#### **ARTICLE**



# Helper-dependent adenovirus-mediated gene transfer of a secreted LDL receptor/transferrin chimeric protein reduces aortic atherosclerosis in LDL receptor-deficient mice

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### **Abstract**

Familial hypercholesterolemia (FH) is a genetic hyperlipidemia characterized by elevated concentrations of plasma LDL cholesterol. Statins are not always effective for the treatment of FH patients; unresponsive patients have poor prognosis and rely on LDL apheresis. In the past, we developed safe and effective gene therapy strategies for the expression of antiatherogenic proteins using PEGylated helper-dependent adenoviral (HD-Ad) vectors. We recently developed a HD-Ad vector for the expression of the soluble form of the extracellular portion of the human LDL receptor (LDLR) fused with a rabbit transferrin dimer (LDLR-TF). We evaluated the efficacy of the LDLR-TF chimeric protein in CHOLDLA7, a cell line lacking LDLR expression, restoring the ability to uptake LDL. Subsequently, we administered intravenously 1 × 10E13 vp/kg of this vector in LDLR-deficient mice and observed amelioration of lipid profile and reduction of aortic atherosclerosis. Finally, we studied LDL distribution after HD-Ad vector-mediated expression of LDLR-TF in LDLR-deficient mice and found LDL accumulation in liver, and in heart and intestine. These results support the possibility of lowering LDL-C levels and reducing aortic atherosclerosis using a secreted therapeutic transgene; the present strategy potentially can be modified and adapted to non-systemic gene transfer with expression of the secreted chimeric protein in muscle or other tissues. Intramuscular or local administration strategies could improve the safety profile of this strategy and facilitate applicability.

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### Introduction

Coronary artery disease is the leading cause of death in western countries. One of the major predisposing factors is the increase of pro-atherogenic lipoproteins. The most characterized genetic hyperlipidemia is familial hypercholesterolemia (FH) [1, 2] mainly due to mutations in the low-density lipoprotein (LDL) receptor (*LDLR*) gene with consequent increase in LDL cholesterol (LDL-C) and lipid accumulation in tendons (xanthomas) and arteries (atheromas) [3]; mutations in other genes such as *APOB* and proprotein convertase subtilisin/kexin type 9 (*PCSK9*), also give rise to a similar phenotype [4, 5].

Treatment for heterozygotes and homozygotes is directed at lowering the plasma level of LDL. In heterozygotes, the most effective therapy is the administration of drugs called statins, whereas homozygotes are relatively resistant to drugs and their plasma LDL levels can be effectively lowered by physical or surgical means. Treatment in heterozygotes and homozygotes can lead to a reduced rate of

progression, and, regression of coronary atherosclerosis but is not effective in all patients, justifying the search for more effective alternatives. New drugs for FH have recently been approved: mipomersen, an antisense oligonucleotide inhibitor of apo B-100 mRNA, administered subcutaneously once weekly, has also been approved for use in the Homozygous familial hypercholesterolaemia (HoFH) population from the Food and Drug Administration [6]: lomitapide is a microsomal triglyceride transfer protein (MTP) inhibitor, binding directly to MTP in the lumen of the endoplasmic reticulum, developed as an orphan drug for treatment of HoFH, and is indicated in addition to low-fat diet and other lipid-lowering agents, with or without LDL apheresis, as an adjunctive treatment for the management of HoFH in the United States and European Union [7]. Currently available PCSK9 inhibitors, human monoclonal antibodies that exert their effect by binding to PCSK9, blocking the lysosomal degradation of LDLRs, and allowing for their repeated intracellular recycling, has been recently introduced and are able to reduce LDL-C in patients with heterozygous or receptor-defective homozygous FH by reducing LDLR degradation [8]. In contrast with the abovementioned therapeutic strategies, gene therapy can be considered as a definitive cure and not a life-long patient care, particularly for homozygous FH patients, aiming possibly at longlasting effects that lead to a definitive therapeutic solution. Significant progresses in gene transfer technology have encouraged to develop LDLR gene therapy approaches for FH treatment [9–12]. In experimental animal models of FH, VLDL receptor overexpression following viral vector-based gene transfer is associated with long-term stable correction of hyperlipidemia, with consequent attenuation of atherosclerosis progression and in certain cases even with lesions regression [13, 14]. We also observed that helper-dependent adenoviral (HD-Ad)-mediated human apolipoprotein A-I (hApoA-I) overexpression reduces aortic atherosclerosis development in ApoE- and LDLR-deficient mice [15, 16].

More recently, an AAV8-derived vector expressing LDLR has been successfully evaluated in the mouse and rhesus monkey FH models [17]. However, albeit at a much lower levels compared to Ad vectors, high concentrations of AAV vectors induce a certain degree of systemic toxicity [18]. On the other hand, both in animal models and in humans, intramuscular (IM) delivery of viral vectors has always resulted in mild local inflammation and toxicity [19].

We therefore decided to evaluate the efficacy of an alternative therapeutic strategy for FH treatment based on a chimeric secreted protein expressed after HD-Ad vector cell transduction; the advantage of this approach compared to LDLR gene transfer consists in the theoretical possibility of the expression of the therapeutic transgene in alternative tissues, such as muscle, to reduce treatment toxicity and to

translate this approach into human application. The secreted chimeric protein specifically binds LDL and is internalized by several mammalian cells, including hepatocytes, with a receptor-mediated endocytosis mechanism functionally similar to that involved in LDL metabolism. Razzini et al., in fact, generated this fusion protein that contains the signal peptide, the LDL-binding domain, the epidermal growth factor (EGF) homology domain, and the O-linked sugar domain of the human LDLR, lacking the membrane anchor as well as the cytoplasmic domain; the LDLR portion is fused with a dimer of rabbit transferrin [20, 21]. Transferrin receptor is also known to be recycled, therefore a mechanism similar to what observed with LDLR should occur [22].

Using a non-viral approach, they demonstrated that this fusion protein binds LDL restoring uptake in LDLRdeficient Watanabe heritable hyperlipidemic rabbit fibroblast in vitro [20]. The chimeric protein induces endocytosis of LDL bound to the extracellular LDLR portion through the interaction of the transferrin dimer portion with the transferrin receptors (TfRs) [20]. In vivo hydrodynamic administration of a plasmid for the expression of the LDLR-TF fusion protein was able to induce liver uptake of labeled LDL in LDLR-deficient mice [20]. In this case, plasmid administration induced only a transient effect in LDLRdeficient mice. In fact, hydrodynamic administration of unmodified plasmids allows low levels of hepatocyte transduction with short duration of expression; therefore, modifications of LDL-C levels and aortic atherosclerosis were not observed with this approach [20]. It was nonetheless relevant that most of the labeled LDL were uptaken by the liver, with limited lipid accumulation in other sites [20].

We have developed a HD-Ad vector for the expression of the above-mentioned chimeric LDLR-TF protein to achieve safe and long-term transgene expression; in the present work, we evaluated the effect of HD-Ad-mediated expression of LDLR-TF in LDLR-deficient mice. We observed amelioration of lipid profiles and reduction of aortic atherosclerosis in treated mice supporting possible developments of this strategy for future clinical applications.

### Materials and methods

### Development of a HD-Ad vector for LDLR-TF expression

We generated a cassette for liver-specific expression of the LDLR-TF fusion protein. At this aim, we used the mouse transthyretin (TTR) promoter to drive LDLR-TF expression. The expression cassette was introduced into the pLPBL1 plasmid and further subcloned into the p $\Delta$ 28 backbone plasmid to generate the HD-AdTTR-LDLR-TF as described [15]. Rescue and amplification of the vector were performed using the Ad-NG163R-2 helper virus as described elsewhere [23–25]. Briefly, a 60-mm dish of 116 cells at 80% confluence was transfected with Lipofectamine 2000 with 5 µg of PmeI-digested parental plasmid. Next day, cells were infected with Ad-NG163R-2 at a multiplicity of infection (MOI) of 1,000 vp/cell. The vector was amplified by serial coinfections of 60-mm dishes of 116 cells at 90% confluence with 10% of the crude lysate from the previous passage and Ad-NG163R-2 at an MOI of 200 vp/cell. After three passages (P) in 60-mm dishes (P1, P2, and P3), one 150-mm dish of 116 cells at 90% confluence was co-infected with 10% of the crude serial P3 lysate and Ad-NG163R-2 at MOI of 200 vp/ cell for P4. Large-scale HD-Ad production was performed in a 3 liter-culture of 116 cells (3-4 × 10E5 cells/ml) in spinner flask co-infected with all crude lysate from the 150-mm dish of serial P4 and Ad-NG163R-2 at a 200 vp/cell MOI; coinfected cells were harvested and resuspended in TM solution (10 mM Tris-HCl and 2 mM MgCl<sub>2</sub>, pH 8.0) 48 hours after infection. The harvested cells were lysed with three freezethaw cycles and incubated in 2 M MgCl<sub>2</sub> and DNase I for 1 h at 37 °C. Subsequently, cellular debris were spun down and lysate was subjected to ultracentrifugation, as described elsewhere [14]. Vector concentration was measured as virus particle number determining absorbance at 260 nm. Helper virus contamination and vector characterization were performed as previously described [26].

### Cell cultures and transfections

The 116 cells (provided by Dr. Ng from Baylor College of Medicine, Houston, Texas) were cultured in minimum essential medium containing 10% fetal bovine serum, 50 U penicillin/50 µg streptomycin in a humidified incubator at 37 °C with 5% CO<sub>2</sub>. Cells were seeded in 60-mm dishes and transfected the following day with Lipofectamine 2000 transfection reagent (Life Technologies, USA), according to the manufacturer's instructions.

### Fluorescent LDL incorporation assay

Fluorescently labeled LDL (DiI-LDL, Molecular Probe by Life Technologies, USA) are added to CHOldlA7 cells (gift from Dr. Krieger) [27], a cell line deficient for the LDLR, at a 10 µg/ml concentration for 5 hours, after infection with either HD-AdTTR-LDLR-TF vector (10 vp/cell) or with mock vector (10 vp/cell). Complementation of CHOldlA7 cell defect was evaluated by assessing incorporation of fluorescently labeled LDL (Dil-LDL) with intracellular fluorescence with a confocal microscopy, LSM (Laser Scanning Microscope) 510 Meta (Zeiss).

### Immunofluorescence staining

293 cells were washed with phosphate-buffered saline (PBS), fixed using 4% paraformaldehyde for 15 min at room temperature, and washed again with PBS. Cells were permeabilized using 0.2% Triton X-100 in PBS for 15 min at room temperature, followed by washing the samples with PBS. Samples were incubated for 1 hour at room temperature in blocking buffer (PBS with 10% bovine serum albumin). Anti-transferrin antibody (Abcam ab9033, England) was used as primary antibody and incubated in blocking for 2 hours at room temperature; after washing with PBS cells were incubated for 45 min at room temperature with an anti-rabbit secondary antibody diluted in blocking buffer [21].

### Western blot analysis

Two microliters of plasma were combined with Laemmli buffer and electrophoresed on a 6.5% PAGE. Samples were then transferred to nitrocellulose filters and subjected to western blotting analysis. We used a primary anti-rabbit transferrin ab (Abcam ab9033, England) diluted 1:500 in blocking buffer; as secondary antibody, we used a horseradish peroxidase (HRP)-conjugated rabbit anti-goat IgGs (Abcam ab97100, England) diluted 1:10,000 in blocking buffer. Antigen-antibody complexes were revealed with an enhanced chemiluminescent substrates for detection of HRP enzyme activity (Thermo Scientific Pierce), according to the manufacturer's instructions [21].

#### **Animal studies**

Animal studies were performed in accordance with National Institutes of Health guidelines and in accordance with ethical and safety rules and guidelines for the use in biomedical research provided by the relevant Italian laws and European Union's directives (no.86/609/EC). All efforts were made to minimize the animal's suffering.

Food and water were provided ad libitum. For all the experiments, we used 8-week-old female LDLR-deficient mice in C57BL/6 background (Jackson Laboratory). Animals were placed under an infrared lamp for 6 min, in order to induce tail vein vasodilation. A unit of  $1 \times 10E13$  vp/kg of HD-Ad vector was diluted in sterile PBS, prewarmed at 37 °C, and injected into tail vein as previously described [28]. Injections were performed in a total volume of 200  $\mu$ l. Control mice were treated with the same volume of PBS. Blood samples (200  $\mu$ l) were withdrawn at 0, 1, 2, 4, 8, and 12 weeks from the retro-orbital plexus, collected in a 1.5 ml tube containing 20  $\mu$ l of 50 mM EDTA (pH 8.0) and centrifuged at low speed to obtain

plasma. Plasma lipid profile (tryglicerides: TG; total cholesterol: TC; HDL-cholesterol: HDL-C; and LDL-C) was determined by dry chemistry technology on a Vitros 250 Analyzer (Ortho Clinical Diagnostics, Johnson & Johnson Co, Rochester, NY). For quality control, we used two commercial control sera (Ortho Clinical Diagnostics) to detect potential inaccuracy of the methods on both low and high analytes concentrations (low concentration control intra assay standard deviation range for the tested analytes: 1.56-4.33; high concentration control intra assay standard deviation range: 2.35-6.30; data from control assay sheets). During Fluorescence molecular tomography (FMT) studies, animals were maintained in general anesthesia at 37 °C body temperature. Anesthesia was initially delivered in an induction chamber saturated with 5% isoflurane (Iso-Vet 1000 mg/g Inhalation Vapor, Piramal Healthcare UK Ltd., Northumberland, UK) in oxygen (21/min) and subsequently maintained during all procedures with 2.5% isoflurane in oxygen (21/min).

### Quantification of aortic atherosclerotic lesions

After sacrifice, aortas were dissected from heart to iliac branching, with particular attention at removing external fat in order to stain exclusively sub-intimal aortic fat [14]. Aortic lesion stainings were performed as previously described [29]; images were acquired using a Nikon Coolscope (Nikon), and fat deposits were quantified with NIS elements software (Nikon). For histological analysis, aortas were formalin-fixed and paraffin-embedded. Representative sections of each specimen were stained with hematoxylineosin and were examined by a pathologist to evaluate atherosclerotic lesions.

### Preparation of VivoTag 750S-labeled LDL nanoparticle

Lyophilized LDL (3,500 kDa, Sigma Aldrich) were dissolved overnight in 50 mM carbonate/bicarbonate buffer, pH 8.5, at a concentration of 1 mg/ml, with gentle rotation. One milligram of VivoTag<sup>®</sup> 750S (PerkinElmer), a near-infrared (NIR) fluorescent imaging labeling agent containing an N-hydroxysuccinimide-ester, was reconstituted with 100 µl of dimethyl sulfoxide in a dark vial. One milliliter of LDL solution was incubated with 10 µl of VivoTag® 750S solution for 1 hour without light exposure at room temperature. After incubation, the reaction mix was purified with a size exclusion chromatography using Sephadex<sup>TM</sup> G-25 Medium PD-10 Desalting (GE Healthcare). Elution was monitored measuring protein concentration spectrophotometrically at 280 and 750 nm. After purification, we obtained a LDL concentration of 0.5 mg/ml with 8.5 µM of fluorophore. The solution was stored at 4 °C in a dark vial.

### Fluorescence in vivo studies with in VivoTag 750Slabeled LDL nanoparticles

LDL biodistribution in vivo was evaluated using the FMT 4000<sup>TM</sup> fluorescence tomography in vivo imaging system (PerkinElmer, Massachusetts, USA). For whole-body imaging, anesthetized mice were placed in a prone position on the imaging cassette, adjusted to the proper depth to gently restrain each mouse, and then inserted into the heated docking system (regulated at ~37 °C). A NIR laser diode was used to transilluminate mouse bodies, with signal detection occurring via a thermoelectrically cooled chargecoupled device camera placed on the opposite side of the animal. Scanning was performed using the 750EX channel. The entire image acquisition sequence took approximately 6-8 min per mouse. The images were reconstructed using the TrueQuant software for qualitative and quantitative three-dimensional analysis of the total amount of fluorescence signal in each mouse (pmoles). Mice blood clearance, organ distribution, and in vivo imaging were investigated.

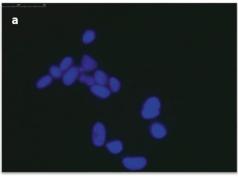
#### **Statistics**

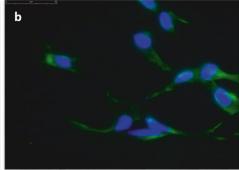
Results were analyzed using the GraphPad Prism 5 Software (Graphpad Software); p < 0.05 was considered statistically significant. Kolmogorov-Smirnov test was used to estimate normal data distribution. Intra-group differences between 0-, 1-, 2-, 4-, 8-, and 12-week time points were evaluated by analysis of variance analysis followed by Bonferroni correction. Inter-group differences between treated and untreated mice were evaluated by Student's t-test. The equality of variances was estimated by the Levene's test. All data are expressed as mean  $\pm$  SD.

#### Results

### HD-Ad vector-mediated LDLR-TF expression induces LDL uptake in a cellular FH model

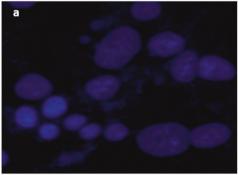
We generated a cassette for a liver-specific expression of the LDLR-TF fusion protein under the control of the mouse TTR promoter to reduce host response to the transgene. We then developed a HD-Ad vector containing this cassette for subsequent in vitro and in vivo evaluation (HD-AdTTR-LDLR-TF). We confirmed expression of the fusion protein, infecting 293 cells with either the HD-AdTTR-LDLR-TF vector (10 vp/cell) or a mock vector (10 vp/cell) as negative control and analyzed cells by immunofluorescence 48 hours after infection with an anti-TF antibody. Cells infected with HD-AdTTR-LDLR-TF showed immunoreactivity in perinuclear structures, likely the endoplasmic reticulum, as expected for a protein channeled through the secretory

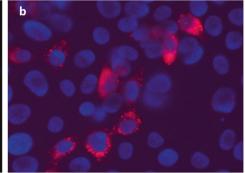




**Fig. 1** In vitro transduction with Hd-AdTTR-LDLR-TF allows expression of the chimeric protein. 293 cells were infected with a control vector (a) or with HD-AdTTR-LDLR-TF (b); immunofluorescence staining was performed with an antibody directed to

rabbit transferrin: association with perinuclear structures in cells infected with HD-AdTTR-LDLR-TF suggests that significant portion of fusion protein is contained in endocytotic vesicles localized to the perinuclear region





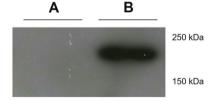
**Fig. 2** Infection with HD-AdTTR-LDLR-TF restores low-density lipoprotein (LDL) uptake in CHOldlA7 cells. CHOldlA7 cells (a cell line deficient for the LDL receptor (LDLR)) were infected with HD-AdTTR-LDLR-TF or with a control vector and supernatants were collected; LDL uptake was determined assessing the incorporation of fluorescently labeled LDL (DiI-LDL) in CHOldlA7 cells treated with

supernatant obtained from cells infected with either a control vector (a) or HD-AdTTR-LDLR-TF vector (b). DiI-LDL-positive endocytic vesicles were observed in cells treated with the LDLR-TF-containing supernatant; the incorporation of DiI-LDL was analyzed by confocal microscopy

pathway (Fig. 1). This result indeed suggests that a significant portion of fusion proteins may be contained in endocytic vesicles. We then evaluated the ability of the LDLR-TF fusion protein to restore LDL uptake in the LDLR-deficient CHOldlA7 cell line. At this purpose, we evaluated the incorporation of fluorescently labeled LDL (Dil-LDL) in CHOldlA7 cells, treated with the supernatant of cells infected with either HD-AdTTR-LDLR-TF or a control vector: as expected, CHOldlA7 cells treated with the supernatant containing the fusion protein recover the ability to uptake LDL (Fig. 2).

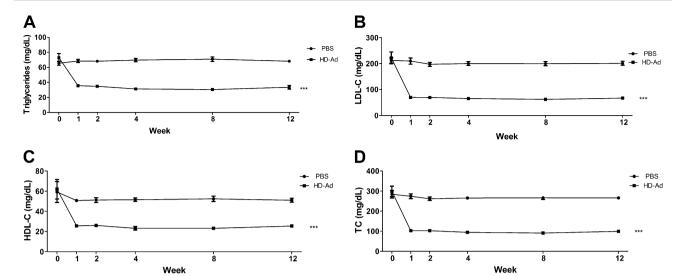
## HD-Ad vector-mediated LDLR-TF expression ameliorates lipid profile and reduces aortic atherosclerosis in LDLR-deficient mice

After assessing the ability of the LDLR-TF transgene in complementing LDLR function in vitro, we decided to evaluate its efficacy in vivo. At this aim, we intravenously



**Fig. 3** HD-AdTTR-LDLR-TF administration induced LDLR-TF expression in low-density lipoprotein receptor (LDLR)-deficient mice. Two groups of LDLR-deficient mice were injected with either  $1 \times 10E13$  vp/kg of HD-AdTTR-LDLR-TF vector or phosphate-buffered saline as control; LDLR-TF fusion protein was detected in blood sample from treated mice by western blot analysis using an anti-rabbit transferrin antibody. A band of approximately 200kD was detected in the plasma of mice treated with the HD-AdTTR-LDLR-TF vector (**b**) that was absent in plasma from control mice (**a**)

administered 1 × 10E13 vp/kg of HD-AdTTR-LDLR-TF to a group of LDLR-deficient mice; a control group was treated with PBS. We collected blood samples from both



**Fig. 4** HD-Ad-mediated LDLR-TF expression ameliorate lipid profiles in low-density lipoprotein receptor (LDLR)-deficient mice. A group of LDLR-deficient mice (n=5) was treated intravenously with  $1\times10E13$  vp/kg of HD-Ad/TTR-LDLR-TF (solid square), whereas a second group of mice (n=5) was treated with the same volume of phosphate-buffered saline (PBS) as control (solid circle). Samples were collected

before treatment and 1, 2, 4, 8, and 12 weeks after treatment. Trigly-ceride (a), total cholesterol (TC) (b), high-density lipoprotein cholesterol (HDL-C) (c), and low-density lipoprotein cholesterol (LDL-C) (d) levels were determined in both groups. TC, HDL-C, and LDL-C levels were significantly different between treated mice and the control group (\*\*\* p < 0.0001). Data are expressed as mean  $\pm$  SD

groups at different time points. We assessed the presence of the fusion protein in plasma from mice treated with HD-AdTTR-LDLR-TF by western blot using a commercially available polyclonal antibody against TF. We detected a prominent band of approximately 200 kD absent in mocktreated mice (Fig. 3); we observed an increase in molecular weight, probably due to glycosylation as previously reported [17]. To evaluate the effect of LDLR-TF expression on lipid metabolism, we determined TG, TC, LDL-C, and HDL-C levels at different time points (0, 1, 2, 4, 8, and 12 weeks) in both groups of mice. Baseline levels of plasma TG were  $72.8 \pm 5.4$  and  $65.4 \pm 2.6$  mg/dl in HD-AdTTR-LDLR-TF-treated animals and in the control group, respectively; we observed a statistically significant reduction of the levels of TG in vector-treated mice compared to controls (Fig. 4a). Basal LDL-C levels were similar in both groups (176.7 mg/dl in HD-Ad-treated mice and 211.2 mg/dl in the control group); in treated mice, LDL-C levels started to decrease 1 week after vector administration and remained significantly lower for the entire duration of the experiment (Fig. 4b). Basal levels of HDL-C were also similar in the two groups  $(62 \pm 9.7 \text{ mg/dl})$ in HD-Ad-treated mice and  $59.2 \pm 10.3$  mg/dl in the control group); however, 1 week after treatment also, HDL-C values decreased in vector-treated mice and remained lower for the entire duration of the experiment (Fig. 4c); this reduction is expected since LDLR-deficient mice have also increased LDL compared to wild-type mice. Basal levels of TC were  $298.4 \pm 26.2$  mg/dl in vector-treated

mice and  $284 \pm 18.5$  mg/dl in the control group; 1 week after administration HD-Ad-treated mice showed a significant decrease in TC levels to  $102 \pm 4.3$  mg/dl; the control group of mice did not show any significant variation of TC levels (Fig. 4d). These data suggest that HD-Ad-mediated LDLR-TF expression induces a long-lasting amelioration of lipid profile in LDLR-deficient mice. We also verified whether the treatment had an effect on the iron status of the mice, but we observed no significant differences of free iron concentration before and after treatment (data not shown).

Lastly, we evaluated whether changes in lipid metabolism affected development of atherosclerotic lesions. Twelve weeks after treatment we sacrificed both group of mice and dissected aortas from heart to iliac branching. At a macroscopic evaluation, we observed deposits of fat at the level of three branches of the aorta as expected (brachiocephalic trunk, left common carotid artery, and left subclavian artery). Fat deposits were much more evident in control mice compared to vector-treated mice (Fig. 5a). To quantify atherosclerotic lesions, we stained the aortas enface with Oil-Red-O after removal of external fat and residual tissues [14]. Lesion areas were significantly smaller in HD-AdTTR-LDLR-TF vector-treated with  $(1.78 \pm 0.48)$ mm<sup>2</sup>) compared to control group  $(5.38 \pm 0.54 \text{ mm}^2)$ Fig. 5b). Histological analysis of aortas, by means of hematoxylin/eosin staining confirmed the reduction of atherosclerotic lesions in animals treated with HD-Ad/TTR-LDLR-TF (Supplemental Fig. 1, panel A), respect to the

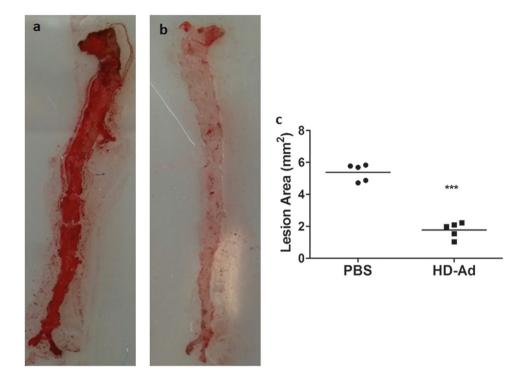


Fig. 5 HD-Ad-mediated LDLR-TF expression reduces aortic atherosclerosis development. A group of LDLR-deficient mice (n=5) were treated intravenously with  $1 \times 10E13$  vp/kg of HD-Ad/TTR-LDLR-TF-treated mice, whereas a second group of mice (n=5) was treated with the same volume of phosphate-buffered saline (PBS) as control. Twelve weeks after treatment, mice were sacrificed and their aortas were dissected. Sub-intimal fat deposits in the aortas were determined after removal of external fat and staining with Oil-Red-O. After gross

examination, we observed that aortic atherosclerosis lesion areas were smaller in mice treated with HD-Ad/TTR-LDLR-TF (b) compared to the control group (a). Total aortic atherosclerotic lesions were determined and compared in mice from both groups; HD-Ad/TTR-LDLR-TF-treated mice (solid square) showed a significantly smaller total atherosclerotic lesion area compared with control mice (solid circle, \*\*\*p < 0.01)

control group (Supplemental Fig. 1, panel B). These results strongly suggest that Hd-AdTTR-LDLR-TF administration leads not only to persistent transgene expression and amelioration of lipid metabolism, but also to a significant reduction in aortic atherosclerosis formation.

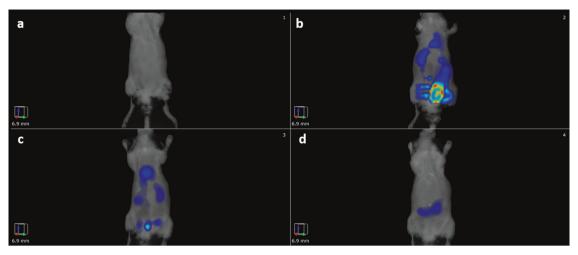
### LDL biodistribution after Hd-AdTTR-LDLR-TF vector administration in LDLR-deficient mice

Razzini et al. previously showed that LDLR-TF expression after hydrodynamic injection of an expression plasmid induced LDL accumulation in several tissues [20]. They also suggested that the prominent observed thyroid signals may be due to Iodine-labeling of LDL. We decided to characterize tissue LDL biodistribution in LDLR-deficient mice treated with the Hd-AdTTR-LDLR-TF vector and therefore expressing the LDLR-TF transgene; we also treated a control group of LDLR-deficient mice with PBS as control. Two weeks after treatment, we administered VivoTag-S 750-labeled LDL nanoparticles to both groups of mice and evaluated fluorescent signals. In mice treated with the LDLR-TF-expressing vector, no signal was present at time 0 (Fig. 6a); 2 hours after fluorescent LDL

administration we observed signals in heart, liver, and intestine (Fig. 6b). After 6 hours, the fluorescent signal was still present in the heart, albeit with a lower intensity; a signal appeared in the kidney probably due to the excretion of free dye (Fig. 6c). Signals disappeared 24 h after fluorescent LDL administration (Fig. 6d). In the control group of mice, we observed a single fluorescence signal in the bladder, probably due to free dye excretion (data not shown).

### **Discussion**

Over the last two decades, several liver-directed gene therapy strategies have been proposed for LDLR deficiency treatment [13–16]. After unsuccessful attempts with ex vivo gene transfer approaches [30, 31], several in vivo gene transfer strategies showed improved efficacy allowing the transduction of a large number of hepatocytes [32]. HD-Ad vector-mediated gene transfer was also successful for this purpose in mice models, showing improved safety and efficacy compared to FG-Ad vectors [33]. After administration of HD-Ad vectors, we and others observed



**Fig. 6** Low-density lipoprotein (LDL) biodistribution in vivo after HD-Ad-mediated LDLR-TF expression. Fluorescence images in coronal view of LDL receptor (LDLR)-deficient mice treated with 1 × 10E13 vp/kg of HD-Ad/TTR-LDLR-TF at time 0 (a); 2 h after labeled-

LDL intravenous injection we observed a FMT signal in liver, heart, and intestine (b); 6 h after injection, we observed a low FMT signal in heart and kidney (c); 24 h after injection, only a signal in the bladder was present, probably due to free dye excretion (d)

extremely prolonged high levels of transgene expression, obtaining life-long disease correction in a number of animal models [13–15]. We have already successfully ameliorated lipid profile and reduced aortic atherosclerosis expressing VLDL receptor in LDLR-deficient mice [15] and hApoA-I in both LDLR- and ApoE-deficient mice [13]. More recently, Greig et al. demonstrated phenotype correction in FH mouse and rhesus monkey models using an AAV8-based vector [17, 18].

AAV8-derived vectors demonstrated an improved hepatocyte transduction efficiency compared to other AAV serotypes; however, levels of transduction that may lead to clinical improvement are obtained only with administration of very high doses that can cause systemic toxicity [34]. Recently Hordeaux et al. reported the potential for serious acute toxicity in non-human primates after systemic administration of high dose of AAV characterized by thrombocytopenia, hyperbilirubinemia, and increased transaminases [35]. On the other hand, delivery to alternative tissues, such as muscle, may increase the therapeutic window and may lead to safer clinical applications as observed for Glybera, approved for familial hypertriglyceridemia [19]. In order to develop a system that may ultimately be administered in muscle or other tissues, we decided to evaluate an alternative strategy for the correction of LDLR deficiency using a chimeric secreted protein previously engineered by Parise et al. [20]. The LDLR-TF fusion protein contains the signal peptide, the LDL-binding domain, the EGF homology domain, and the O-linked sugar domain of the human LDLR fused with the coding sequence of a rabbit transferrin dimer [21]. This protein induces endocytosis of LDL, through binding of the LDLR-TF fusion protein to the, in fibroblasts obtained from the LDLR-deficient Watanabe rabbits [21]. Expression of LDLR-TF after hydrodynamic administration of an expression plasmid restored uptake of labeled LDL in liver and other tissues in LDLR-deficient mice [20]. However, the low plasma levels of LDLR-TF obtained after hydrodynamic plasmid administration did not affect neither lipid profile nor atherosclerosis development. This is expected since alteration of lipid metabolism with gene delivery is significantly hampered by the low bioactivity of apolipoproteins and their receptors; in fact, several molecules, such as erythropoietin, induce phenotypic effects at a picograms/ milliliter serum levels range, whereas gene transfermediated expression of apolipoproteins affects lipid profile only when their serum concentration is at least 0.1-1 mg/ml. Therefore, gene transfer-mediated LDLR expression for phenotype correction in LDLR mice is effective only when a large percentage of the hepatocytes are corrected [36]; in fact, ex vivo retrovirus-mediated hepatocyte correction and infusion did not lead to clinical improvement in patients for the low number of transduced hepatocytes able to engraft the liver [37, 38].

In the present work, we expressed the LDLR-TF fusion protein using a HD-Ad vector to efficiently transduce a large number of hepatocytes and, therefore, reach significant plasma levels able to affect lipid profile and atherosclerotic lesions in treated LDLR-deficient mice. We generated a cassette for the expression of the LDLR-TF fusion protein under the control of the mouse TTR promoter to increase hepatocyte-specific transcription [39] and subsequently developed the HD-AdTTR-LDLR-TF vector; hepatocyte-specific transduction has been shown to increase transgene duration of expression probably reducing host response against the transgene [40, 41].

HD-AdTTR-LDLR-TF allowed expression of a functional protein with perinuclear localization suggesting the implication in the secretory pathway. Expression of LDLR-TF restored DiI-LDL uptake in CHOldlA7 cells complementing LDLR deficiency. This in vitro preliminary evaluation demonstrated that the fusion protein is able to functionally bind LDL and that the HD-AdTTR-LDLR-TF vector allows the expression of a functional LDLR-TF fusion protein. We then evaluated the effect of HD-Ad-LDLR-TF administration on the lipid profile in LDLRdeficient mice. Although a lowering of the HDL was also evident, there was a considerable reduction in the TG, TC, and LDL-C levels, suggesting that the HD-AdTTR-LDLR-TF treatment led to a marked reduction in the atherogenic potential of the originally highly atherogenic plasma lipoproteins in these animals. These results demonstrate that the ability of the HD-Ad-mediated expression of the LDLR-TF fusion in inducing a long-term modifications of lipid metabolism. Moreover, 12 weeks after treatment, we observed in HD-AdTTR-LDLR-TF vector-treated mice significantly reduced atherosclerotic lesion areas compared to the control group. Since TfRs are expressed in several tissues, this result was not obvious.

Finally, we evaluated biodistribution of fluorescently labeled LDL in vivo after HD-AdTTR-LDLR-TF vector administration and subsequent LDLR-TF chimeric protein expression in LDLR-deficient mice. A similar study had been previously performed by Razzini et al. [20] using iodine-labeled LDL: in this experiment, radioactivity was predominantly taken up by liver and bladder; a significant signal was also observed in the thyroid, probably due to free radioactive iodine uptake that was completely absent in our study probably why our in vivo fluorescence determination, is adequately sensitive and avoid artifacts due to the accumulation of free radioactive iodine in specific organs, such as thyroid. We observed strong signals in liver, but also in heart, intestine, and kidney. TfR1 is expressed in a number of tissues and organs with a profile similar to LDLR; TfR2, on the other hand, has a liver-restricted expression. At terminal sacrifice, we did not observe significant organ fat accumulation in vector-treated mice after gross pathology examination. However, extra-hepatic uptake of LDL represents a concern for our approach and possible toxicity due to lipid accumulation in other tissues and organs will be thoroughly evaluated in future studies.

The data generated in this work demonstrated that the HD-AdTTR-LDLR-TF vector express a functional LDLR-TF fusion protein that is able to bind LDL and induce their uptake both in LDLR-deficient cell line and in LDLR-deficient mice; in this animal model, we observed amelioration of lipid profile and more importantly, reduction of aortic atherosclerosis lesions. To potentiate receptor recycling we are addressing our future studies to block

PSCK9 binding with the extracellular portion our chimeric protein [42].

On the other hand, in order to further improve persistence of transgene expression and reduce toxicity, additional strategies, such as vector PEGylation [43, 44] needs to be explored before suggesting a possible clinical application. However, the major advantage of this strategy is due to the expression of a secreted therapeutic transgene; LDLR-TF chimeric protein can, in fact, be produced by non-hepatic tissues, such as muscle, that can be easily transduced by HD-Ad vectors [45]. The IM delivery route can dramatically improve the safety profile and increase the therapeutic window leading to a safer clinical application.

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### Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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