

Rapamycin attenuates atherosclerosis induced by dietary cholesterol in apolipoprotein-deficient mice through a p27^{Kip1}-independent pathway

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Abstract

Activation of immune cells and dysregulated growth and motility of vascular smooth muscle cells contribute to neointimal lesion development during the pathogenesis of vascular obstructive disease. Inhibition of these processes by the immunosuppressant rapamycin is associated with reduced neointimal thickening in the setting of balloon angioplasty and chronic graft vessel disease (CGVD). In this study, we show that rapamycin elicits a marked reduction of aortic atherosclerosis in apolipoprotein E (apoE)-null mice fed a high-fat diet despite sustained hypercholesterolemia. This inhibitory effect of rapamycin coincided with diminished aortic expression of the positive cell cycle regulatory proteins proliferating cell nuclear antigen and cyclin-dependent kinase 2. Moreover, rapamycin prevented the normal upregulation of the proatherogenic monocyte chemoattractant protein-1 (MCP-1, CCL2) seen in the aorta of fat-fed mice. Previous studies have implicated the growth suppressor p27^{Kip1} in the antiproliferative and antimigratory activities of rapamycin in vitro. However, our studies with fat-fed mice doubly deficient for p27^{Kip1} and apoE disclosed an antiatherogenic effect of rapamycin comparable with that found in apoE-null mice with an intact p27^{Kip1} gene. Taken together, these findings extend the therapeutic application of rapamycin from the restenosis and CGVD models to the setting of diet-induced atherosclerosis. Our results suggest that rapamycin-dependent atheroprotection occurs through a p27^{Kip1}-independent pathway that involves reduced expression of positive cell cycle regulators and MCP-1 within the arterial wall.

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1. Introduction

Atherosclerosis and associated cardiovascular disease (e.g. myocardial infarction and stroke) are the major causes of mortality and morbidity in industrialized countries. Neointimal thickening is initiated by transendothelial migration and activation of circulating monocytes and lymphocytes at the sites of vessel injury [1,2]. Recruited leukocytes release inflammatory chemokines and cytokines that promote vascular smooth muscle cell (VSMC)

proliferation and migration towards the atherosclerotic lesion, thus further contributing to neointimal hyperplasia [1–4]. It has become increasingly evident that both adaptive and innate immune mechanisms modulate the inflammatory response induced in atherosclerosis, restenosis after angioplasty, and chronic graft vessel disease (CGVD) [1,5,6].

Rapamycin (Rapamune, Sirolimus), a macrolide antibiotic produced by *Streptomyces hygroscopicus* [7], has potent immunosuppressive, antiproliferative, and antimigratory properties (reviewed in [8,9]). Rapamycin exerts these effects by binding to the cytosolic immunophilin FKBP-12 (FK506 binding protein), thus inhibiting the kinase activity of the mammalian target of rapamycin (mTOR). Proposed mechanisms of rapamycin action include dephosphorylation

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and inactivation of p70 ribosomal protein S6 kinase (p70^{s6k}) and eukaryotic translation initiation factor 4E-binding protein, accumulation of the growth suppressor p27^{Kip1}, inhibition of cyclin-dependent kinase (CDK) activity, accumulation of hypophosphorylated retinoblastoma protein, and inhibition of minichromosome maintenance protein expression [9–21].

By virtue of its potent immunosuppressive activities, rapamycin has been introduced in clinic as a new effective drug for the prevention of allograft rejection [22–24]. Moreover, several animal studies have shown the efficacy of rapamycin in reducing neointimal hyperplasia, both in vessel and cardiac allografts [25–29] and in response to mechanical denudation of the vessel wall [18,26,27,30–33]. These animal studies have led to clinical trials with rapamycin-eluting stents, which have shown a significant reduction in binary restenosis, late lumen loss and repeat revascularization rates as compared with standard coronary stents [34–37].

Cell cycle progression in mammals requires the sequential assembly and activation of different CDK/cyclin holoenzymes at specific phases of the cell cycle [38]. VSMC proliferation in balloon-injured arteries is associated with a temporally and spatially coordinated expression of CDKs and cyclins [20,39]. Importantly, augmented expression of these factors coincides with increased CDK activity [39,40], demonstrating the assembly of functional CDK/cyclin holoenzymes within the injured arterial wall. Moreover, CDK2 and cyclin E expression has been detected in human VSMCs within atherosclerotic and restenotic tissue [39,41,42], suggesting that increased expression (and possibly activation) of positive regulators of cell cycle progression is a characteristic of vascular proliferative disease in humans. CDK activity is negatively regulated by the interaction with specific CDK inhibitory proteins (CKIs) [43]. It has been suggested that the CKI p27^{Kip1} functions as a negative regulator of neointimal thickening during atherosclerosis and at late phases of arterial healing after balloon angioplasty [42,44–48], at least in part via the coordinated suppression of cell proliferation and migration [49]. Exposure of cultured VSMCs and T lymphocytes to rapamycin potentially impairs their growth and migratory capacities, and these inhibitory effects correlate with p27^{Kip1} accumulation *in vitro* and *in vivo* [10,12,14,15,17,18,46,50]. However, both p27^{Kip1}-dependent [51,52] and p27^{Kip1}-independent [20,33] mechanisms of rapamycin action have been suggested (see Section 4).

In the present study, we assessed the effect of rapamycin on atherogenesis induced by dietary cholesterol in apolipoprotein E (apoE)-null mice, which develop atherosclerotic lesions that resemble those seen in humans [53,54]. We demonstrate the efficacy of rapamycin in inhibiting atherosclerosis in fat-fed apoE-null mice through a p27^{Kip1}-independent pathway associated with reduced expression of positive cell cycle regulatory proteins and attenuated monocyte chemoattractant protein-1 (MCP-1) expression within the injured arterial wall.

2. Materials and methods

2.1. Animals

Mice deficient in apoE (C57BL/6J, Taconic M&B) and doubly deficient for p27^{Kip1} and apoE [47] (backcrossed for more than five generations to a C57BL/6J background) were maintained on a low-fat standard diet (2.8% fat, Panlab, Barcelona, Spain) after weaning. At 2 months of age, mice received an atherogenic diet containing 12% fat, 1.25% cholesterol and 0.5% sodium cholate (S8492-S010, Ssniff) (4 and 6 weeks for apoE-p27^{Kip1} doubly deficient and apoE-deficient mice, respectively). Rapamycin (1 and 4 mg/kg of body weight, s.c., q.o.d.) was suspended in a vehicle solution containing 0.2% sodium carboxymethylcellulose/0.25% polysorbate 80. Control mice received vehicle.

2.2. Lipoprotein isolation and quantification of cholesterol

Blood samples were collected from the orbital sinus under anesthesia. Serum very low-density lipoprotein (VLDL) fraction was obtained by sequential-density ultracentrifugation using a fixed-angle rotor. Serum intermediate (IDL)-, low (LDL)-, and high (HDL)-density lipoprotein fractions were obtained by step-gradient ultracentrifugation using a swing-bucket rotor. The concentration of cholesterol in serum and in lipoprotein fractions was determined using an autoanalyzer Cobas Mira (Roche).

2.3. Histomorphometric studies

Fat-fed mice were euthanized and their aortas were perfusion-fixed *in situ* with 4% paraformaldehyde to quantify the extent of atherosclerosis using computerized morphometry essentially as previously described [47]. Briefly, one set of animals was used to quantify the area of Oil Red O-stained tissue in the aortic arch region (from the aortic root up to approximately 1–2 mm beyond the left subclavian artery). In another group of animals, the heart and the proximal aorta were fixed with 4% paraformaldehyde, specimens were paraffin-embedded and mounted in a Micron microtome. Once the three valve cusps were reached, sections throughout the first ~2 mm of the ascending aorta were discarded. Then, ~25 consecutive sections (5 µm thickness) were taken from 2 to 3 regions of the aortic arch separated by ~60 µm. Three cross-sections from each region were stained with hematoxylin/eosin. Specimens were examined with a Zeiss Axiolab stereomicroscope to quantify by computerized morphometry the intima-to-media ratio (I/M). For each animal, I/M was calculated by averaging all independent values.

2.4. Western blot analysis

Snap-frozen aortic tissue from fat-fed mice was pooled ($n = 4$ each group) for the preparation of whole cell

extracts in ice-cold lysis buffer (50 mmol/l Hepes [pH 7.5], 1% Triton X-100, 150 mmol/l NaCl, 1 mmol/l DTT, 0.1 mM orthovanadate, 10 mM β -glycerophosphate, 10 mM sodium fluoride) supplemented with protease inhibitor Complete Mini cocktail (Roche) using an Ultraturrax T25 basic homogenizer (IKA Labortechnik). Fifty micrograms of protein was separated onto 12% SDS-PAGE and transferred to Immobilon P (Millipore). Blots were incubated at room temperature with blocking solution (4% nonfat dry milk in PBS containing 0.1% Tween-20) for 30–40 min, followed by 1 h incubation with the following primary antibodies from Santa Cruz Biotechnology: anti-tubulin (1/200, sc-3035), anti-CDK2 (1/200, sc-163-G), and anti-proliferating cell nuclear antigen (PCNA) (1/200, sc-7907). After extensive washes with 0.1% Tween-20/PBS, the blots were incubated with species-specific secondary antibodies conjugated to horseradish peroxidase. Blots were washed twice with each 0.1% Tween-20/PBS and PBS, and immunocomplexes were detected using the ECL detection system according to the recommendations of the manufacturer (Amersham). Densitometric analysis of the blots was done using the Labimage version 2.6 software.

2.5. Quantitative RT-PCR

Total RNA was obtained from snap-frozen aortic arch tissue using the Ultraspec RNA isolation system (Biotech). Two micrograms of DNaseI-treated RNA were reverse transcribed with MuLV reverse transcriptase (Roche). Expression of MCP-1 and GAPDH mRNA was quantified by real-time PCR following the manufacturer's instructions (Lightcycler rapid thermal cycler, Roche) using the following primers specific for exon sequences: 5'-CACCA-GCAAGATGATCC-3' (MCP-1-forward); 5'-ATAAAGTTGTAGGTTCTGATCTC-3' (MCP-1-reverse); 5'-TGGGTG-TGAACACGA-3' (GAPDH-forward); and 5'-ACAGCT-TTCCAGAGGG-3' (GAPDH-reverse).

2.6. Statistical analysis

Results are reported as mean \pm S.E. In experiments with two groups, differences were evaluated using a two-tailed, unpaired *t*-test. Analyses involving more than two groups were done by ANOVA and Fisher's post-hoc test using the Statview software (SAS institute). Differences were considered significant at $P < 0.05$.

3. Results

3.1. Rapamycin attenuates diet-induced atherosclerosis in apoE-null mice

The apoE-deficient mouse [53,54] has become a valuable tool in elucidating molecular pathways implicated in atherosclerosis and in assessing therapeutic strategies

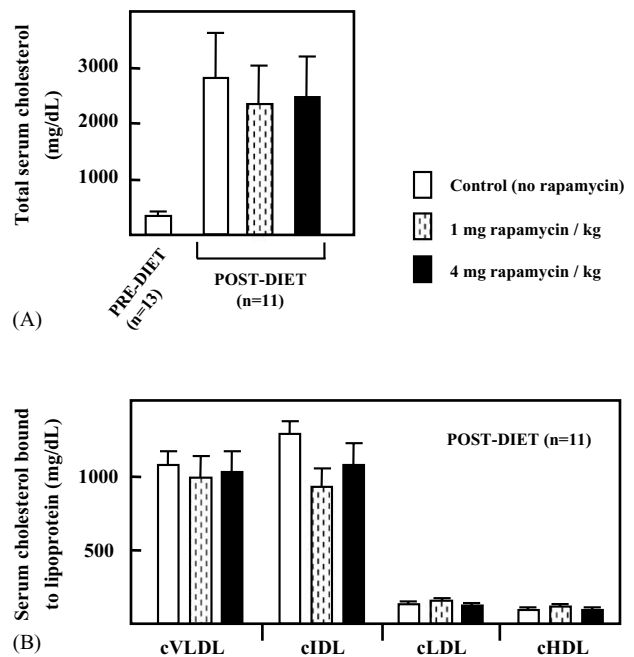


Fig. 1. Rapamycin does not affect lipid profile in fat-fed apoE-null mice. (A) Total serum cholesterol levels were measured in mice fed control chow (pre-diet) or challenged with the atherogenic diet for 6 weeks (post-diet). (B) Cholesterol levels were measured in different lipoprotein fractions isolated from the serum of fat-fed animals. All fat-fed mice displayed similar total cholesterol levels ($P > 0.05$), which were markedly increased compared with pre-diet levels ($P < 0.0001$). Cholesterol content in discrete lipoprotein fractions was also similar in all fat-fed mice ($P > 0.05$ when comparing each lipoprotein fraction among the three groups of mice). Gender distribution in each group of fat-fed mice was six males and five females.

against this disease. As expected, apoE-null mice challenged with a high-fat, cholesterol-rich diet for 4 weeks developed severe hypercholesterolemia compared with pre-diet level ($P < 0.0001$) (Fig. 1A). Importantly, total serum cholesterol level in fat-fed mice was not affected by systemic treatment with rapamycin at 1 and 4 mg/kg (RAPA1 and RAPA4, respectively, $P > 0.05$ versus control fat-fed mice) (Fig. 1A). Likewise, the amount of cholesterol associated with discrete lipoprotein fractions of fat-fed mice was unchanged in rapamycin-treated versus untreated animals (Fig. 1B). Thus, rapamycin does not affect lipid profile in fat-fed apoE-null mice.

We next examined the extent of diet-induced atherosclerosis in aortic tissue stained with Oil Red O. Consistent with numerous studies in apoE-null mice, atherosclerosis prevailed within the aortic arch in all groups of mice included in our studies (not shown). Thus, we quantified the area of atheroma in the aortic arch region by computerized morphometry using two independent approaches: (1) Oil Red O staining of whole-mounted arteries and (2) quantification of the I/M ratio in arterial cross-sections. As shown in Fig. 2A, both groups of rapamycin-treated mice displayed a significant reduction in the area of Oil Red O-stained atherosclerotic plaques as compared with untreated mice (inhibition

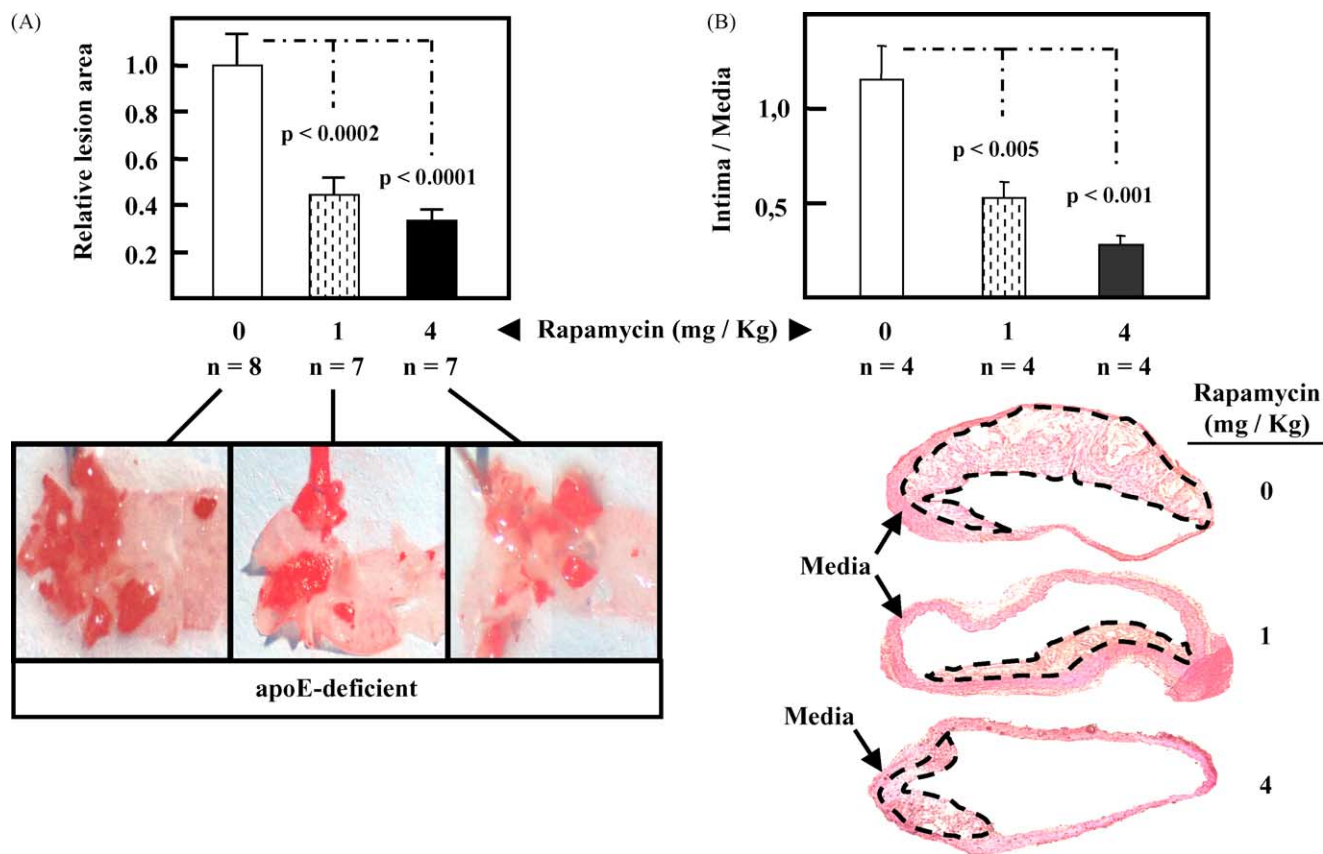


Fig. 2. Rapamycin attenuates diet-induced atherosclerosis. Atheroma development in the aortic arch of apoE-null mice fed the atherogenic diet for 6 weeks was quantified by computerized morphometry. The photomicrographs show representative examples. (A) Arteries were stained with Oil Red O (atherosclerotic lesions are shown in red). Results represent the area of lesion relative to untreated controls (1). Gender distribution was four males/four females (untreated controls), four males/three females (RAPA 1), and three males/four females (RAPA 4). (B) Cross-sections from the aortic arch were stained with hematoxylin and eosin to quantify the average I/M (two males/two females in each group). The edge of the atherosclerotic plaque (intimal lesion) has been drawn with a discontinuous line.

of 56% in RAPA1 and 66% in RAPA4, $P < 0.0002$ and <0.0001 versus control, respectively). Likewise, examination of arterial cross-sections revealed a significant reduction of the I/M in RAPA1 and RAPA4 mice ($P < 0.005$ and <0.001 versus control, respectively) (Fig. 2B). Both studies disclosed a trend towards more protection in RAPA4 versus RAPA1, although the differences between the groups of rapamycin-treated mice did not reach statistical significance. These findings demonstrate a protective effect of rapamycin against atherosclerosis in apoE-null mice challenged with an atherogenic diet in spite of sustained hypercholesterolemia.

3.2. *p27^{Kip1}* is not required for rapamycin-dependent inhibition of atherogenesis

Arterial cell proliferation is thought to contribute to atheroma development [1–4]. Because of the well-established antiproliferative action of rapamycin, we next performed Western blot analysis to examine the effect of rapamycin on the expression of positive cell cycle regulators in the aorta of fat-fed apoE-null mice. Two independent sets of mice were analyzed in these studies (experiments 1 and 2, Fig. 3). We

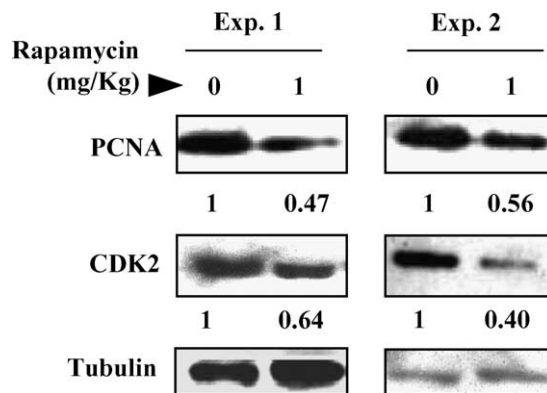


Fig. 3. Effect of rapamycin on aortic expression of cell cycle regulators. Immunoblot analysis of cell lysates prepared from the aorta of control and rapamycin-treated fat-fed apoE-null mice (pool of four arteries in each group). Densitometric analysis was performed and each value was divided by its tubulin loading control. Numbers below the blots indicate the level of expression relative to untreated mice (set as 1). The results of two independent experiments are shown.

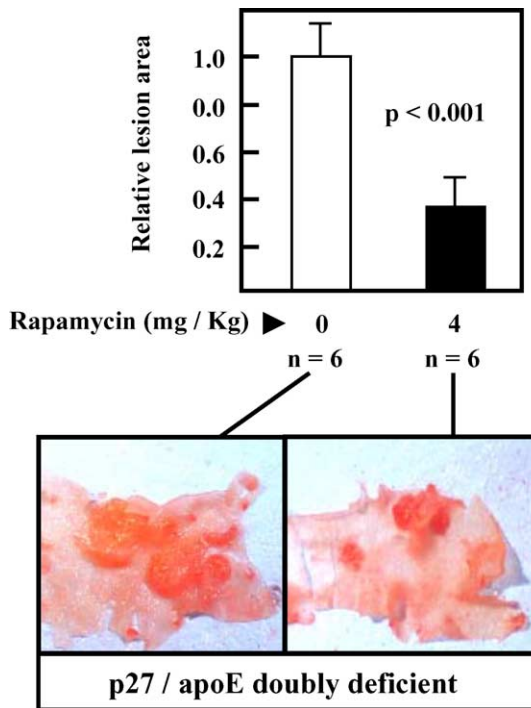


Fig. 4. Genetic disruption of p27^{Kip1} does not impair the atheroprotective effect of rapamycin. Mice doubly deficient for p27^{Kip1} and apoE were fed the atherogenic diet for 4 weeks. Gender distribution in both groups was four males/two females. Atherosclerosis was quantified in Oil Red O-stained arteries (atherosclerotic lesions are shown in red). Results represent the area of lesion relative to untreated controls (1).

found a 50–60% reduction in the expression of the S-phase markers PCNA and CDK2 in rapamycin-treated mice.

The growth suppressor p27^{Kip1} has been implicated in the control of atheroma development [42,45,47,48]. Remarkably, both p27^{Kip1}-dependent [51,52] and p27^{Kip1}-independent [20,33] mechanisms of rapamycin action have been suggested. Thus, we sought to assess the role of p27^{Kip1} on rapamycin-dependent atheroprotection by examining fat-fed mice doubly deficient for apoE and p27^{Kip1}. As shown in Fig. 4, the ability of rapamycin to inhibit atherogenesis in these mice was comparable to that seen in apoE-null mice with an intact p27^{Kip1} gene (63% inhibition, $P < 0.01$ versus control, compare with Fig. 2). These results demonstrate that rapamycin inhibits diet-induced atherosclerosis via a p27-independent mechanism.

3.3. Rapamycin prevents diet-induced aortic MCP-1 upregulation

Chemokines promote the recruitment of immune cells at the sites of vascular injury and the migration of medial VSMC towards the atherosclerotic lesion [1,2]. Gain- and loss-of-function experiments have implicated the chemotactic cytokine MCP-1 and its receptor CCR2 in the development of atherosclerosis [55–60]. Interestingly, rapamycin reportedly inhibits MCP-1 mRNA and protein expression in

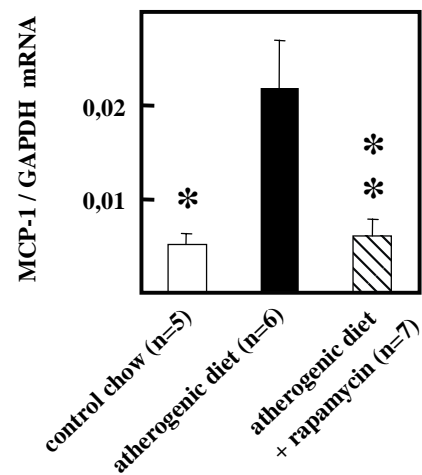


Fig. 5. Rapamycin attenuates diet-induced aortic MCP-1 upregulation. apoE-null mice were fed either control chow or the atherogenic diet for 4 weeks (with or without rapamycin at 1 mg/kg). Total RNA was isolated from the aortic arch for real-time quantitative RT-PCR analysis. The number of animals in each group is indicated (n). Results are given as the ratio of MCP-1/GAPDH. Comparisons vs. atherogenic diet (no rapamycin): * $P < 0.005$; ** $P < 0.006$.

animal models of cardiac [61] and kidney [62] transplantation. Thus, we examined the effect of rapamycin on aortic MCP-1 expression by quantitative RT-PCR analysis of RNA isolated from the aortic arch of apoE-null mice fed control chow or the atherogenic diet. As shown in Fig. 5, rapamycin blocked diet-induced upregulation of MCP-1 compared with untreated fat-fed mice.

4. Discussion

Activation of immune cells and excessive cellular proliferation and migration within the arterial wall are thought to contribute to neointimal thickening in both experimental animals and humans [1–6]. Rapamycin's immunosuppressive, antiproliferative and antimigratory actions are associated with attenuated neointimal thickening in several animal models of alloimmune and mechanical injury [18,25–33]. Moreover, rapamycin has shown promising results in reducing human coronary in-stent restenosis [34–37]. In this study, we examined the effect of rapamycin on the vessel wall response to dietary cholesterol in the apoE-deficient mouse model of atherosclerosis. We found a significant rapamycin-dependent reduction in the severity of aortic atherosclerosis in spite of sustained hypercholesterolemia compared with untreated controls. This atheroprotective effect of rapamycin coincided with reduced aortic expression of the positive cell cycle regulatory proteins CDK2 and PCNA, consistent with previous studies in the rat carotid artery model of balloon angioplasty [20].

Rapamycin induces p27^{Kip1} accumulation in vitro and in vivo [12,15,17,18,46], suggesting that p27^{Kip1} may mediate

the inhibitory effects of this drug. Consistent with this notion, p27^{Kip1} inactivation impairs rapamycin-mediated growth arrest in fibroblasts and T lymphocytes [51], and migration inhibitory responses in VSMCs [52]. However, evidence of p27^{Kip1}-independent mechanisms of rapamycin action have also been provided. First, rapamycin efficiently impaired the growth of p27^{Kip1}-null VSMCs in vitro [33]. Second, rapamycin failed to prevent the in vivo down-regulation of p27^{Kip1} seen 24 h after balloon injury of rat carotid arteries [20]. Third, attenuation of neointimal thickening after mechanical injury was similar in wild-type and p27^{Kip1}-null mice treated with rapamycin [33]. Although we can not rule out that aortic p27^{Kip1} expression may be induced by rapamycin in the present study, we found that the atheroprotective action elicited by this drug was not impaired in fat-fed mice doubly deficient for apoE and p27^{Kip1} versus apoE-null mice with an intact p27^{Kip1} gene (compare Figs. 2 and 4). Thus, p27^{Kip1} is not essential for the therapeutic effect of rapamycin against neointimal thickening induced by both dietary cholesterol and balloon angioplasty.

Human and animal studies suggest that local production of chemokines within the atherosclerotic plaque plays a critical role in atherogenesis [1,2]. Accumulating evidence has implicated MCP-1 as a proatherogenic factor [58]. For instance, several cell types involved in atheroma formation (i.e. endothelial cells, VSMCs, and macrophages) display abundant expression of MCP-1 and its receptor CCR2 [58], and high level of MCP-1 expression has been observed within the atherosclerotic plaque in both experimental animals and humans [63–65]. Importantly, genetic inactivation of MCP-1 or CCR2 [55–57] and anti-MCP-1 gene therapy [59] reduce murine atherosclerosis. In marked contrast, local infusion of MCP-1 protein increases plaque formation in apoE-null mice [60]. We found that rapamycin abrogates the upregulation of MCP-1 mRNA expression normally seen in the aortic arch of fat-fed apoE-null mice, consistent with recent studies demonstrating rapamycin-dependent inhibition of MCP-1 expression in animal models of cardiac [61] and kidney [62] allografts.

In conclusion, the present study extends previous reports documenting the therapeutic efficacy of rapamycin against neointimal thickening in the setting of CGVD [25–29] and balloon angioplasty [18,26,27,30–37] by demonstrating rapamycin-dependent reduction of atherosclerosis in apoE-null mice challenged with a high-fat, cholesterol-rich diet. Rapamycin's atheroprotective effects occur through a p27^{Kip1}-independent pathway that coincides with reduced arterial expression of both positive cell cycle regulatory factors (i.e. CDK2 and PCNA) and proatherogenic MCP-1. Because of this novel therapeutic application of rapamycin, gene expression profiling and proteomic studies comparing untreated and rapamycin-treated fat-fed animals are warranted to identify potential therapeutic targets for the prevention and/or treatment of atherosclerosis.

Note added in proofs

While this manuscript was under review, Elloso et al. reported the protective effect of rapamycin against atheroma development in apoE-null mice (Elloso MM, Azrolan N, Sehgal SN, Hsu PL, Phiel KL, Kopec CA, Basso MD, Adelman SJ. Protective effect of the immunosuppressant sirolimus against aortic atherosclerosis in apo E-deficient mice. *Am J Transplant* 2003;3:562–9).

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