

Deletion of MFGE8 inhibits neointima formation upon arterial damage

Viola: MFGE8 promotes neointima formation upon arterial damage

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Dear Sirs,

Atherosclerosis, a chronic inflammation of the vessel wall, is a major cause for vascular mortality due to narrowing (stenosis) of the arterial wall. In part, artery stenosis is therapeutically addressed by widening the vessel through angioplasty. Although well-established, angioplasty can damage the arterial endothelium, giving rise to an inflammatory response that leads to a neointimal hyperplasia with consequent recurrence of stenosis (1, 2). Key players in this process are leukocytes and smooth muscle cells (SMCs), as leukocyte recruitment and smooth muscle cell proliferation and migration are determinants of neointimal hyperplasia (3).

MFGE8 (Milk Fat Globule-Epidermal Growth factor 8) or lactadherin is mostly regarded as a bridging molecule with a critical function in efferocytosis and, hence, during resolution of inflammation (4). However, MFGE8 also plays a major role in promoting neovascularization (5), and, more recently, arterial MFGE8 expression emerged as a molecular hallmark of adverse cardiovascular remodeling upon ageing (6). In this context, MFGE8 is directly associated with SMC proliferation and migration (6, 7), suggesting its participation in neointima formation. We here show that MFGE8 negatively impacts on arterial restenosis and its neutralization may therefore be a potential therapeutic strategy.

To study the role of MFGE8 in arterial restenosis we subjected two groups of mice, *Apoe*^{-/-} and *Apoe*^{-/-}*Mfge8*^{-/-}, to wire injury of the left carotid artery. To simulate hypercholesterolemia, a condition often present in atherosclerotic patients, the mice were fed a high fat diet (HFD), starting one week prior to injury. Two weeks after the injury the mice were euthanized and the blood and carotids were collected. To assess neointima formation, the carotids, were stained for elastic tissue fibers with Verhoeff's van Gieson stain (EVG). Neointima area was significantly smaller throughout the injured carotid in mice lacking MFGE8 as compared to the *Apoe*^{-/-} group (**Figure 1, A-C**). No differences in blood counts were observed between the groups (**Table 1, Supplemental Data**), suggesting that the distinct neointima areas are mediated by local cells. However, cholesterol levels in the blood were higher in *Apoe*^{-/-}*Mfge8*^{-/-} mice as compared to *Apoe*^{-/-}; likely a consequence of decreased fatty acid uptake by the liver as well as small intestine, as MFGE8 has been reported to promote fatty acid uptake (8, 9). To determine which cells contributed the most for the larger neointima observed in *Apoe*^{-/-} mice, we stained the carotids with antibodies against macrophages and SMCs, since MFGE8 has been reported to be expressed in these cells (10, 11). Lack of MFGE8 did not affect neointimal macrophages, while SMC areas were vastly reduced (**Figure 1D, E**). To assess whether this was the result of reduced cell proliferation, the carotid arteries were stained with an antibody against Ki67, a cellular marker of proliferation. The staining revealed significantly less SMC proliferation in mice lacking MFGE8 as compared to their wild type littermates, suggesting a determinant role of lactadherin in restenosis formation. To confirm this observation, and verify MFGE8 as a therapeutic target, we subjected *Apoe*^{-/-} mice to arterial injury combined with the local application of siRNA, either against MFGE8 or scrambled. siRNA against MFGE8 resulted in 50% reduced expression of the protein both in the intima and the neointima (lesion) area of the carotid artery (**Figure 1, Supplemental Data**). Similar to what was observed in the knockout animal models, analysis of the injured carotids of mice treated with siRNA directed to MFGE8 showed a decreased neointima sizes as compared to mice treated with control siRNA (**Figure 1, G-I**). Equally in accordance to the studies in the knockout animal models, no differences in blood counts were observed between the groups (**Table 1, Supplemental Data**). Blood cholesterol levels remained unchanged (**Figure 2, Supplemental Data**) supporting the argument *supra* presented for the difference observed in the blood of the knockout animal models, since the siRNA effect is strictly local it does not affect fatty acid uptake. Consistent with the observations in the carotids of *Apoe*^{-/-} vs. *Apoe*^{-/-}*Mfge8*^{-/-} mice, the administration of siRNA against MFGE8 affected SMCs content in the neointima (**Figure 1K**) and its proliferation (**Figure 1L**) but the macrophage composition remained unchanged (**Figure 1J**).

Overall these results strongly point towards a relevant role of MFGE8 in post-injury arterial wall remodeling, with the potential to be exploited for therapeutic purposes. Our studies suggest these effects to be SMC-mediated, more specifically: by stimulating SMC proliferation MFGE8 promotes the formation of neointima possibly leading to hyperplasia and consequent stenosis.

Conflicts of interest

None declared.

Acknowledgements

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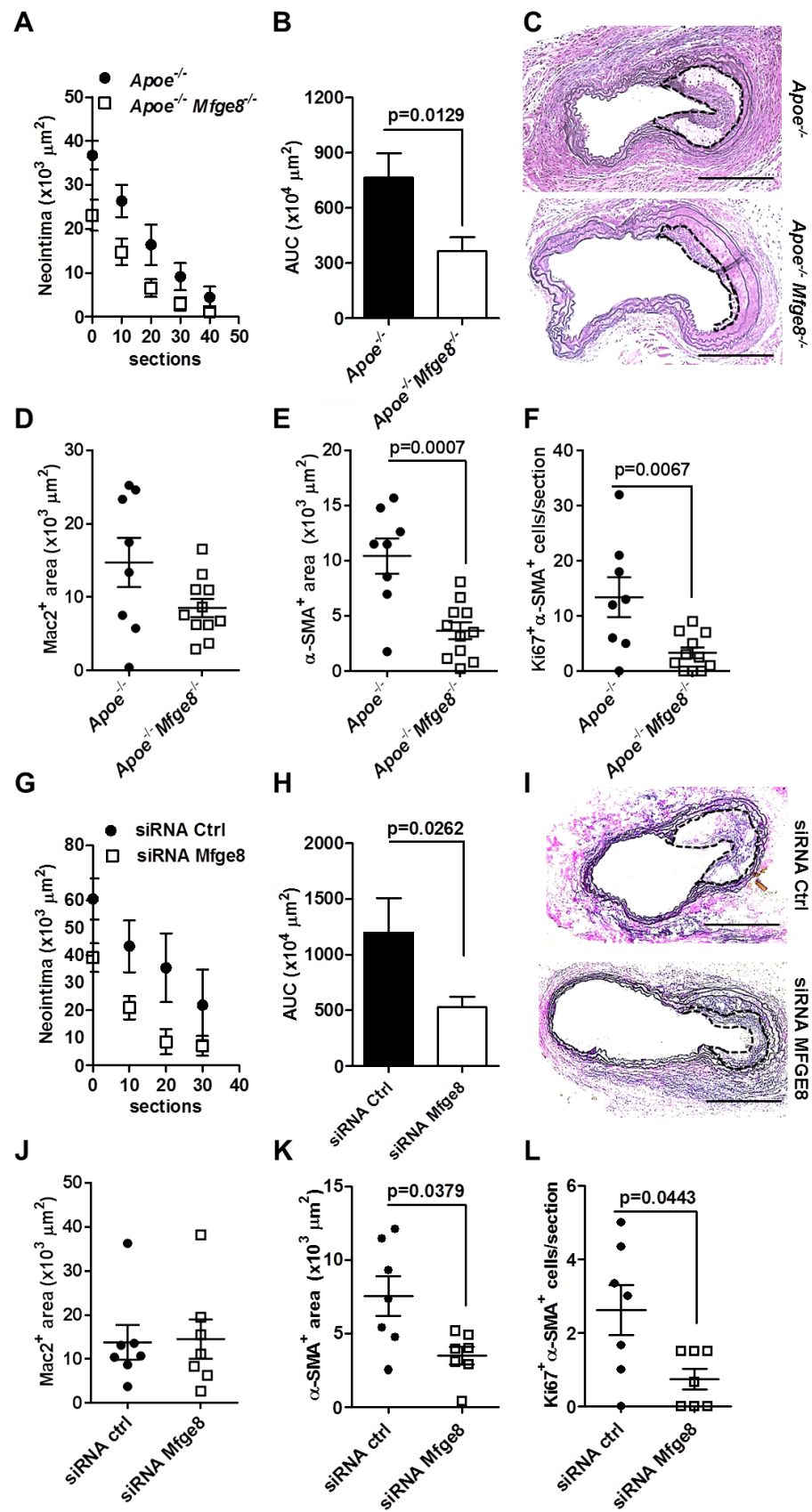


Figure 1. Mfge8 promotes post-injury neointima formation by fostering smooth muscle cell proliferation. Mice were fed a high fat diet (HFD) starting one week prior to carotid injury induction. Two weeks after injury, mice were euthanized and the blood and carotids were collected. Carotids were fixed in paraformaldehyde (PFA) and embedded in paraffin. For each antibody staining three sections per mouse were analyzed. **(A-F)** Assessment of neointima size and composition in *Apoe*^{-/-} versus *Apoe*^{-/-} *Mfge8*^{-/-} mice. (A) Neointima area was analyzed by elastic tissue fibers staining, with Verhoeff's van Gieson stain (EVG), and quantified throughout the carotids. (B) To better quantify the difference in neointima area between the two groups, the area under the curve (AUC), corresponding to the data points presented in (A), was calculated. (C) Representative images of EVG stained injured carotids. To identify possible differences in neointima composition between the two mice groups, carotids were stained with antibodies against (D) macrophages (anti-Galactin-3 or anti-Mac2) or (E) smooth muscle cells (SMCs) (anti- α -SMA). Each data point indicates the average of positive area per section in one mouse. To determine whether the increased numbers of SMCs in the neointima of *Apoe*^{-/-} *Mfge8*^{-/-} mice originated from cell proliferation or migration, carotids were stained with an antibody against a cell proliferation marker, Ki67 (F). n = 8 (*Apoe*^{-/-} mice) or n = 11 (*Apoe*^{-/-} *Mfge8*^{-/-} mice) throughout panels A-F. Statistical comparisons were made with t-test following D'Agostino Pearson omnibus normality test. **(G-L)** Assessment of neointima size and composition in *Apoe*^{-/-} mice treated with control siRNA or directed to Mfge8. (G) Neointima area was quantified throughout the carotids, and (H) the AUC, corresponding to the data points presented in (G), was calculated. (I) Representative images of EVG stained injured carotids. Carotids were stained with antibodies against (D) macrophages (anti-Galactin-3 or anti-Mac2) or (E) SMCs (anti- α -SMA). Each data point indicates the average of positive area per section in one mouse. (F) Carotids were stained with an antibody against a cell proliferation marker, Ki67. n = 7 mice per group throughout panels G-L. Statistical comparisons were made with t-test following Kolmogorov-Smirnov (KS) normality test. All data is represented as mean \pm SEM. Scale bar represents 200 μ m.

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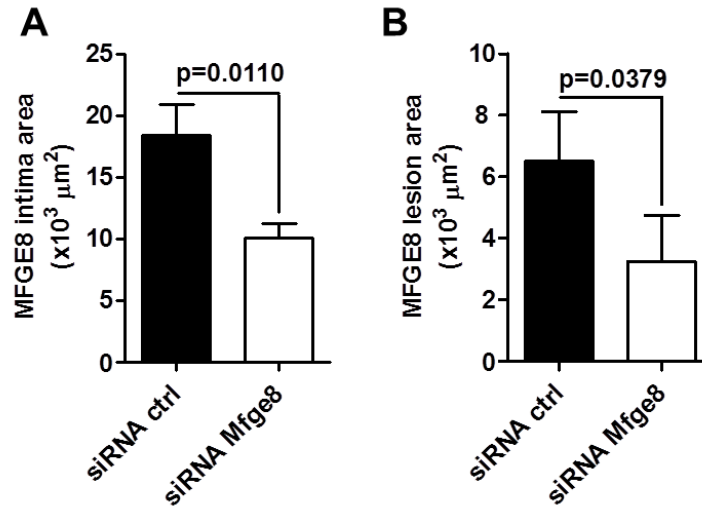
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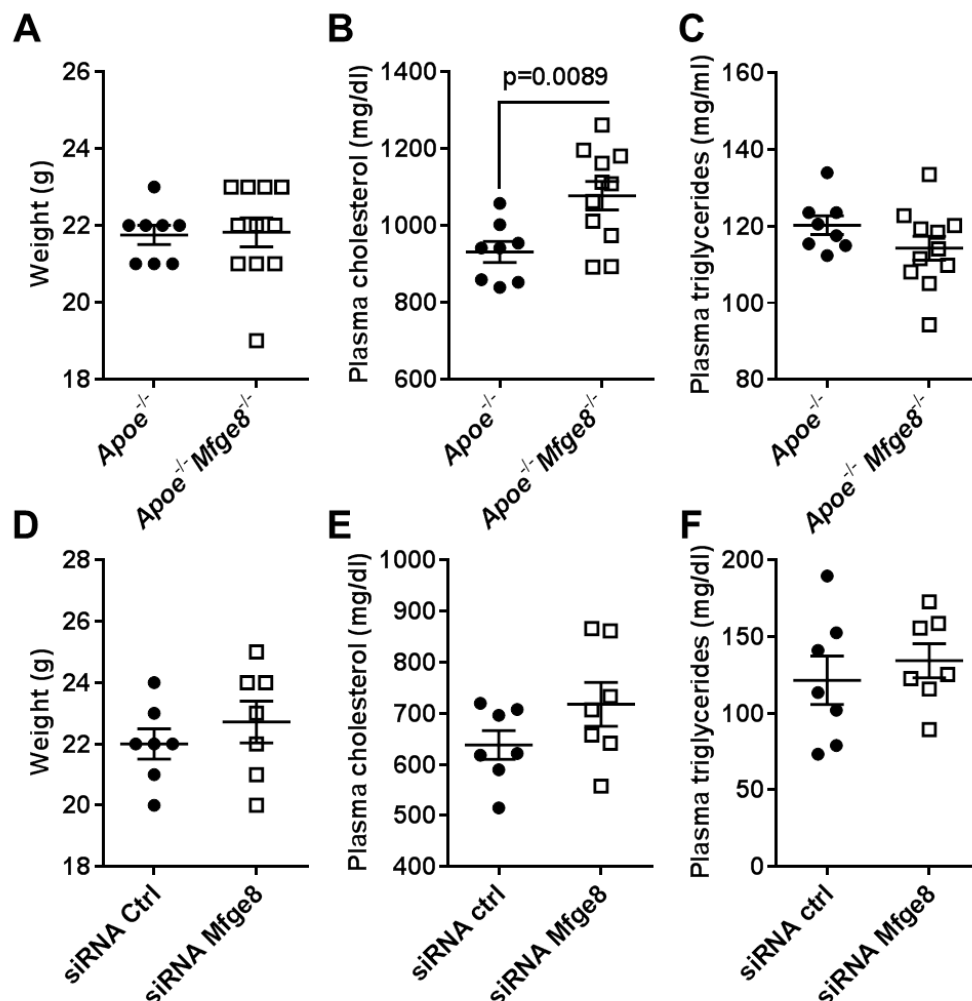
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- Supplementary Information -



Supplementary Figure 1. Delivery of siRNA against MFGE8 decreases MFGE8 expression in the carotids by 50%. Mice were fed a high fat diet (HFD) starting one week prior to carotid injury induction. siRNA (Accell siRNA, 4 nmol/treatment, Dharmacon), formulated with pluronic gel (35%) as previously described (1), was locally administered one week after injury. Two weeks after injury, mice were euthanized and the blood and carotids were collected. Carotids were fixed in paraformaldehyde (PFA) and embedded in paraffin. Three sections, per mouse, along the carotids were stained with antibody against MFGE8. The protein expression was quantified in **(A)** the intima and **(B)** in the lesion (neointima area) and is presented as positive area per section per mouse. n=7. Statistical comparisons were made with t-test following Kolmogorov-Smirnov (KS) normality test. All data is represented as mean \pm SEM.



Supplementary Figure 2. Lack of MFGE8 does not affect body weight and plasma lipid levels. Mice were fed a high fat diet (HFD) starting one week prior to carotid injury induction. Two weeks after injury, mice were euthanized and the blood and carotids were collected. (A-C) Initial studies were conducted with knockout murine models: *Apoe*^{-/-} mice versus *Apoe*^{-/-} *Mfge8*^{-/-} mice. (A) Prior to euthanasia, mice were weighted. Plasma levels of (B) cholesterol and (C) triglycerides were quantified. n=8 (*Apoe*^{-/-} mice) or n=11 (*Apoe*^{-/-} *Mfge8*^{-/-} mice). Statistical comparisons were made with t-test following D'Agostino Pearson omnibus normality test. (D-F) To confirm the initial findings, and verify MFGE8 as a therapeutic target, we subjected *Apoe*^{-/-} mice to carotid injury in combination with local administration of siRNA, either control or directed to MFGE8. (D) Prior to euthanasia mice were weighted. Thereafter, plasma levels of (B) cholesterol and (C) triglycerides were quantified. n=7. Statistical comparisons were made with t-test following Kolmogorov-Smirnov (KS) normality test. All data is represented as mean ± SEM.

| | CD45 | Monocytes | Gr1 high Monocytes | Gr1 low Monocytes | Neutrophils | CD3 | CD4 | CD8 |
|---|---------|-----------|-----------------------|----------------------|-------------|--------|--------|--------|
| <i>Apoe</i> ^{-/-} | 4833177 | 457995 | 358806 | 99258 | 1268575 | 678270 | 345503 | 193256 |
| | ± | ± | ± | ± | ± | ± | ± | ± |
| | 1874553 | 147590 | 109115 | 39903 | 567122 | 368306 | 181959 | 138842 |
| <i>Apoe</i> ^{-/-} <i>Mfge8</i> ^{-/-} | 5208352 | 379764 | 287607 | 92215 | 1037433 | 828128 | 434248 | 171364 |
| | ± | ± | ± | ± | ± | ± | ± | ± |
| | 2575619 | 322766 | 261031 | 64074 | 1329042 | 338130 | 235834 | 174823 |
| siRNA | 3933710 | 387717 | 293614 | 94226 | 1269987 | 592991 | 292864 | 216502 |
| Ctrl | ± | ± | ± | ± | ± | ± | ± | ± |
| | 1596807 | 181276 | 123926 | 71226 | 834292 | 216242 | 119140 | 86610 |
| siRNA | 3732792 | 312409 | 251532 | 60894 | 1366741 | 495049 | 239870 | 174706 |
| MFGE8 | ± | ± | ± | ± | ± | ± | ± | ± |
| | 599833 | 46083 | 40343 | 10226 | 398906 | 81113 | 32941 | 40178 |

Supplementary Table 1. Lack of MFGE8 does not affect white blood cell counts. Mice were subjected to injury in the left carotid and euthanized two weeks later. Blood was collected and blood cell counts was evaluated by flow cytometry. Cells were stained with a combination of antibodies (CD45, CD115, CD11b, CD3, CD4,CD8, Gr1). Results are presented as number of cells/ml.

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