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A CD31-derived peptide prevents angiotensin II-induced atherosclerosis progression and aneurysm formation

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INTRODUCTION

Complex inflammatory processes underlie the progression of atherosclerosis towards its complications¹ such as coronary plaque thrombosis² and abdominal aortic aneurysms (AAA)³. Several antigens become immune targets in atherosclerotic patients^{4, 5} possibly because the regulation of immune responses is defective^{6, 7}. The trans-homophilic⁸ inhibitory immunoreceptor CD31 (PECAM-1)⁹ is expressed exclusively and constitutively by the cells of the blood-vessel interface and may therefore play a major role in vascular homeostasis¹⁰. Notably, experimental studies have shown that CD31 signaling is necessary to prevent blood leukocyte cell-cell adhesion¹¹, chronic inflammatory diseases¹², and platelet thrombosis¹³.

We previously observed that a reinforcement of the physiologic CD31 T-cell regulatory pathway, operating before the development of the disease, prevents plaque development in atherosclerosis-prone mice¹⁴. However, this approach requires the presence of the trans-homophilic CD31 extracellular domains¹⁴ which are typically lost on peripheral T-cells of mice¹⁵ and patients¹⁶ that have already developed atherosclerotic complications. Recently, a new therapeutic option has arisen from our latest data showing that a truncated extracellular CD31 fragment is indeed expressed by T-cells that apparently lack CD31¹⁷ and that a CD31-derived peptide is able to engage this fragment. In particular this peptide showed an immunosuppressive effect *in vivo* through restoration of the CD31 inhibitory pathway¹⁷. The aim of this study was therefore to evaluate whether restoring the CD31-mediated regulatory pathway with this peptide could harness the inflammatory responses underlying atherosclerosis progression and aneurysmal complication in an experimental model. We chose to test this hypothesis in aged apolipoprotein E knockout (ApoE KO) mice submitted to chronic infusion of angiotensin II because, as in patients¹⁸, angiotensin II promotes atherosclerotic progression and AAA formation via its pro-inflammatory effect in this model^{19, 20}.

METHODS

Peptides. The murine synthetic CD31 peptide (aa 551-574, MW 2606.0) (purified or 5,6-FAM-conjugated, >95% pure) was synthesized by Genosphere (France) or Mimotopes (Australia) and dissolved at 1 mg/ml in sterile PBS (DMSO 0.5%). Endotoxin levels were consistently < 0.01 ng/μg of peptide as determined by the LAL test.

Mice. Male, 28-week old ApoE KO mice (B6.129P2-Apoe^{tm1Unc}/Crl, Charles River France) were maintained on a regular chow diet under standard conditions. The experiments were repeated four times and included 2 groups (n =8-10 mice/group) assigned to the administration of 50 μl of either the peptide solution (“peptide” group) or of vehicle alone (“control” group). The dose of the murine CD31 peptide (1.5 mg/Kg/d) was chosen on the basis of our previous *in vivo* studies¹⁷. The treatment was administered subcutaneously, for 28 days during which Angiotensin II (Sigma, #A9525) was infused (1 mg/kg/d) using osmotic mini-pumps (Alzet, #2004) as previously described²¹. At the end of the study, mice were euthanized by exsanguination under anesthesia (i.p. injection of Ketamine-HCl 100mg/Kg and Xylazine 20 mg/kg). Blood was withdrawn from the right heart ventricle and collected in heparinized and EDTA tubes for blood cell and plasma analysis. The heart and the aorta were dissected for AAA assessment, measurement of plaque size and phenotypic analysis. Four additional mice of the “control” group received a single injection of 1.5 mg/Kg fluorescent (5,6-FAM-conjugated) peptide, 30 minutes before euthanasia (day 28). All the investigations conformed to the Directive 2010/63/EU of the European Parliament and formal approval was granted by the Local Animal Ethics Committee (Comité d'éthique Bichat – Debré).

AAA and atherosclerotic lesions. The presence of an AAA was blindly assessed by two investigators (A-T. G. and A. N.). Plaque size was measured on oil-red-O stained frozen cross-sections of the aortic root, as previously described¹⁵. Morphometric analysis was performed on Masson's trichrome stained slides. Adventitial cell infiltrate was calculated as the fraction of the

total surface area of the tissue outside the external elastic lamina occupied by nuclei (black); plaque extracellular matrix density was calculated as the ratio between the green-stained surface area and the surface of the plaque. All computer-assisted image analyses were performed using the Leica Qwin® software.

Blood cells and plasma analysis. Blood was centrifuged at 900 g for 30 minutes to separate plasma. The Th1/Th2/Th17 CBA kit (BD Biosciences) was used to measure cytokines in the plasma samples. Cholesterol was measured using Infinity® Cholesterol-Liquid reagents and an Olympus AU-400 multiparametric analyzer. Leukocyte pellets were stained with CD3-Alexa Fluor®700, CD4-PerCP, CD8-Pacific Blue, CD19-APC, CD69-PE-Cy7, CD115-PE, Ly-6G-APC-Cy7, CD31-FITC, (all from BD Biosciences) or with anti-mouse CD4-PerCP, CD25-APC (clone PC61, BD) and intracellular FoxP3-PE (clone PE-FJK-16s, eBiosciences). Analysis was performed using a LSRII® flow cytometer and BD FACSDiva® Software 6.0.

T-lymphocyte function. TCR stimulation. CD4⁺ splenocytes (CD4 T Lymphocyte Enrichment Set, BD Biosciences) were plated at 0.2×10^6 cells/well in U bottom 96-well plates (Costar®) pre-coated with anti-mouse CD3 ϵ antibody (BD Biosciences) and cultured for 3 days in the presence of 1 μ g/ml anti-mouse CD28 antibody (BD Biosciences) and different concentrations of the CD31 peptide. Cytokines were measured in the supernatants using the Th1/Th2/Th17 CBA kit (BD, Biosciences) and T-cell proliferation was assessed using the CellTrace™ CFSE Cell Proliferation Kit (Invitrogen). Phospho-SHP2 was evaluated by flow cytometry on TCR-stimulated splenocytes as detailed in the supplementary method section. **Antigen-specific stimulation.** Four individual ApoE KO male mice aged 24 weeks were immunized (4 footpads) with 100 μ g autologous oxidized (ox) LDL (prepared as previously described²²) in complete Freund's adjuvant. Ten days later, the animals were euthanized and the popliteal and axillary lymph nodes draining the immunization sites were collected. Single cell suspensions obtained from pooled lymph nodes from each mouse were stained with CFSE (5 μ M) and plated in complete medium in 96-well plates

(round bottom, 2×10^5 cells/well) in the presence of oxLDL (1 $\mu\text{g/ml}$) and in the presence of increasing doses of CD31 peptide (0, 12.5, 25, 50 and 100 $\mu\text{g/ml}$). Culture supernatants were harvested four days later for soluble CD31 analysis and cells were submitted to cytometry to analyze the CFSE-based proliferation index within the CD4⁺ lymphocyte gate. The mean fluorescent intensity of surface CD31 was analyzed in the same wells and soluble CD31 concentration was assessed in the supernatants (supplementary methods).

Macrophage function. Bone marrow-derived macrophages were differentiated and IFN γ -primed as previously described²³. Cells were then cultured in complete medium and stimulated with 100 ng/ml angiotensin II in the presence of increasing doses (0, 25, 50 $\mu\text{g/ml}$) of CD31 peptide. Macrophage intracellular MMP-2/9 activity was quantified by flow cytometric detection of a fluorescent enzyme substrate, according to the manufacturer's instructions (Invitrogen). Briefly, 10 $\mu\text{g/ml}$ fluorescein-conjugated DQTM type IV collagen, analog of the MMP-2/9 natural substrate, were added to stimulated macrophages for six hours and the fluorescence derived from the enzyme-driven hydrolysis of the fluorescein-conjugated DQTM was quantified by flow cytometry. IL-1 α , IL-1 β , IL-12/IL-23p40, RANTES, IL-6, MCP-1 and MIP-1 α and MIP-1 β (CBA flex sets, BD Biosciences) were measured in the culture supernatants from each condition. Individual mouse data (n =4), expressed as Median Fluorescence Intensity (MFI) have been used for statistical analysis.

Statistical Analysis. Data are expressed as means \pm SD unless otherwise indicated in the text. Differences between groups were analyzed by Student t-test or chi-squared test, as appropriate. Differences were considered statistically significant when the p value was ≤ 0.05 .

RESULTS

Loss of T-cell extracellular CD31 and development of AAA

Plaque-infiltrating leukocyte analysis showed that CD4⁺ cells were all positive for intracellular CD31 but not all of them co-expressed extracellular CD31 (supplementary Figure 1a). The lack of extracellular CD31 was also specifically observed on peripheral blood CD4⁺ T lymphocytes and associated with the development of an AAA (supplementary Figure 1b and 1c).

CD31 peptide treatment reduces AAA formation and plaque growth

Since AAA development was associated with the loss of extracellular CD31 on CD4⁺ T-cells we reasoned that this receptor could be shed from mouse leucocytes as we have previously described for human cells¹⁷. Accordingly, we evaluated the therapeutic potential of a CD31-derived peptide, which is able to restore the lost inhibitory function *in vitro* and *in vivo*¹⁷ in this model. CD31 peptide treatment significantly reduced the incidence of AAA formation (Figure 1a) and the extent of atherosclerotic lesions (Figure 1b and 1c). Aortic root adventitial cell infiltrate was associated with the presence of an AAA (Figure 1d and 1e) and was also reduced in peptide-treated mice. The peptide stabilized the phenotype of the plaques as determined by higher extracellular matrix density (44.52 ± 5.5 vs 31.61 ± 4.9 a.u, $p=0.05$), lower expression of VCAM-1 and higher α SMA content (supplementary Figure 2). Fibrillar collagen was more abundant in the aortic root plaques and significantly higher at the site of the abdominal aorta that is prone to dissection in this model, suggesting that the peptide treatment confers a higher resistance against rupture to the arterial wall (supplementary Figure 3). Furthermore, the presence of T-cells (CD4⁺), macrophages (Mac3⁺) and MMP9 within the plaques was also blunted by the treatment (supplementary Figure 2). CD31-treatment did not affect body weight (34.8 ± 0.8 vs 36.4 ± 0.9 g), lipoprotein profile (supplementary Figure 4) or plasma levels of total cholesterol (11.9 ± 0.75 vs 10.9 ± 0.6 mmol/l), HDL (1.2 ± 0.1 vs 1.1 ± 0.1 mmol/l), and triglycerides (1.1 ± 0.1 vs 1.2 ± 0.1 mmol/l). No

differences could be found between the two groups in terms of total IgG and IgM and of specific anti-oxLDL IgG and IgM serum levels (supplementary data, Figure 5) or circulating blood cell count (supplementary Table 1).

CD31 peptide modulates the peripheral T-cell compartment *in vivo* and directly inhibits T-cell responses *in vitro*

Peptide-treated mice showed increased CD4⁺CD25⁺FoxP3⁺ (Tregs) and reduced CD69⁺ (recently activated) T-cell % in the peripheral blood (Figure 2a). The treatment decreased plasma levels of IL-2, IFN γ , IL-4 and IL-6 (Figure 2b) but not those of TNF α , IL-10 or IL-17A (data not shown).

We therefore assessed if the peptide exerts direct T-cell suppressive effects *in vitro*. Our data show that, as for human cells ¹⁷, the peptide reduces mouse T-cell proliferation in response to TCR engagement (Figure 3a). In addition, it inhibited the production of IL-2, IFN γ , IL-4, IL-6, TNF and IL-10 (Figure 3b). To assess whether the peptide triggered the physiological T-cell CD31 inhibitory signaling, we evaluated the levels of the SHP2 phosphorylation [tyrosine (Y) 542] in TCR-stimulated splenocytes. Similar to the specific control (antibody-mediated cross-linking of surface CD31) a significant increase in intracellular SHP2 pY542 was induced by the peptide, in a dose-dependent manner (Figure 3c).

The peptide was also able to suppress the CD4⁺-specific immune response directed against ox-LDL, a major atherosclerosis-related antigen (Figure 4a). Interestingly, the expression of CD31 extracellular domains at the cell surface was reduced after antigenic T-cell stimulation, concomitantly with an increase in soluble CD31 concentration in the supernatants, further adding to the evidence that CD31 shedding is driven by (antigen-specific) TCR-mediated T-cell activation (Figure 4b and 4c).

CD31 peptide prevents angiotensin II-induced macrophage activity *in vitro*

While T cells are certainly involved in the acceleration of plaque formation in the angiotensin II infusion model, macrophages may play a more important role in AAA formation^{24, 25}. We therefore assessed whether the peptide is able to bind to macrophages *in vivo* and to exert a direct effect on angiotensin II-induced macrophage activation, *in vitro*. Fluorescent peptide tracking showed that plaque-infiltrating and peri-aneurysmal macrophages could indeed be targeted by the peptide, *in vivo* (Figure 5). *In vitro*, the CD31 peptide reduced intracellular MMP-2/9 activity (Figure 6a and 6b) and release of IL-6, MCP-1, MIP-1 α and MIP-1 β in the culture medium (Figure 6c). The concentrations of IL-1 α , IL-1 β , IL-12 and RANTES were not affected by the peptide (data not shown).

DISCUSSION

The present study provides further evidence for a protective role of T-cell CD31 against atherosclerotic complications, such as AAA. Indeed, as in patients¹⁶, the presence of an AAA was linked to the loss of CD31 on both plaque-infiltrating and peripheral T-cells in angiotensin II-infused apoE KO mice. We also found that the incidence of AAA correlated with adventitial cell infiltration at the level of the aortic root, distant from the abdominal aorta. While such a perivascular leukocyte infiltration had previously been described at the site of aneurysm formation²⁴, our findings suggest that the adventitial inflammatory process induced by angiotensin II extends to the whole arterial tree and contributes to both the acceleration of plaque-growth (aortic root) and the occurrence of wall dissection/aneurysm formation (abdominal aorta) depending on site-specific hemodynamics and leukocyte environment. In support of this hypothesis, the CD31-peptide treatment effectively reduced perivascular inflammation and equally protected against aneurysmal complications and atherosclerosis progression in this study.

Although the CD31 peptide can potentially act on CD31-positive cells other than leukocytes (platelets and endothelial cells), in this study we focused our attention on its putative effect on the cells of the immune system. Indeed T-cell activation is a key cellular process linking the hypertensive and pro-inflammatory effects of angiotensin II^{19, 26, 27}. Our data suggest that the inhibition of T-cell activation achieved *in vivo*^{17, 28} by the CD31 peptide prevents the angiotensin II-driven inflammatory vascular damage. However, the T-cell suppression cannot explain, alone, the anti-inflammatory and protective effect of the peptide in this model because angiotensin II receptors are expressed both by T lymphocytes and macrophages²⁹. The latter are particularly relevant in angiotensin II-driven arterial wall dissection and AAA formation^{24, 25}. Interestingly, the peptide was found to bind to both plaque and adventitial macrophages, which were both reduced by the treatment. In addition, the peptide inhibited macrophage collagenolytic activity and the release

of IL-6 and MCP-1 by angiotensin II-stimulated inflammatory macrophages, possibly explaining the resistance of the aortic wall to dissection in peptide-treated mice ³⁰.

In agreement with our findings in human lymphocytes ¹⁷, our *in vitro* data suggest that restoring CD31-driven intracellular SHP-2 phosphorylation in activated leukocytes constitutes a novel immunomodulatory strategy. Interestingly, it has recently been shown that the CD31-SHP2 pathway simultaneously inhibits tyrosin kinase-dependent activation while promoting T-cell survival via the Erk/MAP kinase pathway ³¹. The latter is also involved in extrathymic Treg induction ³² and function ³³. This may account for the enrichment of the Treg peripheral pool that we have observed in treated mice and represent an additional immunoregulatory mechanism exerted by the peptide.

As compared to other potential immunosuppressive atheroprotective agents, such mycophenolate mofetil ³⁴, anti-CD3 ^{35,36} and anti-CD20 ³⁷ antibodies, the peptide that we used in this study induces a “targeted” immunosuppressor effect because it does not cause lymphocyte depletion and reduces the release of selected disease-related cytokines . Moreover, since disease severity is linked to the extent of CD31 shedding from leukocytes ¹⁶, which can be easily measured in plasma samples ¹⁷, the dosing of the peptide could be finely adjusted to optimize the benefit/risk ratio of its use as a drug.

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CONFLICT OF INTEREST

None declared.

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LEGENDS

Figure 1. CD31 peptide-treatment prevents plaque growth and aneurysm formation. **a.** The incidence of AAA (% of mice/group) was significantly reduced by CD31 treatment (14.5 ± 3 vs 75 ± 7 , $*p < 0.001$). **b.** Quantification of atherosclerotic lesion surface area in serial cross-sections of the aortic root (200, 400, 600, 800 μm from the appearance of the first cusp) in control (■, n=10) and CD31 peptide-treated (□, n=8) mice. $*p < 0.05$ vs control. **c.** Representative micrographs of oil-red-O stained sections. **d.** Representative images showing Masson's trichrome staining of aortic root cross-sections. Control mice showed increased adventitial cell infiltration (nuclear staining) as compared to peptide-treated mice. **e.** Quantitative analysis of the adventitial cell infiltration. Peptide-treated mice (n=8) showed significantly reduced ($*p < 0.01$) cell infiltration as compared to control mice (n=8). Interestingly, a higher adventitial nuclear density was correlated with the presence of abdominal aortic aneurysms (AAA, red dots; no AAA, green dots).

Figure 2. Immunoregulatory effects of the CD31 peptide *in vivo*. **a.** The percentage of Tregs ($\text{CD}25^+ \text{FoxP}3^+$) was increased while the relative fraction of activated T-cells ($\text{CD}69^+$) $\text{CD}4^+$ T-cells among total blood $\text{CD}3^+$ T-cells was diminished by the peptide treatment ($*p < 0.05$). **b.** Plasma IL-2, $\text{IFN}\gamma$, IL-4 and IL-6 (a.u.) were significantly decreased by the peptide treatment *in vivo* ($*p < 0.05$).

Figure 3. The CD31 peptide inhibits TCR-stimulated T cell responses *in vitro* and drives SHP2 phosphorylation. **a.** Proliferation in response to TCR engagement is inhibited as a function of dose by the CD31 peptide ($*p < 0.05$ vs dose "0"). Proliferation was analyzed by the Modfit® software and expressed as "Prolif Index". The dotted line represents baseline proliferation (Prolif Index for unstimulated $\text{CD}4^+$ cells). **b.** Cytokine concentrations (pg/ml) in the supernatants of stimulated $\text{CD}4^+$ T-cells were significantly reduced by the peptide. ($*p < 0.05$ vs dose "0"). **c.** Flow

cytometric quantification of SHP2 pY542 in stimulated splenocytes. Dotted line (baseline) represents SHP2 phosphorylation induced by TCR engagement alone. Crosslinking of the TCR with surface CD31 molecules induced SHP2 phosphorylation (* $p < 0.01$ vs baseline); Increasing doses of CD31 peptide induced further increments of SHP2 phosphorylation (* $p < 0.01$ vs baseline). Data are expressed as Median Fluorescent Intensity (MFI).

Figure 4. oxLDL stimulation drives the cleavage and shedding of CD31 and the CD31 peptide inhibits oxLDL-specific T-cell responses. **a** The peptide inhibits the proliferation of oxLDL-specific CD4⁺ splenocytes derived from oxLDL-immunized apoE KO mice (* $p < 0.05$ vs dose “0”). Proliferation was analyzed at 4 days after a challenge with oxLDL *in vitro* by the Modfit® software and expressed as “Prolif Index”. **b.** Representative density plot and **c.** quantitative analysis of CD31 by flow cytometry on a CD4⁺ splenocyte surface (MFI) and by ELISA in matched culture supernatants (ng/ml). Upon oxLDL stimulation *in vitro*, the expression of CD31 decreases at the surface of CD4⁺ cells while its detection in the soluble form increases in parallel (* $p < 0.01$).

Figure 5. The CD31 peptide binds to plaque-infiltrating and peri-aneurysmal adventitial macrophages. Representative *in vivo* tracking immunofluorescent cross-section micrographs. The fluorescent peptide (green) bound to cells in clusters (blue = nuclei) including CD68⁺ cells (red, macrophages) within and surrounding abdominal aortic aneurysm (left panel) as well as within aortic root plaques (right panel). Numbered white boxes in the top panels indicate the position of the magnified insets. Scale bar=100 μ m. Hem=intramural hematoma; Adv=adventitia, EL=elastin, L=lumen.

Figure 6. The CD31 peptide inhibits macrophage response to angiotensin II. **a.** Representative phase contrast and fluorescent images of angiotensin II-stimulated macrophages in the absence (left panel) or presence (50 μ g/ml) of the CD31 peptide (right panel). Collagenolytic activity (MMP, green) was detected by the presence of the fluorogenic degradation product of DQ collagen. Blue = nuclei. Scale bar = 100 μ m. **b.** Flow cytometric quantification of the collagenolytic activity

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shown in **a**. At 50 $\mu\text{g/ml}$, the peptide significantly inhibits collagen degradation ($*p < 0.05$ vs dose “0”) MFI = median fluorescent intensity of viable macrophages in the fluorescein channel. **c**. The concentration of IL-6, MCP-1, MIP-1 α and MIP-1 β in the culture supernatants was significantly reduced by the peptide. ($*p < 0.05$ vs dose “0”).