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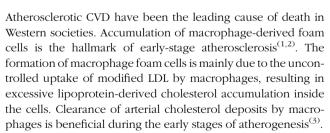
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Abstract

It has been suggested that retinoic acid (RA) has a potential role in the prevention of atherosclerotic CVD. In the present study, we used J774A.1 cell lines and primary peritoneal macrophages to investigate the protective effects of RA on foam cell formation and atherogenesis in apoE-deficient (apoE $^{-/-}$) mice. A total of twenty male apoE $^{-/-}$ mice (n 10 animals per group), aged 8 weeks, were fed on a high-fat diet (HFD) and treated with vehicle or 9-cis-RA for 8 weeks. The atherosclerotic plaque area in the aortic sinus of mice in the 9-cis-RA group was 40.7% less than that of mice in the control group (P < 0.01). Mouse peritoneal macrophages from the 9-cis-RA group had higher protein expression levels of ATP-binding cassette transporter A1 (ABCA1) and G1 (ABCG1) than those from the control group. Serum total and LDL-cholesterol concentrations were lower in the 9-cis-RA group than in the control group (P<0.05). In vitro studies showed that incubation of cholesterol-loaded J774A.1 macrophages with 9-cis-RA (0·1, 1 and 10 µmol/l) induced cholesterol efflux in a dose-dependent manner. The 9-cis-RA treatment markedly attenuated lipid accumulation in macrophages exposed to oxidised LDL. Moreover, treatment with 9-cis-RA significantly increased the protein expression levels of ABCA1 and ABCG1 in J774A.1 macrophages in a dose-dependent manner. Furthermore, 9-cis-RA dose-dependently enhanced the protein expression level of liver X receptor-α (LXRa), the upstream regulator of ABCA1 and ABCG1. Taken together, the present results show that 9-cis-RA suppresses foam cell formation and prevents HFD-induced atherogenesis via the LXRα-dependent up-regulation of ABCA1 and ABCG1.

Key words: Retinoic acid: Macrophage cholesterol efflux: ApoE-deficient mice: Atherosclerosis



Cholesterol efflux from macrophages is the first and potentially the most important step in reverse cholesterol transport, a process especially relevant to atherosclerosis and to the regression of atherosclerotic plaques (4,5). The ATP-binding cassette transporter family, including ABCA1 and ABCG1, is highly expressed in lipid-loaded macrophages and mediates cholesterol efflux⁽⁶⁻⁸⁾. ABCA1 is a key regulator of cholesterol and phospholipid export to lipid-free apolipoproteins, forming nascent HDL⁽⁹⁾. ABCG1 was identified as a more likely mediator of cholesterol transport to HDL⁽¹⁰⁾. Both ABCA1 and ABCG1 have been reported to be mediated by liver X receptors (LXR)^(11,12), the key regulators of lipid and cholesterol metabolism.

The effects of vitamin A on growth and development are achieved through the action of biologically active metabolites, retinoic acids (RA), mainly including all-trans-RA and 9-cis-retinoic acid (9-cis-RA)(13,14). RA is synthesised primarily intracellularly from retinaldehyde, which itself can be produced from retinol or provitamin A carotenoids such as

Abbreviations: 9-cis-RA, 9-cis-retinoic acid; ABCA1, ATP-binding cassette transporter A1; ABCG1, ATP-binding cassette transporter G1; FXR, farnesoid X receptor; LXR, liver X receptor; LXRE, liver X receptor response element; MPM, mouse peritoneal macrophages; PXR, pregnane X receptor; RA, retinoic acid; RPMI, Roswell Park Memorial Institute; RXR, retinoid X receptor; siRNA, small interfering RNA.

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β-carotene⁽¹⁵⁾. The broad range of biological effects exerted by RA is mainly attributable to the control of specific gene expression. RA binds to and activates the members of the nuclear receptor family including retinoic acid receptor and retinoid X receptor (RXR), which then form heterodimers with potential partners and bind to the response element in the promoter of target genes to initiate gene transcription (16-18). LXR, farnesoid X receptor (FXR) and pregnane X receptor (PXR) are three major potential partners of RXR, which modulate systematic lipid metabolism. Recent studies have confirmed the regulatory role of RA in lipid metabolism in various cell types including macrophages. RA effectively regulates the expression of ABCA1 in many macrophage cell types (19). All-trans-RA increased the protein expression levels of ABCG1 and cholesterol efflux from the macrophages⁽²⁰⁾. 9-cis-RA has also been reported to induce ABCA1 expression and cholesterol efflux in brain cells⁽²¹⁾. Although the critical role of RA in cholesterol efflux in vitro has been fully demonstrated, the impact of 9-cis-RA on macrophage foam cell formation and atherosclerosis in vivo and the underlying molecular mechanisms are still obscure, and need to be further explored.

The purpose of the present study was to investigate the protective effects of 9-cis-RA on foam cell formation and atherosclerosis in apoE $^{-/-}$ mice, and to uncover the underlying molecular mechanism.

Materials and methods

Materials

9-cis-RA was obtained from Enzo Life Sciences. apoAI was obtained from human plasma and other reagents were purchased from Sigma-Aldrich, unless otherwise indicated. liver X receptor response element (LXRE)-Luc and farnesoid X receptor response element (FXRE)-Luc reporter plasmids were purchased from Genomeditech. Pregnane X receptor response element (PXRE)-Luc reporter plasmid was constructed by cloning PXR binding sequences in the promoter area of ABCA1 and ABCG1 into the vector pEZX-PG02 upstream of the firefly luciferase reporter gene (GeneCopoeia).

Animal experiments

Male homozygous apoE^{-/-} mice (C57BL/6 genetic background), aged 8 weeks, were obtained from Jackson Laboratories and randomly divided into two groups (*n* 10 animals per group). Mice were fed on a Western-type diet (containing 0·15% cholesterol and 41% saturated fat, MD12015; Medicience Limited; Table 1) for 8 weeks. 9-cis-RA was dissolved in dimethyl sulfoxide, and after dilution with PBS containing 0·5% bovine serum albumin, a dose of 2 mg/kg was administered intraperitoneally four to five times per week (9-cis-RA group). PBS containing 0·5% bovine serum albumin was administered as vehicle to the control group. Body weight and food intake were recorded weekly. At the end of the experiment, mice were killed by CO₂ inhalation and serum and tissue samples were collected and stored at

Table 1. Composition of the experimental diet

Nutrients	High-fat diet (MD12015)
Energy	
kcal/g	4.7
kJ/g	19.7
Ingredients (g/kg)	
Casein	195.0
Maltodextrin	60-0
Sucrose	340.0
Anhydrous milk fat	210.0
Soyabean oil	20.0
Starch	55-46
Cellulose	50.0
Mineral mix*	43.0
Vitamin mix*	19.0
L-Cys	3.0
Choline bitartrate	3.0
Cholesterol	1.5
TBHQ, antioxidant	0.04
Total (g)	1000

TBHQ, tertiary butylhydroquinone.

 -80°C until analysis. All animal experiments were approved by the Institutional Animal Care and Use Committee of Sun Yat-sen University.

Measurement of serum lipid concentrations

Serum samples were collected at the end of the experiment. After 12 h fasting, venous blood was collected from the lateral saphenous vein into Eppendorf tubes. Serum was isolated by centrifugation at 2800 **g** for 20 min at 4°C, and stored at -80° C until analysis. Enzymatic colorimetric methods with diagnostic reagents from Wako on an Automatic Biochemistry Analyzer (Hitachi 7600-010) were used to determine total cholesterol (Cholesterol E, catalogue no. 439-17 501), LDL-cholesterol (L-Type LDL-C, catalogue no. 993-00 404), HDL-cholesterol (L-Type HDL-C, catalogue no. 997-72 591) and TAG (L-Type TAG M, catalogue no. 461-08 992).

Measurement of serum total retinoic acid concentrations

Serum RA concentration was measured using a commercially available ELISA kit (Biosamite), according to the manufacturer's protocol.

Evaluation of atherosclerotic lesions

Mice were perfused with 10% buffered formalin, and the heart was then dissected out. The base of each heart was frozen in O.C.T. tissue-freezing medium (Leica) and cryosectioned, as described previously⁽²²⁾. For quantification of the maximum lesion area within the aortic sinus, the cross-sectional area of Oil red O staining was measured from the points of emergence to the points of disappearance of the aortic valves.

Isolation of primary mouse macrophages

Mouse peritoneal macrophages (MPM) from the two experimental groups were isolated 3 d after intraperitoneal injection



^{*} From Reeves et al. (49).



with 4% thioglycolate solution, as described previously (23). After massage of the abdomen, MPM were collected and then centrifuged at 2000 rpm for 5 min, and resuspended in serum-free Roswell Park Memorial Institute (RPMI)-1640 medium at 37°C in a humidified atmosphere of 5% CO2 for at least 2 h. Then, cells were washed with PBS three times to remove non-adherent cells.

Cell culture and treatments

The J774A.1 cell line was obtained from the American Type Culture Collection. Cells were seeded in six-well plates at a density of 1.0×10^6 cells per well in RPMI-1640 medium (Gibco; Life Technologies) containing 10% fetal bovine serum, and were maintained at 37°C in a humidified atmosphere of 5% CO₂. Cells were treated with vehicle (dimethyl sulfoxide), 9-cis-RA (0·1, 1 and 10 µmol/l) or Z-guggulsterone (100 µmol/l) for 24 h before analysis.

Cholesterol efflux assay

J774A.1 cells were equilibrated with 12-[(7-nitro-2,1,3-benzoxadiazol-4-yl)amino]-(3S,10R,13R)-3-methoxy-10,13-dimethyl-17-((R)-6-methylheptan-2-yl)-tetradecahydro-1H-cyclopenta[α] phenanthrene (NBD-cholesterol) (1 µg/ml; Cayman Chemical) for 6 h. NBD-cholesterol-labelled cells were washed with PBS and incubated in serum-free RPMI-1640 medium containing apoAI (10 μg/ml; Calbiochem) or HDL (50 μg/ml; Calbiochem) for another 6 h. The fluorescence-labelled cholesterol released from the cells into the medium was measured with a microplate reader (Tecan Infinite F200). Cholesterol efflux was expressed as a percentage of fluorescence in the medium relative to the total amounts of fluorescence detected in the cells and the medium. Each experiment was performed in triplicate for at least three times⁽²⁴⁾.

Confocal microscopy analysis

Cells were fixed with 4% paraformaldehyde for 15 min at room temperature. Then, cells were incubated with the fluorescence probe BODIPY®493/503 (4,4-difluoro-1,3,5,7,8-pentamethyl-4-bora-3a,4a-diaza-s-indacene, 1 mg/ml; Invitrogen) for 1 h in the dark, followed by incubation with the nuclei staining 4',6-diamidino-2-phenylindole (DAPI, 5 μg/ml; Beyotime) at 37°C for another 30 min. Images were captured using the laser scanning confocal microscope (Leica TCS SP5 II). The 488-line argon laser was used to capture BODIPY[®] 493/503 lipid staining, and the 405 nm laser diode was used to capture the DAPI nuclear stain. Intracellular fluorescence intensity was quantified by ImageJ software (National Institutes of Health)⁽²⁵⁾.

Quantitative real-time RT-PCR

Total RNA was isolated using TRIzol Reagent (Invitrogen). Complementary DNA was synthesised from total RNA by RT-PCR (Takara). The resulting complementary DNA was subjected to real-time RT-PCR analysis with gene-specific primers in the presence of the SYBR Green PCR master mix in an ABI Prism VR7 instrument (Applied Biosystems). The primer sequences used were as follows: ABCA1 – forward primer 5'-AAA ACC GCA GAC ATC CTT CAG-3', reverse primer 5'-CAT ACC GAA ACT CGT TCA CCC-3'; ABCG1 – forward primer 5'-CTT TCC TAC TCT GTA CCC GAG G-3', reverse primer 5'-CGG GGC ATT CCA TTG ATA AGG-3'; LXRa - forward primer 5'-CTC AAT GCC TGA TGT TTC TCC T-3', reverse primer 5'-TCC AAC CCT ATC CCT AAA GCA A-3'; LXR\beta - forward primer 5'-ATG TCT TCC CCC ACA AGT TCT-3', reverse primer 5'-GAC CAC GAT GTA GGC AGA GC-3'; FXR - forward primer 5'-GCT TGA TGT GCT ACA AAA GCT G-3', reverse primer 5'-CGT GGT GAT GGT TGA ATG TCC-3'; PXR - forward primer 5'-TAG GGA CCT GCC TAT TGA GGA-3', reverse primer 5'-CCG TTT CCG TGT CGA ACA TC-3'; glyceraldehyde 3-phosphate dehydrogenase – forward primer 5'-CCA CAG CTG AGA GGG AAA TC-3', reverse primer 5'-AAG GAA GGC TGG AAA AGA GC-3'. Data were analysed using the $2^{-\Delta\Delta C_T}$ threshold cycle method. Glyceraldehyde 3-phosphate dehydrogenase was used as an invariant internal control.

Western blot

Cells were harvested and protein extracts were prepared as described previously (22). Aliquots (40 µg) of protein extracts were separated by SDS-PAGE and then transferred onto an Immobilon-P transfer membrane (Millipore). After blocking, the blots were incubated with various primary antibodies followed by horseradish peroxidase-labelled secondary antibodies. Protein bands were detected with an ECL Western Blotting Substrate Kit (Pierce). The primary antibodies used were as follows: anti-ABCA1 (Novus); anti-ABCG1 (Novus); anti-LXRα/β (Santa Cruz Biotechnology); anti-FXR (Bioss); anti-PXR (Proteintech); anti-glyceraldehyde 3-phosphate dehydrogenase (Cell Signaling Technology).

Luciferase assay

The 293T cell line was obtained from the American Type Culture Collection. Cells were grown in Dulbecco's modified Eagle's medium (Gibco; Life Technologies) containing 10% fetal bovine serum and were maintained at 37°C in a humidified atmosphere of 5% CO2. Cells were seeded in forty-eightwell plates and transiently co-transfected with LXRE/FXRE/ PXRE-Luc reporter plasmid (0.4 µg) and pRL-SV40 Vector (0.02 µg) with Lipofectamine 2000 (Invitrogen), according to the manufacturer's instructions. After transfection for 24h, cells were treated with vehicle (dimethyl sulfoxide) or 9-cis-RA (0·1, 1 and 10 µmol/l) for another 24 h. Subsequently, the amount of Luciferase activity was measured and normalised to the amount of Renilla luciferase activity using the Dual-Luciferase Reporter Assay System (Promega).

Liver X receptor- α small interfering RNA transfection

For the small interfering RNA (siRNA)-mediated downregulation of LXRα, LXRα-specific siRNA and non-targeting control siRNA were purchased from Santa Cruz Biotechnology. For transfection with siRNA, J774A.1 cells were plated at a density of 2.0×10^6 cells per well. Cells were transfected with LXR α



Table 2. Serum retinoic acid (RA) and lipid concentrations in apoE-deficient mice

(Mean values and standard deviations; n 10)

Variables	Control		9- <i>cis</i> -RA	
	Mean	SD	Mean	SD
TC (mmol/l) TAG (mmol/l) HDL-C (mmol/l) LDL-C (mmol/l) RA (nmol/l)	18·31 ^a 1·10 0·56 ^a 3·95 ^a – ^a *	2·77 0·35 0·09 0·76	11·74 ^b 1·16 0·74 ^b 2·92 ^b 339 ^b	3.96 0.33 0.28 0.22 34.19

TC, total cholesterol; HDL-C, HDL-cholesterol; LDL-C, LDL-cholesterol.

siRNA at a concentration of 100 nmol/l for 48 h followed by treatment with 9-cis-RA (10 \mumol/l) for an additional 24 h.

Statistical analysis

All experiments were carried out for at least three times. Data are presented as means and standard deviations. Statistical analysis was performed using SPSS (version 17.0, IBM SPSS Statistics; IBM Corporation). Significance was determined by two-tailed Student's t test or ANOVA. P < 0.05 was considered as statistically significant.

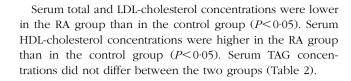
Results

Body weight and food intake

Body weight did not differ between the two groups at baseline (18·5 (sp 1·74) g for the control group and 17·8 (sp 2·91) g for the RA group) or at the end of the experiment (31·23 (sp 3·56) g for the control group and 28·50 (sp 2·90) g for the RA group). There was no difference in daily food intake between the control group (2·50 (sp 0·07) g/d) and the RA group (2·55 (sp 0·07) g/d). The amount of RA injected was 0·05 (sp 0·01) mg/d per mouse.

Serum retinoic acid and lipid concentrations

The serum concentration of RA was <10 nmol/l (assay sensitivity 10 nmol/l) for the control group and 339 (sp. 34·19) nmol/l for the RA group (Table 2).



Atherosclerotic plaque area

The mean atherosclerotic lesion area in the aortic sinus of the RA group (29 288·77 (sp. 1472·92) μ m²) was 40·7% less than that of the control group (49 397·86 (sp. 1179·69) μ m²) (P<0·01) (Fig. 1(a) and (b)).

ATP-binding cassette transporter A1/ATP-binding cassette transporter G1 expression and lipid accumulation in mouse peritoneal macrophages

MPM isolated from apoE^{-/-} mice in the RA group had higher protein expression levels of ABCA1, ABCG1 and LXRα compared with those in the control group (all P<0.05; Fig. 2(a) and (b)). The results of BODIPY[®]493/503 lipid staining showed that the lipid content in MPM from the RA group was significantly reduced compared with the control group (P<0.01; Fig. 2(c) and (d)).

Retinoic acid induces cholesterol efflux from macrophages

Incubation of J774A.1 macrophages with increasing concentrations of 9-cis-RA dose-dependently increased NBD-cholesterol efflux to apoAI compared with the control treatment (P<0·01; Fig. 3(a)). Moreover, the 9-cis-RA treatment also dose-dependently promoted NBD-cholesterol efflux to HDL to a less extent compared with the control treatment (P<0·05; Fig. 3(b)).

Retinoic acid decreases intracellular lipid accumulation in macrophages

Incubation of cholesterol-preloaded J774A.1 macrophages with 9-cis-RA significantly decreased intracellular lipid accumulation compared with the control treatment (P<0·01; Fig. 4(a) and (b)).

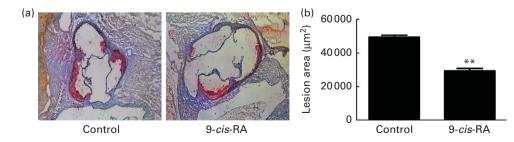


Fig. 1. Atherosclerotic lesions in the aortic sinus of apoE-deficient mice fed a high-fat diet for 8 weeks and treated with 9-cis-retinoic acid (9-cis-RA) or vehicle. (a) Representative photomicrographs of the aortic sinus stained with Oil red O. Magnification $400 \times$. (b) The lesion area in the aortic sinus was calculated using Leica software version 3.8. Values are means (n 10), with their standard deviations represented by vertical bars. ** Mean value was significantly different from that of the control group (P<0.01). (A colour version of this figure can be found online at http://www.journals.cambridge.org/bjn).



a.bMean values within a row with unlike superscript letters were significantly different (P<0.05).</p>

^{*} Serum 9-cis-RA concentration for the control group was below the detection limit.

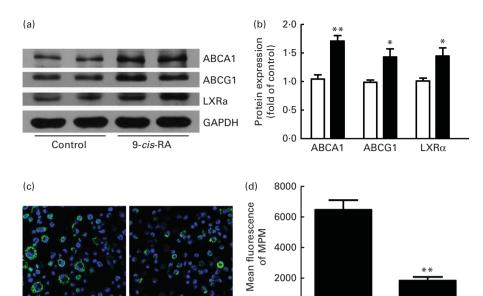


Fig. 2. ATP-binding cassette transporter A1 (ABCA1)/ATP-binding cassette transporter G1 (ABCG1) expression and intracellular lipid accumulation. ApoE-deficient mice fed a high-fat diet were treated with 9-cis-retinoic acid (9-cis-RA; 🔳) or vehicle (control; 🗆) for 8 weeks. (a) Protein expression levels of ABCA1, ABCG1 and liver X receptor-α (LXRα) in mouse peritoneal macrophages (MPM). (b) Quantification of ABCA1, ABCG1 and LXRα protein density normalised to the expression level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH), and expressed as optical density relative to that of the control. (c) Overlap images of BODIPY®493/503 and DAPI (4',6-diamidino-2-phenylindole) of MPM. Magnification 400×. (d) Mean fluorescence was calculated using ImageJ software. Values are means (n 10), with their standard deviations represented by vertical bars. Mean value was significantly different from that of the control group: *P<0.05, **P<0.01. (A colour version of this figure can be found online at http://www.journals.cambridge.org/bjn).

9-cis-RA

O

Control

Retinoic acid increases ATP-binding cassette transporter A1 and ATP-binding cassette transporter G1 expression in macrophages

Control

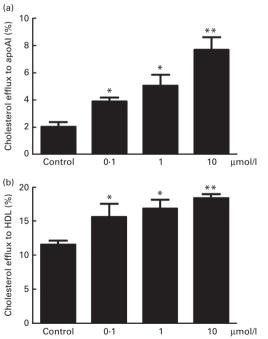
Compared with the control treatment, 9-cis-RA-treated J774A.1 macrophages increased the mRNA expression levels of ABCA1, ABCG1 and LXRα in a dose-dependent manner (P < 0.05) (Fig. 4(c)). The treatment of J774A.1 macrophages with 9-cis-RA dose-dependently increased the protein expression levels of ABCA1, ABCG1 and LXRα relative to the control treatment (P < 0.05; Fig. 4(d) and (e)). Furthermore, the transcriptional activity of LXRE-Luc was markedly induced in response to the 9-cis-RA treatment in 293T cells in a dose-dependent manner (P < 0.01; Fig. 4(f)).

Liver X receptor- α mediates retinoic acid-induced macrophage cholesterol efflux

Compared with the cells treated with control siRNA, reduced cholesterol accumulation induced by 9-cis-RA was largely abolished in the presence of LXR α siRNA (P<0.01; Fig. 5(a) and (b)). The knockdown efficiency of LXR α siRNA is shown in Fig. 5(c) and (d) (P<0·01). In addition, LXRα knockdown significantly suppressed 9-cis-RA-induced expression levels of both ABCA1 and ABCG1 (P<0.05; Fig. 5(e) and (f)), leading to reduced cholesterol efflux (P < 0.05; Fig. 5(g) and (h)).

Retinoic acid-induced macrophage cholesterol efflux is farnesoid X receptor and pregnane X receptor independent

To explore the potential role of other heterodimeric partners of RXR, expression levels of LXRB, FXR and PXR were examined.



9-cis-RA

Fig. 3. Effects of 9-cis-retinoic acid (9-cis-RA) on NBD-cholesterol efflux from macrophages. J774A.1 macrophages were treated with vehicle or 9-cis-RA (0·1, 1 and 10 μmol/l) for 24 h, before equilibration with NBD-cholesterol for another 6 h. Fluorescence in the cells and the medium was detected. Efflux of NBD-cholesterol to (a) apoAl and (b) HDL. Values are means, with their standard deviations represented by vertical bars. Mean value was significantly different from that of the control group: * P<0.05, ** P<0.01.



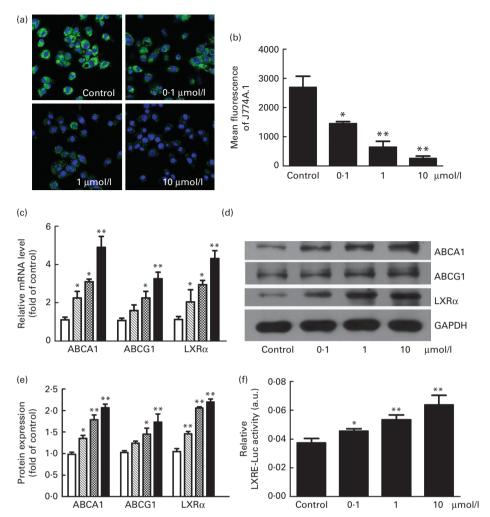


Fig. 4. Effects of 9-cis-retinoic acid (9-cis-RA) on the related gene expression levels of cholesterol transporters. J774A.1 cells were treated with vehicle or 9-cis-RA (0·1, 1 and 10 μmol/l) for 24 h before analysis. (a) Overlap images of BODIPY®493/503 and DAPI (4',6-diamidino-2-phenylindole). Magnification 400×. (b) Mean fluorescence was calculated using ImageJ software. (c) The mRNA levels of ATP-binding cassette transporter A1 (ABCA1), ATP-binding cassette transporter G1 (ABCG1) and liver X receptor-α (LXRα) were determined by quantitative real-time PCR. (d) Protein expression levels of ABCA1, ABCG1 and LXRα. (e) Quantification of ABCA1, ABCG1 and LXRα protein density normalised to the expression level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH), and expressed as optical density relative to that of the control. (f) Liver X receptor response element (LXRE)-Luc activity in 293T cells. Values are means, with their standard deviations represented by vertical bars. Mean value was significantly different from that of the control group: *P<0.05, **P<0.01. □, Control; □, 0.1 μmol/l; ■, 10 μmol/l. a.u., Arbitrary units. (A colour version of this figure can be found online at http://www.journals.cambridge.org/bjn).

Incubation with 9-cis-RA did not induce either the mRNA levels (Fig. 6(a)) or protein expression levels (Fig. 6(b) and (c)). Furthermore, the transcriptional activity of PXR was not influenced by 9-cis-RA (Fig. 6(d)). Although the transcriptional activity of FXR was stimulated by 9-cis-RA (P<0.05; Fig. 6(d)), inhibition of FXR by a specific inhibitor exerted no obvious effect on the stimulation of ABCA1 and ABCG1 expression (P<0.05; Fig. 6(e) and (f)).

Discussion

In the present study, we have presented a novel finding of 9-cis-RA and its underlying mechanism for suppressing macrophage foam cell formation and atherosclerosis. First, we showed that 9-cis-RA prevented the acceleration of atherosclerosis in HFD-fed apoE^{-/-} mice. MPM isolated from the RA group had higher expression levels of ABCA1/ABCG1

and lower cholesterol accumulation than those from the control group. The *in vitro* study further demonstrated that incubation with 9-cis-RA ameliorated oxidised LDL-induced lipid accumulation by dose-dependently inducing ABCA1/ABCG1 protein expression and cholesterol efflux via the up-regulation of LXR α transcriptional activity, and the effects were independent of FXR and PXR.

RA is an active metabolite of vitamin A that inhibits cell migration, regulates extrinsic coagulation, and promotes cell differentiation^(26,27). Its anti-tumour effects have been extensively studied for a long time^(28,29). Recently, studies have shown that RA limited restenosis after balloon angioplasty in a rabbit model⁽³⁰⁾, suggesting a potential role in the prevention of atherosclerotic CVD. A previous study has shown that all-*trans*-RA ameliorated high-fat diet-induced atherosclerosis in rabbits by inhibiting platelet activation and inflammation⁽³¹⁾. However, the rabbit model was criticised



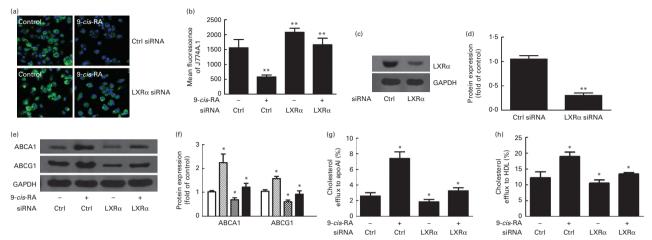


Fig. 5. Liver X receptor-α (LXRα) mediates the induction of cholesterol efflux by 9-cis-retinoic acid (9-cis-RA). After transfection with LXRα small interfering RNA (siRNA) for 48 h, 1774A 1 cells were incubated with 9-c/s-RA at a dose of 10 mmol/l for another 24 h before analysis. (a) Overlap images of BODIPY®493/503 and DAPI (4',6-diamidino-2-phenylindole). Magnification 400×. (b) Mean fluorescence was calculated using ImageJ software. (c) Knockdown efficiency of LXRα siRNA. (d) Quantification of LXRa protein density normalised to the expression level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH), and expressed as optical density relative to that of the control. (e) Protein expression levels of ATP-binding cassette transporter A1 (ABCA1) and ATP-binding cassette transporter G1 (ABCG1) were measured by Western blot. (f) Quantification of ABCA1 and ABCG1 protein density normalised to the expression level of GAPDH, and expressed as optical density relative to that of the control. 🗆 Ctrl siRNA. control: 🖫 Ctrl siRNA. 10 μmol/l: 🛤 LXRα siRNA. control: 🔳 LXRα siRNA. 10 μmol/l. (g, h) After transfection with specific siRNA, cells were equilibrated with NBD-cholesterol. Efflux of NBD-cholesterol to (g) apoAl and (h) HDL. Values are means, with their standard deviations represented by vertical bars. Mean value was significantly different from that of the control siRNA: *P<0.05, **P<0.01 (control siRNA/9-cis-RA v. LXRa siRNA/9-cis-RA). (A colour version of this figure can be found online at http://www.journals.cambridge.org/bjn).

for its use in atherosclerosis-nutritional studies because the majority of circulating cholesterol was transported in β-VLDL⁽³²⁾, which does not mimic human lipoprotein metabolism and the development of aortic lesions. Studies have also indicated that RA reduced Chlamydia pneumoniae-accelerated foam cell lesion formation in hyperlipidaemic mice, but no exact mechanism has been elucidated (33). Besides, retinoic acid-inducible gene-I (RIG-I) has been reported to be involved in atherosclerosis by regulating the differentiation of macrophages (34). Studies on retinoic acid receptor-related orphan receptor-α have also revealed that RA might be beneficial in the prevention of atherosclerosis via receptorrelated orphan receptor- $\alpha^{(35)}$. However, these two studies provided no direct link between RA and protection against

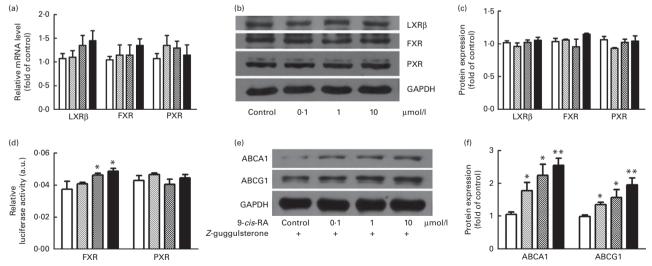


Fig. 6. Effects of 9-cis-retinoic acid (9-cis-RA) on the up-regulation of ATP-binding cassette transporter A1 (ABCA1) and ATP-binding cassette transporter G1 (ABCG1). The ABC transporters induced by 9-cis-BA were independent of farnesoid X receptor (FXB) and pregnane X receptor (PXB). (a-d) Cells were incubated with 9-cis-RA for 24h before analysis. (a) mRNA levels of LXRB, FXR and PXR were determined by quantitative real-time PCR. (b) Protein expression levels of LXRB, FXR and PXR. (c) Quantification of LXRB, FXR and PXR protein density normalised to the expression level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH), and expressed as optical density relative to that of the control. (d) Farnesoid X receptor response element/pregnane X receptor response element (FXRE/PXRE)-Luc activity in 293T cells. (e, f) J774A.1 macrophages were incubated with the combination of 9-cis-RA and Z-guggulsterone for 24 h before analysis. (e) Expression levels of ABCA1 and ABCG1. (f) Quantification of ABCA1 and ABCG1 protein density normalised to the expression level of GAPDH, and expressed as optical density relative to that of the control. Values are means, with their standard deviations represented by vertical bars. Mean value was significantly different from that of the control group: *P<0·05, **P<0·01. □, Control; ⊠, 0·1 μmol/l; ⊠, 1 μmol/l; ■, 10 μmol/l. a.u., Arbitrary units.



516



atherosclerosis. ApoE^{-/-} mice fed a Western-type diet represent a well-characterised animal model of the progression of atherosclerosis in humans (36). Using this model, we observed that 9-cis-RA significantly ameliorated atherosclerotic lesion areas in the aortic sinus compared with the control treatment. To our knowledge, this is the first in vivo study demonstrating a direct anti-atherogenic effect of 9-cis-RA.

Atherogenic lipid abnormalities, characterised by low HDL-cholesterol levels and high TAG levels together with the presence of small, dense, atherogenic LDL particles, contribute substantially to the development of atherosclerosis (37,38). In the present study, apoE^{-/-} mice treated with RA exhibited lower serum levels of total cholesterol (35.9% reduction) and LDL-cholesterol (26·1% reduction), but higher levels of HDL-cholesterol (32·1% increase) compared with those treated with vehicle (control). Elevated serum HDL-cholesterol levels were correlated with a reduced risk of atherosclerosis⁽³⁹⁾, on the basis that HDL play an important role in transporting cholesterol or cholesterol ester from peripheral tissues to the liver where cholesterol is metabolised and transformed into bile acids. The LDL receptor (Ldlr) gene, a direct target gene of LXR, has been proved to play a key role in mediating the endocytic uptake of LDL-cholesterol in the liver (40). In the present study, we showed that the 9-cis-RA treatment significantly induced the transcriptional activity of LXRE, which might induce the expression levels of hepatic LDLR and lead to a decrease in serum total and LDL-cholesterol concentrations. This pathway represents one critically important mechanism by which 9-cis-RA reduces cholesterol concentrations in both blood and peripheral tissues, thus protecting against atherosclerotic plaque formation in apo $E^{-/-}$ mice.

ABCA1 and ABCG1 are the members of the ATP-binding cassette transporter family (ABC transporters), and have been shown to regulate cellular lipid metabolism^(8–10). Studies using gene-manipulated mice have demonstrated that foam cell accumulation and atherosclerotic lesions were significantly promoted in individual transporter-deficient mice^(41,42). Therefore, we examined the regulatory role of 9-cis-RA, and showed that protein expression levels of ABCA1 and ABCG1 were higher in the MPM of apoE^{-/-} mice treated with 9-cis-RA compared with those treated with vehicle. Thus, our findings indicate that the induction of ABCA1 and ABCG1 by 9-cis-RA may contribute to the protection of atherogenesis by affecting cholesterol efflux during the transformation of foam cells. Moreover, we demonstrated that RA treatment mitigated lipid accumulation in cholesterol-preloaded macrophages, as evidenced by lower BODIPY®493/503 immunofluorescent staining in MPM from the RA group compared with the control group. Retinoids have been implicated as potent inducers of genes related to human lipid metabolism in various macrophage cell lines (43). Retinoic acid receptor activators (all-trans-RA and 4-[E-2-5,6,7,8-tetrahydro-5,5,8,8-tetramethyl-2-naphthalenyl-1-propenyl] benzoic acid (TTNPB)) were found to increase the mRNA and protein expression levels of ABCA1 in macrophages (44). However, these studies failed to specify the key role of LXR, FXR and PXR in the mediation of 9-cis-RA-induced ABCA1 and ABCG1 expression and reduced foam cell formation, a crucial early stage in atherogenesis. In accordance with previous studies that demonstrated a stimulatory role of LXR α and/or LXR β and RXR in the activation of ABCA1(45,46), our results revealed that 9-cis-RA increased the transcriptional activity of LXRE in the promoter of ABCA1/ABCG1. Consistent with the observation that in macrophages, LXR regulates cholesterol efflux by the LXRα gene rather than by $LXR\beta^{(47,48)}$, we observed no significant change in $LXR\beta$ expression at both mRNA and protein levels, and conclude that increased LXR activity was mainly attributable to LXRa. In addition, the mRNA and protein expression levels of LXRB, FXR and PXR were not obviously increased by 9-cis-RA. Although the 9-cis-RA treatment had a slight effect on the transcriptional activity of FXR, the protein expression levels of ABCA1 and ABCG1 were still increased dose-dependently when treated with a specific FXR inhibitor, indicating that FXR may not play an important role in the 9-cis-RA-induced up-regulation of ABCA1 and ABCG1 in macrophages. 9-cis-RA did not have an effect on the transcriptional activity of PXR. Collectively, these results excluded the role of FXR and PXR, another two potential heterodimeric partners of RXR modulating systematic lipid metabolism, in the 9-cis-RAinduced up-regulation of ABCA1 and ABCG1. All these data imply that the 9-cis-RA-induced up-regulation of ABCA1 and ABCG1 and cholesterol efflux are only dependent on the activation of LXR α .

Taken together, the present study is the first to reveal that 9-cis-RA significantly ameliorates HFD-induced atherosclerosis in vivo, and presents a detailed molecular mechanism by which 9-cis-RA protects against HFD-induced atherosclerosis through inducing cholesterol efflux from macrophages via the LXRα-dependent up-regulation of ABCA1 and ABCG1.

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None of the authors has any conflict of interest to declare.

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