and IL-1β/TNFα levels were measured by ELISA as described above.

To assess the effects of InflammAb on atherosclerosis development, 10 to 12 weeks old female *ApoE*^{-/-} mice, were placed on a WTD containing 15% cocoa butter and 0.25% cholesterol. The mice were randomly divided over the treatment groups based on body weight, age and plasma total cholesterol levels, and treatment groups were equally distributed over the different cages. From that moment onward, mice were treated with InflammAb (n=11) or isotype control antibody (n=14) 3 times per week intraperitoneally for a total of 6 weeks. At the 2-week timepoint, collars were placed as described previously.³³ In short, mice were anesthetized by subcutaneous injection of ketamine (60 mg/kg), fentanyl citrate, and fluanisone (1.26 mg/kg and 2 mg/kg, respectively), after which sedation was monitored by toe pinch. Access to the anterior cervical triangles was gained through a sagittal anterior neck incision and both carotid arteries were carefully dissected free from the surrounding tissue. Silastic collars (Dow Corning) were placed around both carotid arteries and fixed with 3 circumferential silk ties. Subsequently, the entry wound was closed, and the animals were returned to their cage for recovery from anesthesia. Blood was collected from the tail vein at week 0 and 2. At week 6, mice were anesthetized as described above, after which blood was collected via orbital bleeding. Subsequently, the mice were perfused with PBS and organs isolated.

To determine the effects of InflammAb on advanced atherosclerosis, 15 to 20 weeks old male *ApoE*^{-/-} mice were placed on a WTD for 8 weeks and allocated to an experimental group as described above, after which a baseline group was sacrificed (n=7). The remainder of the mice were kept on WTD for another 6 weeks and at the same time treated with 100 µg InflammAb (n=9) or isotype control (n=10) 3 times per week intraperitoneally. Blood was collected from the tail vein at weeks 8, 10 and 12. The mice were sacrificed at week 14 as described above.

Blood measurements

Collected blood samples were centrifuged for 10 min at 6000 g at 4 °C, after which plasma was stored at -80 °C. The total plasma cholesterol levels were measured with an enzymatic colorimetric assay using Precipath standardized serum (Roche) as the internal standard. Plasma glucose was measured using an Accu-Chek Instant test (Roche).

Histology

The carotid arteries and aortic roots were frozen in Tissue-Tek O.C.T. Compound (Sakura) and kept at -80°C until analysis. $10\ \mu\text{m}$ cryosections of the carotid arteries and aortic roots were prepared for histological analyses. For each carotid artery ($n=23$, one artery was excluded due to thrombosis, one lost due to technical issue), collection of the sections started immediately on the proximal side of the collar, and for each slide, the sections were collected every $90\ \mu\text{m}$ until the complete disappearance of the plaque. Mean plaque size, plaque size at the site of maximal stenosis, plaque volume, and necrotic areas were measured using hematoxylin and eosin (H&E) staining. The necrotic area was defined as the a-cellular, debris-rich plaque area, and was measured as absolute area and relative as percentage of total plaque area. A MOMA-2 antibody (1:1000, isotype IgG2b) and biotinylated Rabbit α -Rat (1:200 vector#BA-4001) as a secondary antibody was used to stain the macrophage content in the plaque.

Cryosections of the aortic root ($n=24$, two excluded due to technical issues), collected every $80\ \mu\text{m}$, were histologically stained with Oil-Red-O (ORO) and 3-4 sections covering the three-valve area were analyzed to measure the lipid content. The same staining as for the carotid arteries was used (MOMA-2) to measure the macrophage content in the atherosclerotic plaques. A Weigert's hematoxylin nuclei stain followed by a Sirius Red staining was performed to determine the collagen content and the necrotic core size. Histological analyses were performed with Leica QWin or the sections were imaged using a Panoramic 250 Flash III slide scanner (3DHISTECH, Hungary) and analyzed with ImageJ software. Mast cells were stained with Naphthol as-d chloroacetate esterase (Sigma-Aldrich). Resting and activated mast cells were counted directly under the microscope near the atherosclerotic plaque in the perivascular tissue of the aortic root. The FAM-FLICA Caspase-1 Assay #97 (ImmunoChemistry Technologies) was used to stain the active caspase-1 in the aortic root³⁴. Images were obtained with the slide scanner and analyzed manually by a blinded operator.

Flow cytometry

A red blood cell lysis was performed on whole blood with ACK lysis buffer to obtain a single white blood cell suspension. Aortic arches were cut into smaller pieces and incubated in a digestion mixture (Collagenase I (450U/mL), Collagenase XI(250U/mL), DNase (120U/mL), Hyaluronidase (120U/mL); all Sigma–Aldrich, in PBS) for 30 minutes at 37°C while shaking. The obtained mixture was filtered through a $70\ \mu\text{m}$ cell strainer to obtain a single cell suspension. To assess the level of NLRP3⁺IL-1R1⁺ double positive cells, the aortic arches of three individual mice were pooled. Aortic root and PC samples were assessed per individual mouse. Next, cell suspensions were stained for specific extracellular markers with flow cytometry antibodies (Table I) for 30 minutes. For the intracellular NLRP3 antibody staining, the corresponding samples were first incubated with fixation/permeabilization (Invitrogen) for 20 min and washed with permeabilization buffer (Invitrogen), after which the samples were stained with the antibody for 10 min. All samples were analyzed using flow cytometry on the

CytoFLEX (Beckman Coulter). The obtained data were analyzed with FlowJo 10.10.0 software. One peritoneal cell sample from the control group was excluded due to technical issues.

Table I. Flow cytometry antibodies.

Target antigen	Fluorochrome	Supplier	Catalog #	Working concentration
Fc block (aCD16/32)		Biolegend	101320	1:250
Fixable Viability dye	eFluor780	eBioscience	15383562	1:2000 or 1:1000
NLRP3⁺IL-1R1⁺ double positive cells				
<i>Extracellular</i>				
Ly-6C	Brilliant Violet 510	BioLegend	128033	1 :500
CD11b	Pacific Blue	BioLegend	101224	1 :500
I-A/I-E	Brilliant Violet 650	BioLegend	107641	1 :800
F4/80	FITC	BioLegend	123108	1 :400
Ly-6G	PerCP/Cyanine5.5	BioLegend	127616	1 :400
CD121a (IL-1 R, Type I/p80)	PE	BioLegend	113505	1 :50
CD45	Alexa Fluor 700	BioLegend	103128	1 :1000
<i>Intracellular</i>				
NLRP3 (NALP3)	APC	Miltenyi Biotec	Order no. : 130-111-397	1 :50
REA Control Antibody (I)	APC	Miltenyi Biotec	Order no. : 130-120-709	1 :50
Initial atherosclerosis				
CD11b	PE	eBioscience	12-0112-83	1:1000 or 1:400
F4/80	BV421	Biolegend	123137	1:500 or 1:400

MHC-II	eVolve 655	Thermo Fisher Scientific	86-5321-42	1:800
CD11c	FITC	Biolegend	117306	1:800
Ly6G	PerCP	Biolegend	127654	1:500
Ly6C	PE-CF594	BD Biosciences	562728	1:800
CD19	BV605	Biolegend	115539	1:500
CD4	V500	BD Horizon	560782	1:1000
CD8	AF700	Biolegend	100730	1:500
CD45	Alexa Fluor 700	Thermo Fisher Scientific	56-0451-82	1:3200
Advanced atherosclerosis				
CD4	Brilliant Violet 510	Biolegend	100559	1:1000
CD8a	Alexa Fluor 700	Biolegend	100730	1:500
CD19	Brilliant Violet 605	Biolegend	115540	1:500
CD3e	FITC	eBioscience	11-0031-85	1:500
Ly6G	PerCP	Biolegend	127654	1:500
Ly6C	PE-Dazzle 594	Biolegend	128044	1:800
NK1.1, CD161	Brilliant Violet 650	BD Horizon	564143	1:500
CD45	eFluor 450	eBioscience	48-0451-82	1:500
Cd11b	PE	Biolegend	101208	1:1000
CD11b	Brilliant Violet 605	Biolegend	101257	1:500
F4/80	FITC	Biolegend	123107	1:400
CD45	PE	Biolegend	103106	1:500
Ly6C	Brilliant Violet 510	Biolegend	128033	1:500
Ly6G	PerCP-Cy5.5	Biolegend	127616	1:400
I-A/I-E	Brilliant Violet 650	Biolegend	107641	1:800
CD11c	APC	eBioscience	17-0114-82	1:500

RNA isolation, cDNA synthesis and qPCR

Left carotid arteries of the initial atherosclerosis experiment were pooled (2-4 per sample) and homogenized in guanidine thiocyanate (GTC) with a tissue homogenizer, after which total RNA was extracted.³⁵ Reverse transcription of RNA was performed by M-MuLV reverse

transcriptase (RevertAid, MBI Fermentas) and qPCR was performed using a 7500 fast real-time PCR system (Applied Biosystems). Primer sequences are displayed in Table II.

Table II. Primer sequences used for the quantitative real-time PCR analysis, including three housekeeping genes (*36B4*, *Rpl27* and *Rpl37*).

Gene	Forward primer (3'-5')	Reverse primer (3'-5')
<i>36B4</i>	ctgagtacaccttcccactactga	cgactcttctttgcttcagcttt
<i>Rpl27</i>	cgccaagcgatccaagatcaagtcc	agctgggtccctgaacacatccttg
<i>Rpl37</i>	agagacgaaacactaccgggactgg	cttgggttccggcgttgttcctc
<i>Nlrp3</i>	cttctgcaccggactgtaaact	gaaggctgtggttgggtca
<i>IL-1R1</i>	agggactcctgctctggtttctcc	tcctccaagacctcaggcaacag
<i>Caspase-1</i>	tacctggcaggaattctggagcttc	gtcagtcctggaaatgtgccatcttc
<i>CD68</i>	ttgacctgctctctaaaggctacag	aggaccaggccaatgatgagagg
<i>CD86</i>	gtagagcgggatagtaacgctga	tgcacttctatttcaggcaaaagca
<i>CD206</i>	tctagcttcatcttcgggccttgg	tgaggatccatcttccttggctcagc
<i>Arg1</i>	tggcagaggtccagaagaatgg	gtgagcatccaccaaatgacac
<i>Tlr4</i>	ctgatcatggcactgttcttctcctg	ggaatgtcatcagggactttgctgag
<i>Mmp9</i>	tgtatagctacctcagggccttccc	ggacacatagtggagggtgctgtc
<i>Tlr9</i>	cctatactgcaccatctctcggct	gcgctctgtgccttatcgaacacc
<i>CD163</i>	cagtgccctcgtcacctg	gatctccacacgtccagaacagtc
<i>Il-8</i>	ttgttggatcctgatgctccatgg	gaagcttcattgccggtggaaattc
<i>Ccl2</i>	ctgaagccagctctcttctc	gggtaagtagtagcagcaggtga
<i>Il-6</i>	agcctggaggaggaaaggct	accgggtaagaccttcacacgagc
<i>Tnf-α</i>	acgctcttctgctactgaactcgg	actccagctgctcctccactg
<i>Sting</i>	tggcctggcactactacattgggtac	cctgcaccactgagcatgttgtatg

Statistical analysis

Data were analyzed using Prism 9.0 (GraphPad Software, Inc. San Diego, CA, USA). Data are presented using individual data points and/or the mean \pm standard error of the mean (SEM) for each group. Outlier tests were performed using Grubbs' test, after which significant outliers were removed from the analysis. To check for normality a Shapiro-Wilk test was performed. Data were compared using unpaired Student's t-test (2 groups) or one-way analysis of variance (ANOVA, >2 groups) if normally distributed and Mann-Whitney U (2 groups) or Kruskal-Wallis test (>2 groups) otherwise. Probability values of $P < 0.05$ were considered statistically significant.

3. Results

Presence of NLRP3⁺IL-1R1⁺ cells in atherosclerotic plaques

First, we established that the cells responsive to InflammAb, IL-1R1⁺NLRP3⁺ double positive cells, are present in atherosclerotic aortic arch and aortic root tissue, as well as in the PC of *ApoE*^{-/-} mice upon hyperlipidemia (Supplemental Figure 1A, B). The majority of the IL-1R1⁺NLRP3⁺ double positive cells in the atherosclerotic tissue are positive for the myeloid marker CD11b (Supplemental Figure 1C). Of these myeloid CD11b⁺ cells, especially in the aortic root, a proportion was positive for MHCII, suggesting that these cells are antigen presenting cells such as macrophages or dendritic cells (Supplemental Figure 1D). In addition, using existing scRNA-seq datasets we observed that also in human atherosclerotic plaques a proportion of the myeloid cell populations co-expressed IL-1R1 and NLRP3 (Supplemental Figure 1E).

InflammAb reduces IL-1 β levels in vitro and in vivo upon inflammasome activation

Next, we determined the efficacy of InflammAb *in vitro* and *in vivo*. The *in vitro* efficacy of InflammAb was evaluated in mouse bone marrow-derived macrophages. As shown in Figure 1A, both dosages of InflammAb significantly inhibited the LPS/Alum induced IL-1 β response, whereas TNF- α levels were not affected by InflammAb treatment (Supplemental Figure 2). *In vivo*, the efficacy of InflammAb was studied in a hyperlipidemic environment by treating Western-type diet fed *ApoE*^{-/-} mice with InflammAb and challenge the mice with LPS after 2 weeks. InflammAb significantly reduced circulating IL-1 β levels (Figure 1B; $P=0.022$ at $t=2h$, $P=0.010$ at $t=4h$), while the TNF- α levels were not affected by InflammAb treatment (Figure 1B), demonstrating the efficacy as well as the specificity of InflammAb *in vivo* to the IL-1 β pathway. Similarly, InflammAb significantly inhibited LPS and Nigericin induced IL-1 β secretion from human THP-1 macrophages (Figure 1C), while an IL-1R1 antibody alone, the NLRP3 ScFv fragment and a bispecific control, consisting of an inactive IL-1R1 antibody combined with the anti-NLRP3 ScFv, did not significantly reduce IL-1 β secretion. Similarly, InflammAb significantly inhibited LPS and Nigericin induced caspase-1 activity in THP-1 cells, while the bispecific control did not (Figure 1D). Binding of InflammAb to recombinant human NLRP3 protein was confirmed as well (Figure 1E).

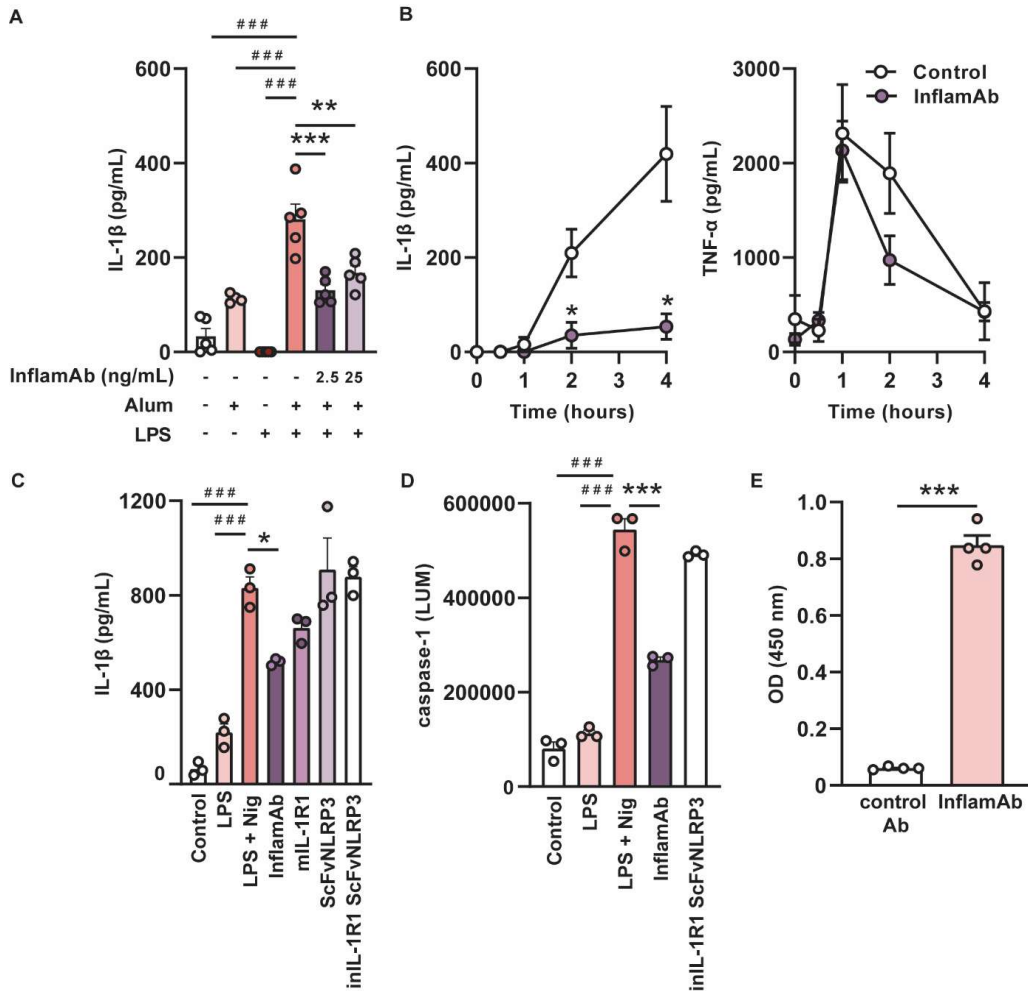


Figure 1. Inflammation Ab inhibits NLRP3 inflammasome-mediated IL-1 β release *in vitro* and *in vivo*. (A) The NLRP3 inflammasome (using LPS and Alum) induced IL-1 β secretion by BMDMs was significantly inhibited by both 2.5 and 25 ng/mL Inflammation Ab: n=4-5 replicates per condition, ordinary one-way ANOVA. (B) Inflammation Ab significantly inhibited the NLRP3 inflammasome induced IL-1 β response in WTD fed *ApoE*^{-/-} mice, compared to isotype control (left panel, n=4 both unpaired t test). (C) Inflammation Ab inhibited the NLRP3 inflammasome (using LPS and Nigericin; LPS + Nig) induced IL-1 β secretion by THP-1 cells. Single mIL-1R1 or ScFvNLRP3 antibodies, as well as a bispecific but inactive IL-1R1 ScFvNLRP3 (inIL-1R1 ScFvNLRP3) control did not significantly affect IL-1 β secretion. (D) LPS and Nigericin induced caspase-1 activity was significantly inhibited by Inflammation Ab, but not by the bispecific control antibody. (E) Inflammation Ab, but not a control antibody, significantly binds to recombinant human NLRP3 protein. Mean \pm SEM, *P<0.05, **P<0.01, ***P<0.001, ####P<0.001.

Inflammation Ab does not affect circulating leukocyte populations but reduces levels of peritoneal innate immune cells

After establishing that Inflammation Ab efficiently inhibits inflammasome activation both *in vitro* and *in vivo*, we next studied the ability of the bispecific antibody to inhibit atherosclerotic lesion

development by treating *ApoE*^{-/-} mice that were equipped with perivascular collars with InflamAb (Supplemental Figure 3A). During the study, InflamAb treatment did not affect total body weight (Supplemental Figure 3B) and total cholesterol levels (Supplemental Figure 3C). Also, total body score levels were not affected and we did not observe any signs of infection in the control or InflamAb treated mice. At the endpoint of the study, we measured circulating leukocyte populations, which were not affected (Figure 2 A-E). InflamAb treatment did reduce innate immune cell populations in the PC as illustrated by a non-significant lower percentage of macrophages (CD11b⁺F4/80⁺, control: 29±2% versus InflamAb: 22±2%; Figure 2F; P=0.05) and a significantly lower percentage of dendritic cells (CD11b⁺CD11c⁺MHC-II⁺, control: 1.4±0.1% versus InflamAb: 0.8±0.1%; Figure 2G; P=0.00002). Also, we observed a reduction in nonclassical myeloid cells (CD11b⁺Ly6C^{lo}, control: 0.25±0.03% versus InflamAb: 0.12±0.03%; Figure 2H; P=0.005) and Ly6C^{mid} myeloid cells (CD11b⁺Ly6C^{mid}, control: 0.7±0.1% versus InflamAb: 0.3±0.1%; Figure 2H; P=0.003).

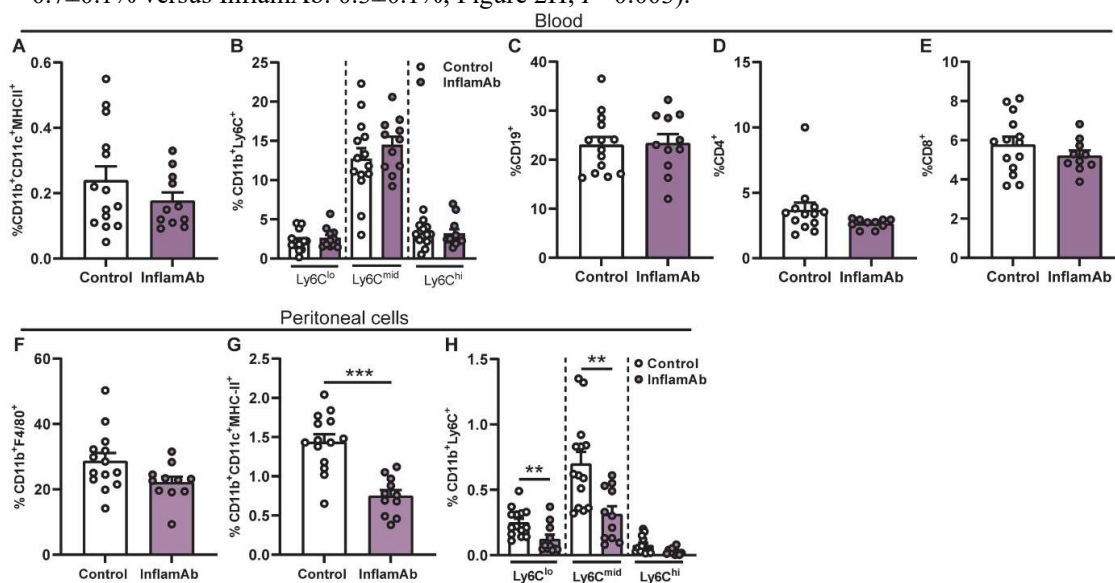


Figure 2. InflamAb treatment of WTD diet fed *ApoE*^{-/-} mice during atherosclerosis development did not affect leukocyte populations in the blood and reduced peritoneal innate immune cell populations. (A) The percentage of blood CD11b⁺CD11c⁺MHC-II⁺, (B) CD11b⁺Ly6C^{lo}, CD11b⁺Ly6C^{mid}, CD11b⁺Ly6C^{hi}, (C) CD19⁺, (D) CD4⁺, (E) CD8⁺ cells were not affected by InflamAb treatment. (F) The percentage of peritoneal CD11b⁺F4/80⁺ cells was reduced by InflamAb compared to isotype control (P=0.05). (G) CD11b⁺CD11c⁺MHC-II⁺, (H) CD11b⁺Ly6C^{lo} and CD11b⁺Ly6C^{mid} peritoneal cell populations were significantly lower upon InflamAb treatment, while the CD11b⁺Ly6C^{hi} content was not affected.

Panels A, B (Ly6C^{mid}), C, F, G, H (Ly6C^{mid}): control n=14, InflamAb n=11, unpaired t test. Panel B (Ly6C^{lo} and Ly6C^{hi}): control n=14, InflamAb n=11, Mann Whitney test. Panel D: control n=13, InflamAb n=10, Mann-Whitney test. Panel E: control n=14, InflamAb n=10, unpaired t test. Panel H (Ly6C^{lo}): control n=14, InflamAb n=11, Mann Whitney test. Panel H (Ly6C^{hi}): control n=13, InflamAb n=11, Mann-Whitney test.

All n-values represent individual animals. Mean ± SEM, all percentage of live cells, **P<0.01, ***P<0.001.

InflamAb inhibited collar-induced atherosclerotic plaque development by reducing macrophage levels and necrotic core area

Next, we assessed carotid artery lesion size and composition. InflamAb treatment significantly inhibited atherosclerotic plaque development (control: $59 \pm 8 \times 10^3 \mu\text{m}^2$ versus InflamAb: $37 \pm 5 \times 10^3 \mu\text{m}^2$; Figure 3A; $P=0.043$), while InflamAb did not affect the media size (Figure 3B). Both the relative necrotic core content (control: $14 \pm 2\%$ versus InflamAb: $8 \pm 1\%$; Figure 3C; $P=0.018$) and the absolute necrotic core area (control: $10 \pm 2 \times 10^3 \mu\text{m}^2$ versus InflamAb: $4 \pm 1 \times 10^3 \mu\text{m}^2$; Figure 3C; $P=0.030$) were reduced upon treatment with InflamAb compared to the control group. Also, the relative macrophage content was reduced by InflamAb treatment (control: $36 \pm 2\%$ versus InflamAb: $28 \pm 3\%$; Figure 3D; $P=0.048$). Also, a non-significant lower absolute macrophage area was observed (control: $24 \pm 4 \times 10^3 \mu\text{m}^2$ versus InflamAb: $14 \pm 3 \times 10^3 \mu\text{m}^2$; Figure 3D; $P=0.05$). Macrophages as percentage of the total immune cell population in the aortic arch, as measured with flow cytometry, did not significantly differ between the two groups (Figure 3E). Perivascular mast cell numbers (Supplemental Figure 4A) and their activation status (Supplemental Figure 4B) were not affected by InflamAb treatment.

Similarly as was shown previously for MMC950²², InflamAb did not affect the total mRNA expression levels of NLRP3 (Supplemental Figure 5A) or the IL-1R1 (Supplemental Figure 5B) in carotid plaques. In addition, we did not observe significant differences in genes related to macrophage phenotype or general inflammation, albeit that the expression of CD86, involved in antigen presentation and a marker of cellular activation, was almost 50% reduced (Supplemental Table I).

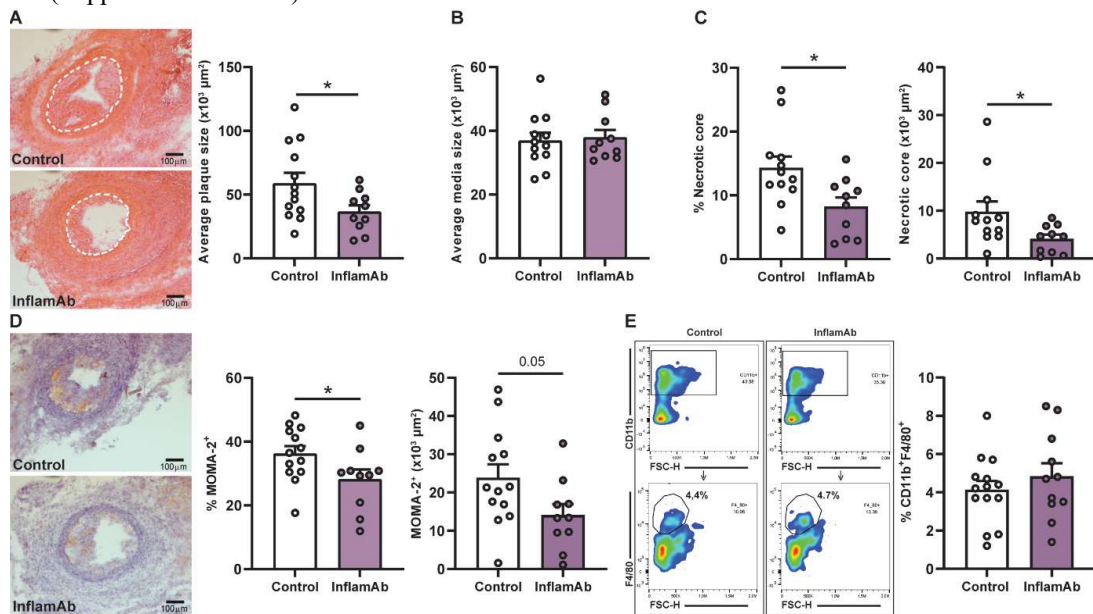


Figure 3. InflammAb treatment inhibited collar-induced atherosclerotic plaque development in WTD fed *ApoE*^{-/-} mice. (A) Representative images of H&E-stained carotid artery plaques (white dotted line = media outline, left panel) illustrate the reduction in average lesion size upon InflammAb treatment (right panel, n=13 versus n=10, unpaired t-test). (B) The average media size was not affected (n=12 versus n=10, unpaired t-test). (C) The relative necrotic core content (n=12 versus n=10, unpaired t-test), and the absolute necrotic core content (n=12 versus n=10, Mann-Whitney test) were also significantly smaller in the InflammAb treatment group compared to isotype control. (D) In the left panel representative images of the MOMA-2⁺ staining (red) are presented. The relative macrophage (MOMA-2⁺) content was shown to be significantly reduced by InflammAb. Also, a non-significant (P=0.05) lower absolute macrophage content was observed upon InflammAb treatment (both: n=13 versus n=10, unpaired t-test). (E) Representative flow cytometry plots of CD11b⁺ cells within the aortic CD45⁺ population (upper panels), followed by a F4/80⁺ gating within this CD11b⁺ population (lower panels). The CD11b⁺F4/80⁺ macrophage content in the aortic arch as a percentage of CD45⁺ live cells was not affected by InflammAb (n=14 versus n=11, unpaired t-test). All n-values represent individual animals. Mean ± SEM, *P<0.05, scale bars = 100 μm.

InflammAb increases markers of stability in established atherosclerotic plaques by reducing macrophage levels and necrotic core area

In a pre-existing atherosclerosis setup (Supplemental Figure 6A), body weight (Supplemental Figure 6B), total cholesterol levels (Supplemental Figure 6C), spleen weight (Supplemental Figure 6D) and plasma glucose levels (Supplemental Figure 6E) were not affected by InflammAb. Similarly, we did not observe signs of infection in this study. At the endpoint of the study, the circulating leukocyte populations did not differ between the groups (Figure 4 A-D). The peritoneal macrophage (CD11b⁺F4/80⁺, Figure 4E) and dendritic cell (CD11b⁺CD11c⁺MHC-II⁺, Figure 4F) percentages were not affected by InflammAb. However, as seen in the initial atherosclerosis study, the nonclassical myeloid cells (CD11b⁺Ly6C^{lo}, control: 0.28±0.06% versus InflammAb 0.12±0.01%; Figure 4G; P=0.013) and Ly6C^{mid} myeloid cells (CD11b⁺Ly6C^{mid}, control: 1.17±0.28% versus InflammAb: 0.45±0.05%; Figure 4G; P=0.008) were significantly reduced by InflammAb treatment compared to the controls.

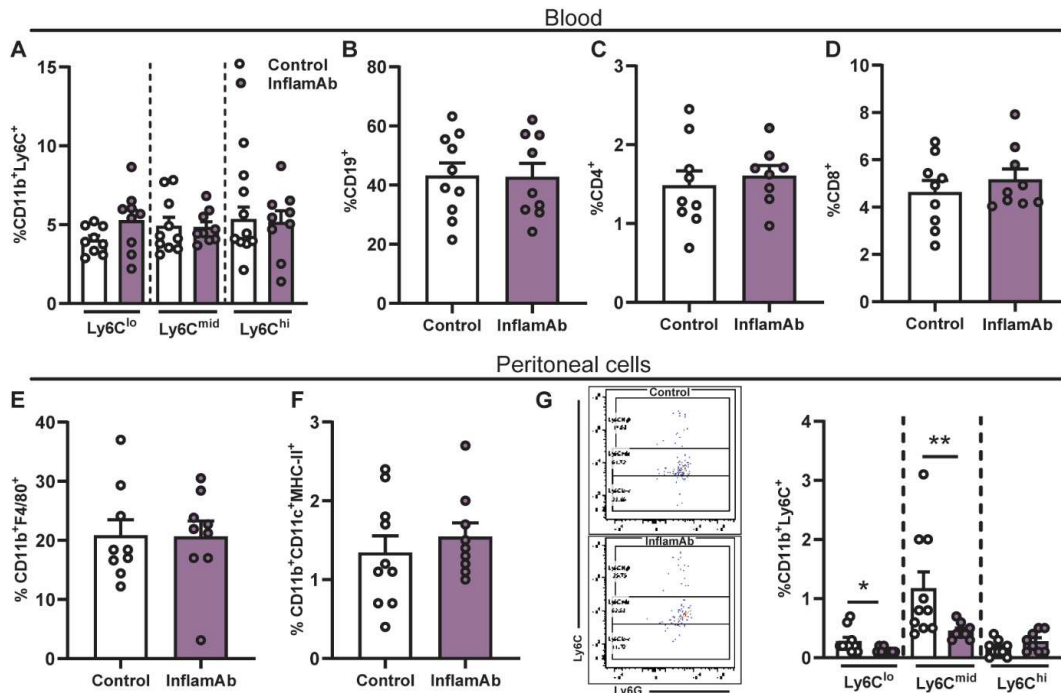


Figure 4. InflammAb treatment of advanced atherosclerotic lesions in *Apoe*^{-/-} mice did not affect leukocyte populations in the blood and reduced myeloid cell population in the PC. (A) The percentage of CD11b⁺Ly6C^{lo} and CD11b⁺Ly6C^{mid}, (B) CD19⁺, (C) CD4⁺ and (D) CD8⁺ cells were not affected by InflammAb treatment. (E) The percentage of peritoneal CD11b⁺F4/80⁺ and (F) CD11b⁺CD11c⁺MHC-II⁺ cells did not differ between the InflammAb and control groups. (G) Representative flow charts and quantification of the CD11b⁺Ly6C^{lo}, CD11b⁺Ly6C^{mid} and CD11b⁺Ly6C^{hi} peritoneal cells. CD11b⁺Ly6C^{lo} and CD11b⁺Ly6C^{mid} peritoneal cell percentages were significantly lowered by InflammAb, while the CD11b⁺Ly6C^{hi} content was not affected.

Panels A, E (Ly6C^{lo}): control n=9, InflammAb n=9, unpaired t test. Panels A (Ly6C^{mid} and Ly6C^{hi}), B, F: control n=10, InflammAb n=9, unpaired t test. Panels C, G (Ly6C^{hi}): control n=9, InflammAb n=8, unpaired t test. Panel D: control n=9, InflammAb n=9, Mann-Whitney test. Panel G (Ly6C^{lo}, Ly6C^{mid}): control n=10, InflammAb n=8-9, Mann-Whitney. All n-values represent individual animals. Mean ± SEM, *P<0.05, **P<0.01, all percentage of CD45⁺live cells.

To assess efficacy of InflammAb in this study, we measured caspase-1 activity using a Fluorescent Labeled Inhibitors of CASpases (FLICA) substrate, which showed significantly reduced caspase-1 activity in the InflammAb treated mice as compared to the controls (Supplemental Figure 7). InflammAb treatment of pre-existing lesions did not affect the absolute atherosclerotic plaque size and vessel occlusion parameters, measured by Oil-Red-O staining (Figure 5A). However, these plaques displayed increased plaque stability parameters upon treatment with InflammAb, as the relative necrotic core content was significantly reduced (control: 21±1% versus InflammAb: 18±1%; Figure 5B; P=0.019; Supplemental Figure 8 shows representative images of the necrotic core analysis). Collagen content, measured by Sirius Red staining, tended to be reduced upon InflammAb treatment (Figure 5B). In addition, the relative

macrophage content, stained with MOMA-2⁺, was significantly reduced (control: 48±2% versus InflamAb: 42±2%; Figure 5C; *P*=0.031). In the aortic arch the macrophage content measured with flow cytometry did not significantly differ between the two groups (Figure 5D). Similarly as in the plaque development study, adventitial mast cell numbers (Supplemental Figure 9A) and activation status (Supplemental Figure 9B) in the aortic root were not affected by InflamAb treatment.

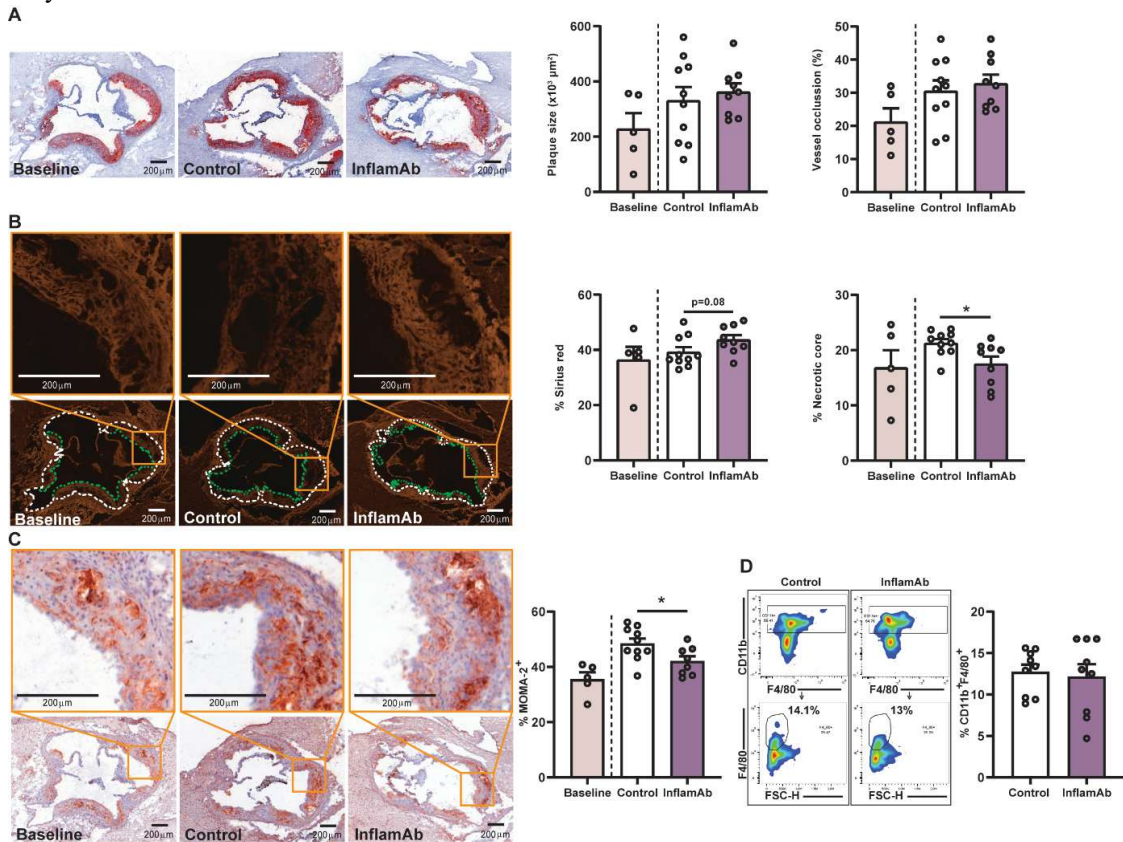


Figure 5. InflamAb treatment increased markers of plaque stability in advanced atherosclerotic lesions in WTD fed *ApoE*^{-/-} mice. (A) Representative images of the Oil-Red-O staining of the aortic root. Assessment of Oil-Red-O shows that InflamAb treatment did not affect absolute plaque size and the vessel occlusion (baseline n=5, control n=10 and InflamAb n=9, unpaired t test). (B) Representative images of the Sirius Red staining of the aortic root visualized in orange (white = vessel outline, green = the plaque outline inside the vessel, yellow = enlarged view of one of the valves). Sirius red assessment shows no effect of InflamAb on the collagen content (baseline n=5, control n=10 and InflamAb n=9, unpaired t test, control versus InflamAb: *P*=0.08). The relative necrotic core content was significantly reduced in the InflamAb treatment group compared to the control treatment (baseline n=5, control n=10 and InflamAb n=9, unpaired t test). (C) Representative images of the MOMA-2⁺ staining (bright red/brown) of the aortic root, including high power views. Assessment of MOMA-2⁺ shows that InflamAb treatment significantly lowered the MOMA-2⁺ percentage in the plaque (baseline n=5, control n=10 and InflamAb n=8, unpaired t test). (D) Representative flow cytometry plots of CD11b⁺ cells within the aortic CD45⁺ live population (upper panels), followed by a F4/80⁺ gating within this CD11b⁺ population (lower panels). This aortic arch macrophage content, as illustrated by CD11b⁺F4/80⁺ cells as a percentage of CD45⁺ live cells, was not affected by InflamAb (n=9 versus n=9, unpaired t test). All n-values represent individual animals. Mean ± SEM, **P*<0.05, scale bars = 200 μm.

4. Discussion

In this study, we investigated the atheroprotective efficacy of the new bispecific antibody InflammAb. The binding of the antibody to IL-1R1 acts as a ferry to deliver the bispecific antibody into the cell, while the therapeutic part (scFV) targets the intracellular NLRP3 inflammasome (Figure 6). The design of InflammAb is thus expected to result in a more specific approach than direct inhibition of IL-1 β or targeting all NLRP3 inflammasomes with a small molecule, because InflammAb inhibits only the NLRP3 inflammasome in cells expressing the IL-1R1.

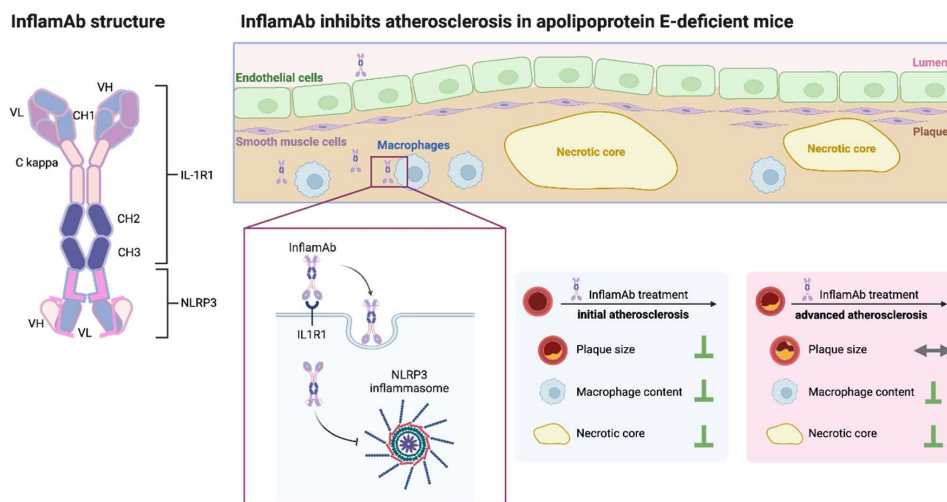


Figure 6. Visual summary of the InflammAb studies. Visual summary of the InflammAb structure and study outcomes. Created in BioRender. Chemaly, M. (2025) <https://BioRender.com/x25m415>.

InflammAb inhibits IL-1 β secretion in vitro and in vivo

We first confirmed that NLRP3⁺IL-1R1⁺ double positive cells are present in plaques of hyperlipidemic *ApoE*^{-/-} mice, confirming the suitability of the atherosclerotic mouse model, and that a subset of human plaque myeloid cells express both NLRP3 and IL-1R1. In these cells expressing both the IL-1R1 and the NLRP3 inflammasome, IL-1 β is able to induce its own synthesis and subsequently create an amplification loop of IL-1 β . Moreover, we clearly establish that InflammAb potently inhibits NLRP3 inflammasome induced IL-1 β production both *in vitro* and upon hyperlipidemia *in vivo*. In addition, we were able to show that InflammAb inhibits caspase-1 activity in a subset of cells in the advanced atherosclerotic plaques, showing efficacy of InflammAb at the target site.

InflammAb inhibits atherosclerotic plaque development

In the plaque development study, InflammAb reduced lesion size, which was caused by a reduction in both necrotic core and macrophage content. The latter finding is fully in line with our previous study, where we inhibited the NLRP3 inflammasome with the small molecule MCC950 in this mouse model.²² The reduction in intraplaque macrophages coincides with a reduction in inflammatory myeloid cells in the PC of these mice. Circulating leukocyte populations, including monocytes, were not affected by InflammAb treatment. This is in contrast with a study performed by Hettwer et al., who showed that inhibition of the NLRP3 inflammasome with MCC950 or treatment with an IL-1 β neutralizing antibody lowered leukocyte numbers in both the circulation and atherosclerotic aortas.³⁶ These findings thus illustrate a more specific and local nature of our approach. InflammAb treatment did not affect the local mRNA expression of NLRP3 and IL-1R1 in the atherosclerotic plaque, which is in line with our previous study where MCC950 did not affect the mRNA expression of NLRP3 and NLRP3-related genes in carotid artery plaques. It remains to be investigated whether the observed reduction in plaque macrophages is directly caused by a reduction in local IL-1 β levels or via reduced adhesion and subsequent influx of immune cells. IL-1 β deficiency in *ApoE*^{-/-} mice for example resulted in decreased vascular cell adhesion molecule (VCAM)-1 and monocyte chemoattractant protein-1 (MCP-1) mRNA expression in the aorta as compared to mice with IL-1 β .³⁷ Also, in our MCC950 study, NLRP3 inflammasome inhibition led to reduced VCAM-1 and ICAM-1 mRNA expression in the carotid artery.²² Similarly, Hettwer et al. showed that treatment with MCC950 reduced endothelial expression of adhesion molecules and chemoattractants in atherosclerotic aortas of *ApoE*^{-/-} mice. Together, these findings may underlie the reduced accumulation of leukocytes, including macrophages, upon inflammasome inhibition in atherosclerosis.³⁶

InflammAb increased plaque stability markers in pre-existing lesions

InflammAb treatment of pre-existing lesions increased the plaque stability markers, illustrated by both a reduced macrophage and necrotic core content as well as by a trend towards increased collagen content. Similarly as in the initiation study, peritoneal myeloid populations were reduced upon InflammAb treatment. During atherosclerotic lesion progression, macrophages differentiate into foam cells following uptake of oxidized or aggregated LDL. These foam cells undergo apoptosis or necrosis, thereby contributing to necrotic core formation, which enhances the probability for the lesion to rupture.² Furthermore, macrophages are known to contribute to the destabilization of plaques via the production of proteases that degrade collagen.² By limiting the macrophage accumulation, InflammAb treatment may limit necrotic core formation, while also reducing degradation of extracellular matrix molecules such as collagen. Additionally, it has been shown that IL-1 β itself increases the production of matrix metalloproteinase (MMP) 1, 8 and 13 in monocytes and macrophages.³⁸ Thus, a reduction in local IL-1 β levels in the plaque can affect the collagen content via a reduced MMP production. In line with these and our findings, increased plaque stability was also observed in the study from Zheng et al., in which the NLRP3 gene was silenced using a lentiviral vector in *ApoE*^{-/-} mice. NLRP3 silencing

reduced the progression of the plaques, while the plaques were less macrophage-rich and contained more collagen and smooth muscle cells.³⁹

Study limitations

Here, we studied the efficacy of a novel bispecific antibody InflammAb in inhibiting plaque development and progression in a preclinical mouse model of atherosclerosis. The cellular trafficking of InflammAb via the IL-1R1 leading to intracellular NLRP3 inflammasome inhibition however remains to be visualized *in vivo*. In addition, detailed studies on cellular migration have not been performed up to date. Such studies would provide more mechanistic insights in the local, but also potential systemic mechanisms involved, which now remain to be elucidated. For example, it is unknown whether InflammAb prevents active recruitment of myeloid cells to the plaque or whether InflammAb may have affected other inflammatory pathways as well which we were unable to detect in our studies. Furthermore, although we observed promising *in vitro* effects of InflammAb using human cells, studies using a more humanized disease model need to be performed to further establish the translational value of our findings.

Conclusion

The novel bispecific anti-NLRP3 antibody InflammAb inhibits plaque development and led to an increase in markers of plaque stability in more advanced plaques (Figure 6). The current encouraging data warrant further development of this therapeutic strategy against atherosclerosis and ACS.

Highlights

- We developed a novel bispecific antibody, InflammAb, designed to target the IL-1R1 for cell entrance and inhibit the intracellular NLRP3 inflammasome.
- InflammAb potently inhibits NLRP3 inflammasome induced IL-1 β production *in vitro* in BMDMs.
- InflammAb potently inhibits NLRP3 inflammasome induced IL-1 β production upon

hyperlipidemia *in vivo* in *ApoE*^{-/-} mice.

- InflamAb inhibits atherosclerotic plaque development in *ApoE*^{-/-} mice.
- InflamAb enhances stabilization parameters of advanced plaques in *ApoE*^{-/-} mice.

Clinical Competency in Medical Knowledge

Acute cardiovascular syndromes are a leading cause of death worldwide and generally caused by rupture or erosion of an atherosclerotic plaque. Recent studies have established that inhibition of the IL-1 β pathway is a powerful therapeutic approach to limit the incidence of acute cardiovascular events. Here, we provide a novel tool to intervene in NLRP3 inflammasome induced IL-1 β production by means of the bispecific antibody InflamAb. In this study, we conclusively demonstrate that treatment of *ApoE*^{-/-} mice with InflamAb inhibits atherosclerotic lesion development and improves the stability of more advanced plaques, rendering InflamAb a promising therapeutic lead for plaque stabilization.

Translational Outlook

From a translational perspective, the increased stability upon InflamAb treatment of preexisting atherosclerosis is a promising finding in relation to patients with established atherosclerosis. In the CANTOS trial, where patients with previous myocardial infarctions and thus established atherosclerotic lesions were treated with an anti-IL-1 β antibody, the potential of targeting this inflammatory pathway to limit secondary cardiovascular events was already demonstrated. Due to the systemic nature of this approach, serious side-effects related to sepsis and deaths due to infection occurred, rendering a more targeted therapy necessary.²¹ Due to its bispecific approach to NLRP3⁺IL-1R1⁺ double positive cells, InflamAb may overcome these adverse effects, at the same time limiting plaque instability parameters. Our preclinical experiments are the first steps towards a clinical application for InflamAb against acute cardiovascular syndromes.

Conflict of interest

None

Author contribution

Lucie Delfos: conceptualization, executing experiments, data acquisition, data analysis, and interpretation, writing – original draft. **Marie A.C. Depuydt:** data acquisition, data analysis, writing – review, and editing. **Melody Chemaly:** conceptualization, writing – review, and editing. **Sophie Coyle:** writing – review, and editing. **Frank H. Schaftenaar:** data analysis. **Peter J. van Santbrink:** executing experiments, data acquisition. **Pier P. Lindenbergh:** data acquisition, data analysis, writing – review, and editing. **Mireia N.A. Bernabé Kleijn:** executing experiments. **Ciara Costello:** writing – review, and editing. **Christine A. Power:** writing – review, and editing. **Rebecca Coll:** writing – review, and editing. **Aaron Peace:** writing – review, and editing. **Meredith Gregory-Ksander:** writing – review, and editing. **Amanda C. Foks:** writing – review, and editing. **Johan Kuiper:** writing – review, and editing. **Victoria McGilligan:** Invention and development of InflammAb, conceptualization, writing – review, and editing. **Ilze Bot:** conceptualization, executing experiments, data acquisition, data analysis, and interpretation, supervision, writing – original draft, funding acquisition.

References

1. Libby P. The changing landscape of atherosclerosis. Review. *Nature*. 2021;592:524-533.
2. Bjorkegren J, Lusis AJ. Atherosclerosis: Recent developments. Review. *Cell*. 2022;185:1630-1645.
3. Fernandez D, Rahmam AH, Fernandez NF, et al. Single-cell immune landscape of human atherosclerotic plaques. *Nature Medicine*. 2019;25:1576–1588.
4. Zernecke A, Winkels H, Cochain C, et al. Meta-Analysis of Leukocyte Diversity in Atherosclerotic Mouse Aortas. *Circulation Research*. 2020;127:402–426.
5. Robbins C, Hilgendorf I, Weber GF, et al. Local proliferation dominates lesional macrophage accumulation in atherosclerosis. *Nature Medicine*. 2013;19:1166-1172.
6. Ensan S, Li A, Besla R, et al. Self-renewing resident arterial macrophages arise from embryonic CX3CR1+ precursors and circulating monocytes immediately after birth. *Nature Immunology*. 2016;17:159-168.
7. Feil S, Fehrenbacher B, Lukowski R, et al. Transdifferentiation of Vascular Smooth Muscle Cells to Macrophage-Like Cells During Atherogenesis. *Circulation Research*. 2014;115:662-667.
8. Cochain C, Zernecke A. Macrophages in vascular inflammation and atherosclerosis. *Pflugers Arch - Eur J Physiol* 2017;469:485-499.
9. Cybulsky M, Cheong C, Robbins CS. Macrophages and Dendritic Cells Partners in Atherogenesis. *Circulation Research* 2016;118:637-652.
10. Wieland E, Kempen LJAP, Donners MMPC, Biessen EAL, Goossens P. Macrophage heterogeneity in atherosclerosis: A matter of context. *Eur J Immunol*. 2023:1-9.
11. Zernecke A, Erhard F, Weinberger T, et al. Integrated single-cell analysis-based classification of vascular mononuclear phagocytes in mouse and human atherosclerosis. *Cardiovascular Research*. 2023;119:1676-1689.

12. Swanson K, Deng M, Ting JP-Y. The NLRP3 inflammasome: molecular activation and regulation to therapeutics. *Nature reviews, Immunology*. 2019;19:477-489.
13. Grebe A, Hos, F, Latz E. NLRP3 Inflammasome and the IL-1 Pathway in Atherosclerosis. Review. *Circulation Research*. 2018;122:1722-1740.
14. Varghese G, Folkersen L, Strawbridge RJ, et al. NLRP3 Inflammasome Expression and Activation in Human Atherosclerosis. *Journal of the American Heart Association*. 2016;5:1-11.
15. Shi X, Xie W-L, Kong W-W, Chen D, Qu P. Expression of the NLRP3 Inflammasome in Carotid Atherosclerosis. *Journal of Stroke and Cerebrovascular Diseases*. 2015;24:2455-2466.
16. Silvis M, Demkes EJ, Fiolet ATL, et al. Immunomodulation of the NLRP3 Inflammasome in Atherosclerosis, Coronary Artery Disease, and Acute Myocardial Infarction. Review. *Journal of Cardiovascular Translational Research*. 2021;14:23-34.
17. Libby P. Interleukin-1 Beta as a Target for Atherosclerosis Therapy. Review. *J Am Coll Cardiol*. 2017;70:2278-2289.
18. Dinarello C, Ikejima T, Warner SJC, et al. Interleukin 1 induces interleukin 1. I. Induction of circulating interleukin 1 in rabbits in vivo and in human mononuclear cells in vitro. *J Immunol*. 1987;139:1902-1910.
19. Duewell P, Kono H, Rayner KJ, et al. NLRP3 inflammasomes are required for atherogenesis and activated by cholesterol crystals. *Nature*. 2010;464:1357-1361.
20. Sheedy F, Grebe A, Rayner KJ, et al. CD36 coordinates NLRP3 inflammasome activation by facilitating intracellular nucleation of soluble ligands into particulate ligands in sterile inflammation. *Nature Immunology*. 2013;14:812-820.
21. Ridker P, Everett BM, Thuren T, et al. Antiinflammatory Therapy with Canakinumab for Atherosclerotic Disease. *The New England Journal of Medicine*. 2017;377:1119-1131.

22. van der Heijden T, Kritikou E, Venema W, et al. NLRP3 Inflammasome Inhibition by MCC950 Reduces Atherosclerotic Lesion Development in Apolipoprotein E–Deficient Mice—Brief Report. *Arterioscler Thromb Vasc Biol.* 2017;37:1457-1461.
23. Duan M, Sun L, He X, Wang Z, Hou Y, Zhao Y. Medicinal chemistry strategies targeting NLRP3 inflammasome pathway: A recent update from 2019 to mid-2023. Review article. *European Journal of Medicinal Chemistry.* 260:115750.
24. Shah F, Leung L, Barton HA, et al. Setting Clinical Exposure Levels of Concern for Drug-Induced Liver Injury (DILI) Using Mechanistic in vitro Assays. *Toxicol Sci.* 2015;147:500-514.
25. Chen M, Borlak J, Tong W. High Lipophilicity and High Daily Dose of Oral Medications Are Associated With Significant Risk for Drug-Induced Liver Injury. *HEPATOLOGY.* 2013;58:388-396.
26. McGilligan V, inventor; BISPECIFIC ANTIBODY TARGETING IL-1R1 AND NLRP3. GB patent application PCT/EP2019/074745. 2020.
27. Wirka RC, Wagh D, Paik DT, et al. Atheroprotective roles of smooth muscle cell phenotypic modulation and the TCF21 disease gene as revealed by single-cell analysis. *Nat Med.* 2019;25:1280-1289.
28. Pan H, Xue C, Auerbach BJ, et al. Single-Cell Genomics Reveals a Novel Cell State During Smooth Muscle Cell Phenotypic Switching and Potential Therapeutic Targets for Atherosclerosis in Mouse and Human. *Circulation.* 2020;142:2060-2075.
29. Alsaigh T, Evans D, Frankel D, Torkamani A. Decoding the transcriptome of calcified atherosclerotic plaque at single-cell resolution. *Commun Biol.* 2022;5:1084.
30. Bashore AC, Yan H, Xue C, Zhu LY, Kim E, Mawson T, Coronel J, Chung A, Sachs N, Ho S, Ross LS, Kissner M, Passegué E, Bauer RC, Maegdefessel L, Li M, Reilly MP. High-Dimensional Single-Cell Multimodal Landscape of Human Carotid Atherosclerosis. *Arterioscler Thromb Vasc Biol.* 2024;44:930-945.

31. Hao Y, Stuart T, Kowalski MH, et al. Dictionary learning for integrative, multimodal and scalable single-cell analysis. *Nature Biotechnology*. 2024;42:293–304.
32. Germain P-L, Lun A, Meixide CG, Macnair W, Robinson MD. Doublet identification in single-cell sequencing data using scDblFinder. *F1000Research*. 2022;10:979.
33. von der Thüsen J, van Berkel TJC, Biessen EAL. Induction of Rapid Atherogenesis by Perivascular Carotid Collar Placement in Apolipoprotein E-Deficient and Low-Density Lipoprotein Receptor-Deficient Mice. *Circulation*. 2001;103:1164-1170.
34. Peikert A, König S, Suchanek D, et al. P2X4 deficiency reduces atherosclerosis and plaque inflammation in mice. *Scientific Reports*. 2022;12:2801.
35. Chomczynski P, Sacchi N. Single-Step Method of RNA Isolation by Acid Guanidinium Thiocyanate-Phenol-Chloroform Extraction. *ANALYTICAL BIOCHEMISTRY*. 1987;162:156-159.
36. Hettwer J, Hinterdobler J, Miritsch B, et al. Interleukin-1b suppression dampens inflammatory leucocyte production and uptake in atherosclerosis. *Cardiovascular Research*. 2022;118:2778-2791.
37. Kirii H, Niwa T, Yamada Y, et al. Lack of Interleukin-1 β Decreases the Severity of Atherosclerosis in ApoE-Deficient Mice. *Arterioscler Thromb Vasc Biol*. 2003;23:656-660.
38. Libby P. Interleukin-1 Beta as a Target for Atherosclerosis Therapy - Biological Basis of CANTOS and Beyond. *J Am Coll Cardiol*. 2017;70:2278-89.
39. Zheng F, Xing S, Gong Z, Mu W, Xing Q. Silence of NLRP3 Suppresses Atherosclerosis and Stabilizes Plaques in Apolipoprotein E-Deficient Mice. *Mediators of Inflammation*. 2014:1-8.

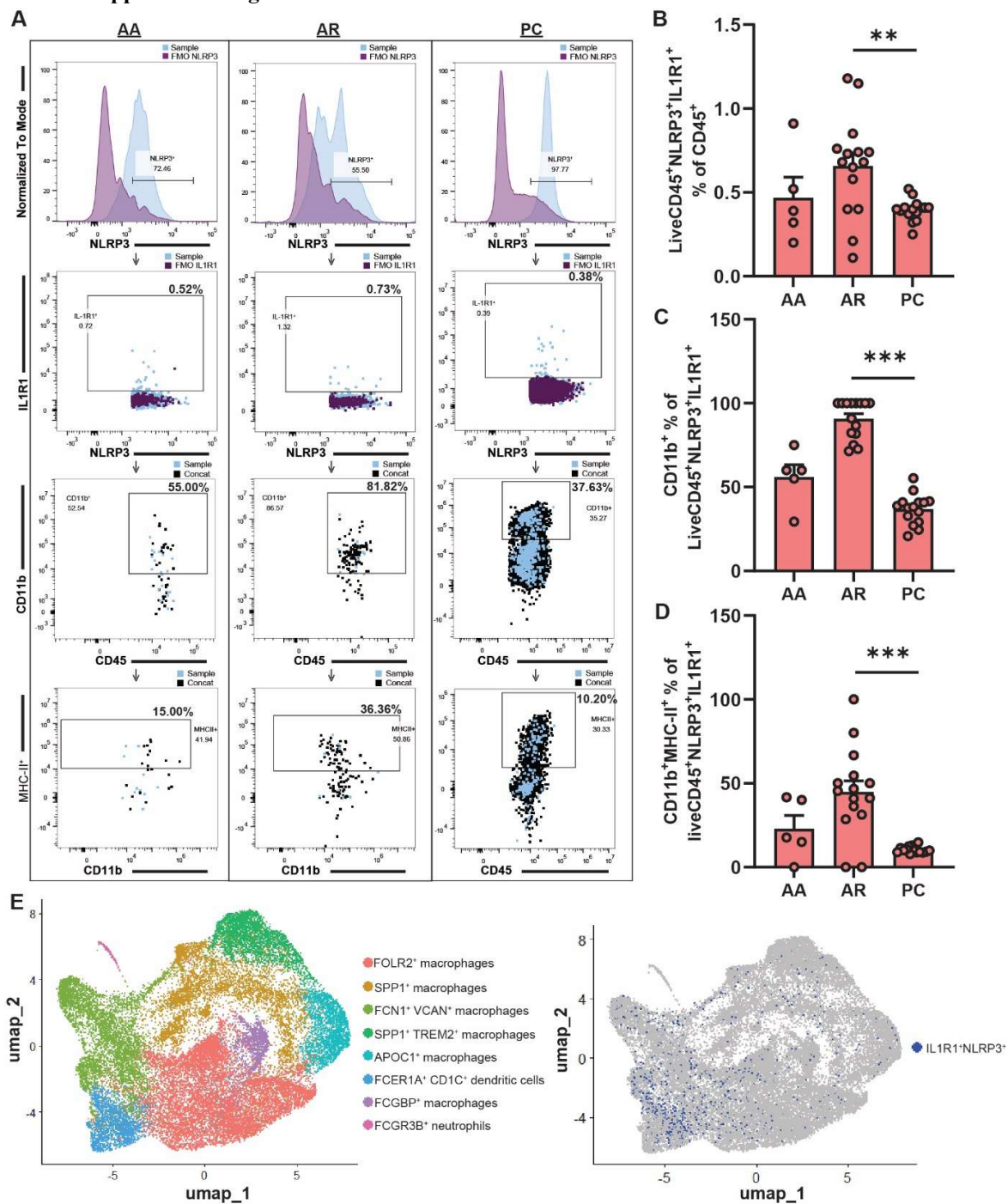
Supplementary Data

Supplementary Table

Supplementary Table I: Relative gene expression in control versus InflamAb treated carotid artery plaques (n=4, pooled samples). Gene expression is displayed relative to the average of three housekeeping genes (*36B4*, *Rpl27* and *Rpl37*). Data are given as mean \pm SEM.

Gene	Control	InflamAb	p-value
<i>Caspase-1</i>	0.0012 \pm 0.0005	0.0007 \pm 0.0002	0.12
<i>CD68</i>	0.083 \pm 0.013	0.094 \pm 0.051	0.91
<i>CD86</i>	0.0008 \pm 0.0003	0.0004 \pm 0.001	0.05
<i>CD206</i>	0.0021 \pm 0.0011	0.0016 \pm 0.0007	0.48
<i>Arg1</i>	0.0014 \pm 0.0015	0.0011 \pm 0.0007	0.83
<i>Tlr4</i>	0.0023 \pm 0.0013	0.0020 \pm 0.0006	0.96
<i>Mmp9</i>	0.0057 \pm 0.0031	0.0130 \pm 0.0110	0.33
<i>Tlr9</i>	0.0029 \pm 0.0009	0.0016 \pm 0.0010	0.14
<i>CD163</i>	0.0005 \pm 0.0004	0.0003 \pm 0.0001	0.35
<i>Il-8</i>	0.00005 \pm 0.00002	0.00010 \pm 0.00009	0.18
<i>Ccl2</i>	0.0068 \pm 0.0028	0.0057 \pm 0.0018	0.63
<i>Il-6</i>	0.00022 \pm 0.00032	0.00011 \pm 0.00005	0.93
<i>Tnf-α</i>	0.0040 \pm 0.0025	0.0047 \pm 0.0062	0.70
<i>Sting</i>	0.0073 \pm 0.0016	0.0089 \pm 0.0027	0.71

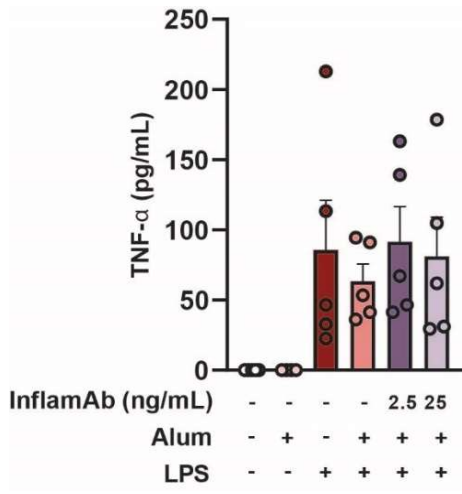
Supplemental Figures



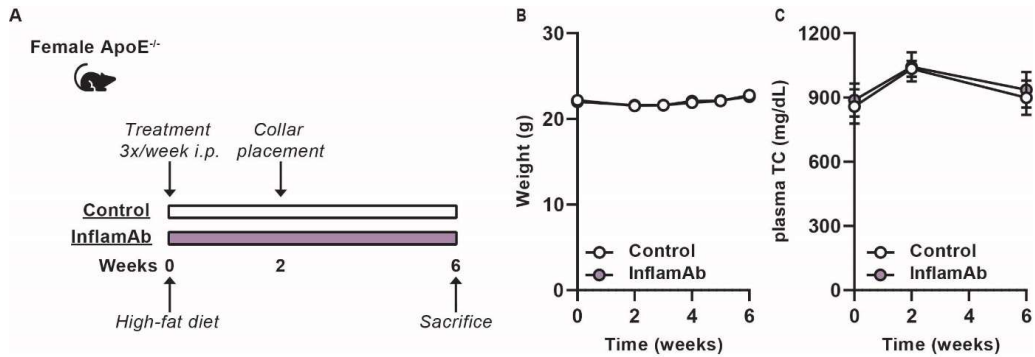
Supplemental Figure 1. NLRP3⁺IL-1R1⁺ cells are present in tissues of *ApoE*^{-/-} mice. (A)

Representative flow cytometry plots of the gating for the aortic arch (AA), aortic root (AR) and

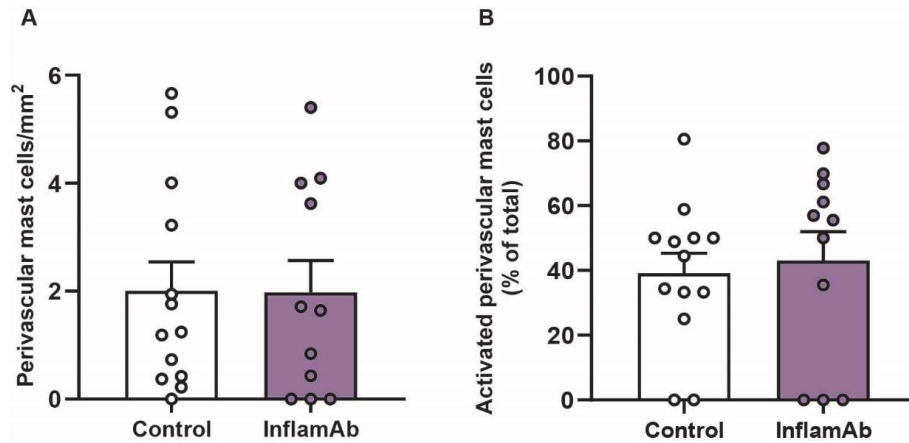
the peritoneal cavity (PC). The upper panels present the NLRP3⁺ gate within the live CD45⁺ cells in a representative sample and the concomitant Fluorescence Minus One (FMO). Normalized To Mode indicates a scaling of the channels as a percentage of the maximum count. Below, the IL-1R1⁺ gate within this NLRP3⁺ population is shown, in a representative sample and the IL-1R1 FMO. In the third lane, flow cytometry plots of the CD11b⁺ gate within the NLRP3⁺IL-1R1⁺ cell population are shown, followed by the MHC-II⁺ gate within the CD11b⁺ population (light blue dots: representative sample, black dots: concatenated sample). (B) Live CD45⁺NLRP3⁺IL-1R1⁺ cells as a percentage of CD45⁺. (C) CD11b⁺ cells as a percentage of the live CD45⁺NLRP3⁺IL-1R1⁺ cells and (D) CD11b⁺MHC-II⁺ cells as a percentage of the live CD45⁺NLRP3⁺IL-1R1⁺ cells in AA, AR and PC. A and B: AA n=5 with 3 pooled aortas per sample, AR n=15 and PC n=15. B and D: Ordinary one-way ANOVA. C: Kruskal-Wallis test. Mean ± SEM, **P<0.01, ***P<0.001. (E) Left panel: a tSNE plot of the clustering of the myeloid population in human atherosclerotic plaques. Right panel: IL-1R1⁺NLRP3⁺ double positive cells in blue.



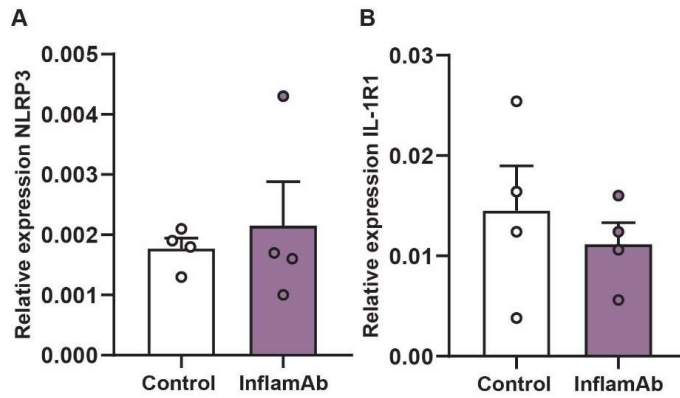
Supplemental Figure 2. InflamAb did not affect TNF- α responses during NLRP3 inflammasome activation using LPS and Alum in BMDMs: n=5 replicates per condition, ordinary one-way ANOVA. Mean \pm SEM.



Supplemental Figure 3. InflamAb treatment during plaque development in *ApoE*^{-/-} mice did not affect the body weight and plasma total cholesterol levels. (A) Schematic overview of the study set-up. Female *ApoE*^{-/-} mice were treated with InflamAb or isotype control antibody *InVivoMAb* 3 times per week intraperitoneally for 6 weeks while being fed a high-fat diet. In week 2 silastic collars were placed around both carotid arteries to induce atherosclerosis. (B) Body weight and (C) plasma total cholesterol levels were not affected by InflamAb during the study. B and C: n=14 control versus n=11 InflamAb treated animals, unpaired t-test. Mean ± SEM.

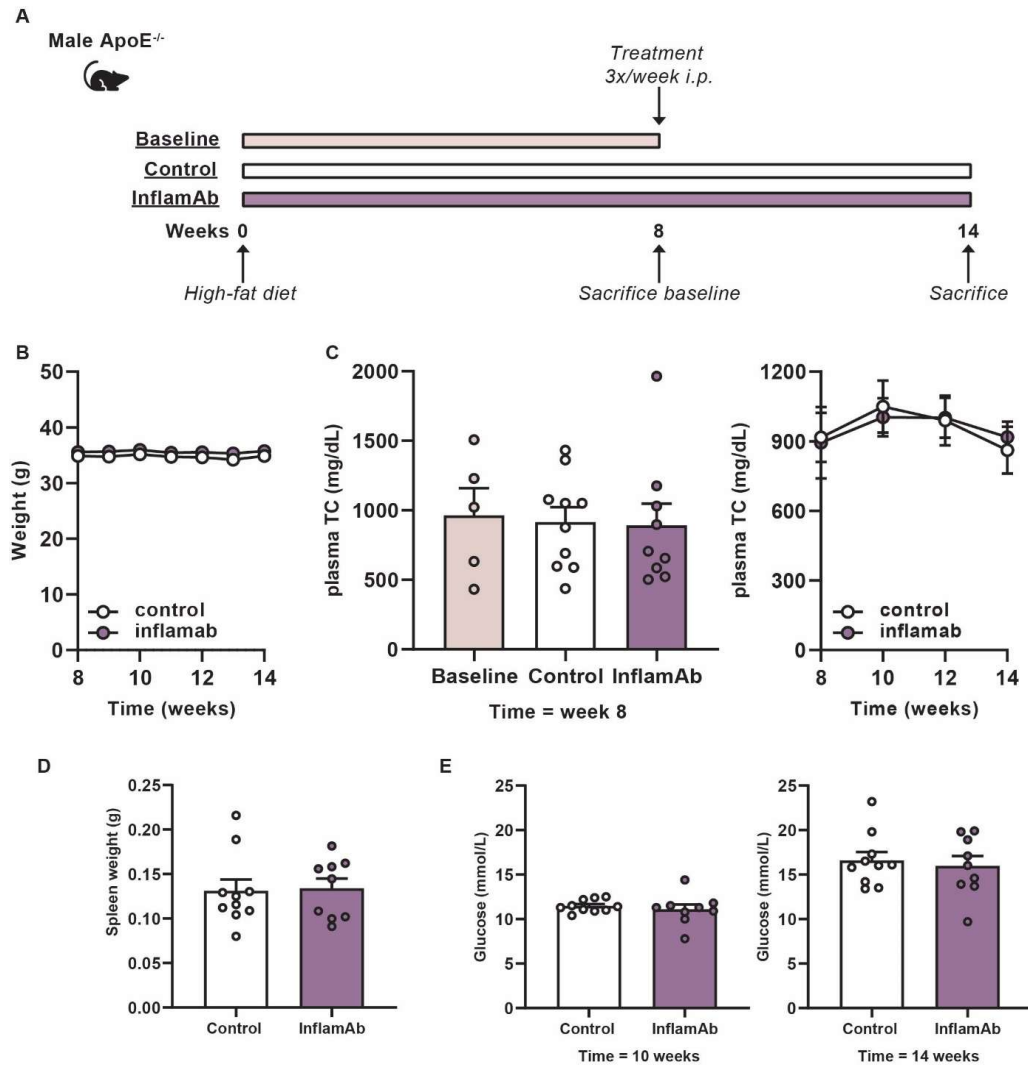


Supplemental Figure 4. InflamAb treatment during atherosclerosis development did not affect (A) mast cell numbers in the perivascular tissue of the carotid artery or (B) the mast cell activation status. A and B: n=13 control versus n=11 InflamAb treated animals, Mann-Whitney test. Mean \pm SEM.



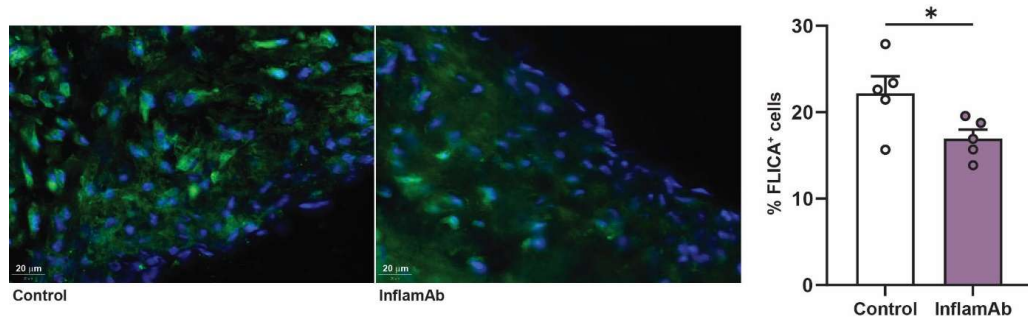
Supplemental Figure 5. Gene expression analysis of the carotid artery plaques of *Apoe*^{-/-} mice.

InflamAb during plaque development did not affect mRNA expression levels of (A) NLRP3 and (B) IL-1R1. A and B: 2-4 pooled carotids per sample, unpaired t-test. Mean ± SEM.



Supplemental Figure 6. InflamAb treatment of advanced atherosclerotic lesions in *ApoE*^{-/-} mice did not affect the body weight and plasma total cholesterol levels. (A) Schematic overview of the study set-up. Male *ApoE*^{-/-} mice were placed on a high-fat diet for 8 weeks to develop advanced lesions. In week 8 a baseline group was sacrificed and the remaining mice were treated with InflamAb or isotype control antibody 3 times per week intraperitoneally for 6 weeks, while being fed a high-fat diet. (B) Body weight was not affected upon InflamAb treatment during the study. (C) Plasma total cholesterol levels did not differ between the three groups at week 8 and were not affected by InflamAb during the treatment period. (D) Spleen

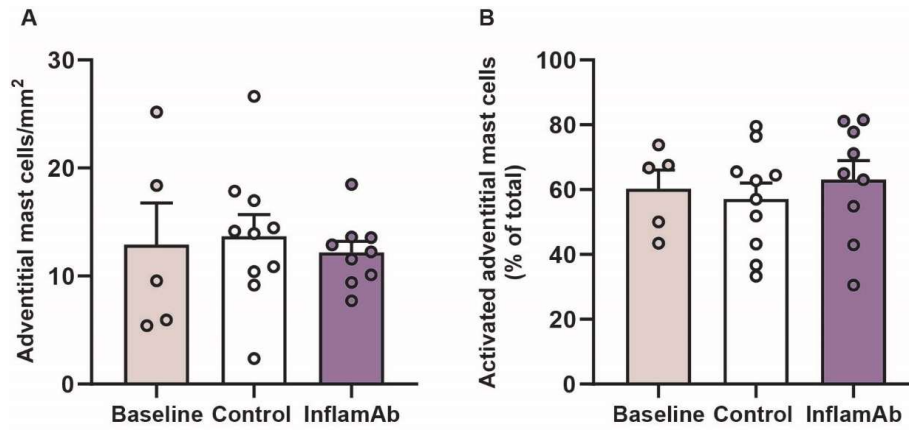
weight was not affected by InflammAb treatment. (E) Plasma glucose levels did not differ between the control and InflammAb treated mice at 10 or 14 weeks of the experiment. B, C (right panel), D and E: n=10 control versus n=9 InflammAb treated animals, unpaired t-test. C (left panel): baseline n=5, control n=10 and InflammAb n=9 animals, Kruskal-Wallis test. Mean \pm SEM.



Supplemental Figure 7. Representative images of the FLICA signal in green and DAPI staining in blue in a control and an InflamAb treated aortic root plaque. InflamAb treatment significantly reduced the percentage of FLICA⁺ cells in the plaques. Control n=5 and InflamAb n=5 animals, unpaired Student's t-test. Mean ± SEM.



Supplemental Figure 8. Representative images of the necrotic core analysis showing vessel outline in white, and necrotic areas outlined in pink. The necrotic area was defined as the acellular, debris-rich plaque area.



Supplemental Figure 9. InflammAb treatment of advanced atherosclerotic lesions in *ApoE*^{-/-} mice did not affect (A) adventitial mast cell numbers in the aortic root or (B) the mast cell activation status. A and B: baseline n=5, n=10 control versus n=9 InflammAb treated animals, ordinary one-way ANOVA. Mean ± SEM.