ORIGINAL ARTICLE
Beneficial effects of selective HDL-raising gene transfer on survival, cardiac remodelling and cardiac function after myocardial infarction in mice

SC Gordts, I Muthuramu, E Nefyodova, F Jacobs, E Van Craeyveld and B De Geest

INTRODUCTION
High-density lipoprotein (HDL) cholesterol levels are inversely correlated with the incidence of ischaemic cardiovascular diseases. This inverse relationship is attributed to the role of HDL in reverse cholesterol transport, but may also be related to its antioxidative, anti-inflammatory, anti-apoptotic and endothelial protective features. These pleiotropic properties offer an interesting perspective to evaluate the therapeutic potential of HDL outside the classical field of atherosclerosis. Apolipoprotein A-I is the main apo of HDL and apo A-I levels strongly correlate with HDL cholesterol concentrations. Human apo A-I gene transfer has been shown to selectively increase HDL.

Post-infarct ejection fraction is lower in patients with low HDL cholesterol levels. The Framingham Heart Study demonstrated an enduring relation between low HDL cholesterol and heart failure incidence after exclusion of baseline coronary heart disease and accounting for interim myocardial infarction (MI). Moreover, low HDL cholesterol levels and low levels of apo A-I indicate an unfavourable prognosis in patients with heart failure, independent of the aetiology.

Direct cellular effects of HDL have been demonstrated in isolated cardiomyocytes in vitro, as evidenced by increased phosphorylation of extracellular signal-regulated kinases 1/2, of the transcription factor signal transducer and activator of transcription 3 and of the pro-survival kinase Akt. Furthermore, Van Linthout et al. have previously shown that HDL inhibits cardiomyocyte apoptosis under conditions of hyperglycaemia. Finally, the development of diabetic cardiomyopathy is inhibited by human apo A-I gene transfer in rats.

Taken together, epidemiological and experimental studies provide converging lines of evidence that HDL may exert direct protective effects on the myocardium. Whereas no selective HDL-raising drugs are currently available in the clinic, hepatocyte-directed human apo A-I gene transfer is an experimental HDL-targeted intervention. The goal of this study is, therefore, to investigate whether selective HDL-raising human apo A-I gene transfer exerts beneficial effects on cardiac remodelling and on cardiac function following MI. We show that human apo A-I gene transfer in C57BL/6 low-density lipoprotein receptor (LDLr)-deficient mice increases survival, decreases infarct expansion, attenuates left ventricular (LV) dilatation and improves cardiac function, following permanent ligation of the left anterior descending coronary artery (LAD).

RESULTS
Human apo A-I gene transfer selectively increases HDL in C57BL/6 LDLr−/− mice. C57BL/6 LDLr−/− mice were fed a diet containing 0.2% (w/w) cholesterol and 10% (v/v) coconut oil starting at the age of 12 weeks. Three weeks after the start of the diet, the mice were treated with the E1E3E4-deleted adenoviral human apo A-I-expressing vector AdA-I. Control mice were injected with the control vector Adnull or with saline buffer. Neither Adnull gene
transfer nor saline injection was shown to alter cholesterol levels compared with pre-injection values, and the data of both control groups were pooled. Gene transfer with the E1E3E4-deleted human apo A-I-expressing vector AdA-I induced human apo A-I levels of 220 ± 27 mg dl⁻¹ (n = 8) and 208 ± 36 mg dl⁻¹ (n = 8) on days 10 and 42, respectively, after injection. This resulted in a 1.5-fold (P < 0.0001) increase of HDL cholesterol on days 10 and 42 after injection, compared with controls (Table 1). No significant alteration of very-low-density lipoprotein cholesterol, intermediate-density lipoprotein cholesterol and LDL cholesterol was observed after AdA-I transfer, compared with the control mice (Table 1).

Increased HDL following AdA-I transfer improves survival after MI in C57BL/6 LDLr⁻/⁻ mice
MI was induced 2 weeks after gene transfer or saline injection by permanent ligation of the LAD. Comparison of Kaplan–Meier survival curves (Figure 1) showed a lower mortality rate in AdA-I-treated MI mice compared with the control MI mice (P < 0.01) (hazard ratio for mortality 0.42, 95% confidence interval 0.24–0.76). Autopsy data revealed that ventricular rupture was the cause of death in all but one mouse, indicating that AdA-I transfer significantly reduced the occurrence of ventricular rupture. To test whether the enhanced survival in AdA-I MI mice may be related to immunomodulatory effects of HDL and to accelerated infarct scar development, interleukin (IL)10, IL1β, transforming growth factor β1, collagen type I, and collagen type III mRNA levels were quantified in the infarct zone on day 3. Data in Supplementary Table 1 indicate that AdA-I gene transfer resulted in immunomodulatory and profibrogenic effects in the infarct zone.

Human apo A-I gene transfer reduces infarct expansion
Infarct area and total area at risk were determined at 24 h after ligation of the LAD in control mice and AdA-I-treated mice to exclude a potential difference in myocardial necrosis (Table 2). The total area at risk and the infarct area were not significantly different at 24 h after ligation of the LAD in control and AdA-I-treated mice. In addition, the LV cavity area and the LV wall area were similar at 24 h in both groups (data not shown).

The time course of infarct length and infarct thickness is illustrated in Figures 2a and b, respectively. Figure 2a shows that a significant increase of infarct length occurred in both the control MI mice and the AdA-I MI mice between days 3 and 14. However, infarct length on day 28 was 24% (P < 0.05) shorter in AdA-I MI mice than in control MI mice. Attenuated infarct expansion in AdA-I MI mice was also evidenced by a 41% (P < 0.0001) increase in the infarct thickness (Figure 2b) on day 28 compared with the control MI mice. Taken together, these results show that AdA-I transfer profoundly mitigates infarct expansion. The delayed divergence of both the infarct length and infarct thickness curves may be explained by the observed difference in mortality in the first 2 weeks (Figure 1), which eliminates MI mice with the most pronounced infarct expansion and consequent cardiac rupture.

Improved infarct healing at the histological level was evidenced by an increase of collagen content on day 28 in the infarct zone of AdA-I MI mice (40.8 ± 1.6%) compared with the control MI mice (37.0 ± 1.5%) (P = 0.057). Furthermore, capillary density was 36% (P < 0.05) higher on day 28 in the infarct zone of AdA-I MI mice than in the control MI mice (Table 3). To evaluate whether enhanced endothelial progenitor cell (EPC) incorporation contributed to increased capillary density in the infarct zone of AdA-I MI mice, transplantations with bone marrow of C57BL/6 β-actin green-fluorescent protein (GFP) mice were performed 4 weeks before the start of the diet. The peripheral blood count in mice 6 weeks after bone marrow transplantation was similar compared with mice without bone marrow transplantation (data not shown). The density of GFP-positive capillaries was not significantly different between chimeric control MI mice and AdA-I MI mice (Table 3), indicating a similar degree of incorporation of bone marrow-derived EPCs.

Increased HDL following AdA-I transfer attenuates LV dilatation after MI
LV cavity area and LV muscle area were determined on day 0 before MI, and on days 3, 14 and 28 after MI (Figure 3). The LV cavity area increased 3.8-fold (P < 0.001) between days 0 and 28 in control MI mice and was 38% (P < 0.05) larger compared with AdA-I MI mice on day 28 (Figure 3a). The delayed divergence of

### Table 1. Lipoprotein cholesterol levels on day 10 and day 42 after saline injection or gene transfer in C57BL/6 LDLr⁻/⁻ mice

|                      | Controls (n = 8) | Day 10 | Day 42 | AdA-I (n = 8) | Day 10 | Day 42 |
|----------------------|-----------------|--------|--------|--------------|--------|--------|
| VLDL cholesterol     |                 |        |        |              |        |        |
| (mg dl⁻¹)            | 805 ± 3.2       | 784 ± 4.3 |        | 76.6 ± 3.5   | 762 ± 5.1 |
| LDL cholesterol      | 157 ± 12        | 161 ± 10 |        | 156 ± 11     | 162 ± 10 |
| HDL cholesterol      | 72.3 ± 3.5      | 69.7 ± 4.1 |        | 108 ± 4     | 106 ± 4  |
| Non-HDL cholesterol  | 403 ± 16        | 409 ± 11 |        | 392 ± 11     | 397 ± 13 |
| Total cholesterol    | 476 ± 13        | 478 ± 8  |        | 500 ± 13     | 503 ± 12 |

Abbreviations: HDL, high-density lipoprotein; IDL, intermediate-density lipoprotein; LDL, low-density lipoprotein; LDLr, low-density lipopr otein receptor; VLDL, very-low-density lipoprotein. Control mice were injected with Adnull (n = 4) or saline (n = 4). Data are expressed as means ± s.e.m. **P < 0.0001 versus controls.
both curves should be interpreted in light of the survival bias. LV muscle area was 44% ($P < 0.01$) and 49% ($P < 0.01$) lower on day 3, compared with day 0 in control MI and AdA-I MI mice, respectively (Figure 3b). This decrease is consistent with data on infarct area on day 1 (Table 2). The subsequent increase of LV muscle area between days 3 and 28 (Figure 3b), consistent with LV hypertrophy, was not significantly different between both groups. Cardiomyocyte hypertrophy was evidenced by a 96% ($P < 0.001$) and 81% ($P < 0.001$) increase of cardiomyocyte cross-sectional area between days 0 and 28 in control MI and AdA-I MI mice, respectively (Figure 3c). This was associated with a reduction of capillary density on day 28 by 37% ($P < 0.001$) and by 23% ($P < 0.001$) in control MI and AdA-I MI mice, respectively, compared with day 0 (Figure 3d).

Representative laminin stainings of the myocardium illustrating development of cardiomyocyte hypertrophy in control MI mice are illustrated in Figure 4.

Data on interstitial fibrosis on day 28 after MI are summarised in Supplementary Table 2. Immature collagen was 28% ($P < 0.05$) lower in AdA-I MI mice compared with the control MI mice, whereas no significant difference in the amount of mature collagen in the interstitium was observed.

Human apo A-I gene transfer improves cardiac function after MI

Hemodynamic parameters in the left ventricle and in the aorta on day 28 in control MI mice and AdA-I MI mice in comparison with reference values in sham C57BL/6 LDLr$^{-/-}$ mice are shown in Table 4. The LV peak systolic pressure was significantly higher in AdA-I MI mice than in control MI mice ($P < 0.05$). The mean peripheral blood pressure ($P < 0.01$), systolic peripheral blood pressure ($P < 0.01$) and diastolic peripheral blood pressure ($P < 0.05$) were significantly higher in AdA-I MI mice compared with the control MI mice. Compared to reference control mice, blood pressure was preserved in AdA-I MI mice, but was significantly lower in control MI mice (Table 4). The increased peak rate of isovolumetric contraction (dP/dt max; $P < 0.01$) in AdA-I MI mice compared with control MI mice points to an improvement of systolic function. The higher peak rate of isovolumetric relaxation (dP/dt min; $P < 0.01$) and the lower time constant of LV relaxation ($t$; $P < 0.05$) are indicative of an improved diastolic function in AdA-I MI mice compared with control MI mice. Taken together, AdA-I gene transfer beneficially affected systolic and diastolic cardiac function in AdA-I MI mice.

Table 2. Infarct area and area at risk 24 h after permanent ligation of the LAD

|                  | Control MI | AdA-I MI |
|------------------|------------|----------|
|                  | n = 6      | n = 5      |
| Total area at risk (％of left ventricular wall area) | 49.0 ± 2.2 | 54.1 ± 3.9 |
| Infarct area (％of left ventricular wall area) | 48.4 ± 2.2 | 53.0 ± 3.7 |
| Non-infarcted area at risk (％of left ventricular wall area) | 0.600 ± 0.200 | 1.09 ± 0.37 |
| Infarct area (％of total area at risk) | 98.8 ± 0.5 | 98.1 ± 0.6 |

Abbreviations: LAD, left anterior descending coronary artery; MI, myocardial infarction. Data are as means ± s.e.m.

Table 3. Quantification of total capillaries and capillaries containing bone marrow-derived GFP-positive-incorporated EPCs in the infarct zone

|                                | No prior bone marrow transplantation | Prior bone marrow transplantation |
|--------------------------------|-------------------------------------|----------------------------------|
|                                | Control MI (n = 16) | AdA-I MI (n = 16) | Control MI (n = 16) | AdA-I MI (n = 14) |
| Capillary density (number mm$^{-2}$) | 145 ± 16 | 197 ± 19* | 144 ± 14 | 198 ± 21* |
| GFP-positive capillary density (number mm$^{-2}$) | N.A. | N.A. | 30.7 ± 8.0 | 35.9 ± 8.6 |
| GFP-positive capillaries (％of total capillaries) | N.A. | N.A. | 19.3 ± 4.2 | 18.0 ± 3.4 |

Abbreviations: EPC, endothelial progenitor cell; GFP, green-fluorescent protein; MI, myocardial infarction; N.A., not applicable. Data are as means ± s.e.m. $P < 0.05$ for control MI versus AdA-I MI.
function after MI, and induced a preservation of blood pressure compared with the sham control mice.

Prior irradiation and bone marrow transplantation enhance survival and attenuate LV dilatation in control MI mice, but beneficial effects of AdA-I transfer on cardiac structure and cardiac function remain. In the experiments designed to evaluate EPC incorporation, we made the startling observation that prior irradiation and bone marrow transplantation significantly reduced mortality in control MI mice ($P < 0.001$) (hazard ratio for mortality 0.23, 95% confidence interval 0.12–0.46) (Figure 5a) and in AdA-I MI mice ($P < 0.05$) (hazard ratio for mortality 0.28, 95% confidence interval 0.081–0.96) (Figure 5b) compared with the respective MI groups without prior bone marrow transplantation. Table 5 summarises infarct parameters and remodelling parameters on day 28 in control MI mice and AdA-I MI mice with and without prior bone marrow transplantation. Infarct area and infarct length were

**Figure 3.** Time course of left ventricular remodelling in control MI mice (grey circles) and AdA-I MI mice (black triangles). Longitudinal changes in left ventricular cavity area, left ventricular muscle area, cardiomyocyte cross-sectional area and capillary density are illustrated in panels (a–d), respectively. Data represent means ± s.e.m. (n is identical as indicated in the Legend to Figure 2). For reasons of clarity, only statistical differences between day 0 and other time points, or differences between control MI mice and AdA-I MI mice are indicated. $^aP < 0.05$; $^{ab}P < 0.01$; $^{abc}P < 0.001$ for longitudinal comparisons versus day 0 control MI. $^{a+}P < 0.01$; $^{ab+}P < 0.001$ for longitudinal comparisons versus day 0 AdA-I MI. $^\dagger P < 0.05$ for control MI versus AdA-I MI.

**Figure 4.** Representative laminin stainings of myocardial tissue illustrating development of cardiomyocyte hypertrophy in control MI mice. Representative photomicrographs show laminin-stained cardiomyocytes on day 0 (panel a), day 3 (panel b), day 14 (panel c) and day 28 (panel d) after MI. Scale bar represents 500 μm.
Survival analysis was performed by log-rank test. Comparison Kaplan–Meier survival curves after myocardial infarction in mice without (grey line) and with (black line) prior bone marrow transplantation during a 28-day follow-up period. Kaplan–Meier survival curves in panel (a) concern control MI mice. At the start of the study, 64 control MI mice without prior bone marrow transplantation and 17 control MI mice with prior bone marrow transplantation were included. Data in panel (b) are based on inclusion of 29 AdA-I MI mice without prior bone marrow transplantation and 15 AdA-I mice with prior bone marrow transplantation. Survival analysis was performed by log-rank test.

Table 4. Hemodynamic parameters in the left ventricle and in the aorta on day 28 after myocardial infarction in comparison with reference values in sham C57BL/6 LDLr⁻/⁻ mice

|                      | Sham (n = 19) | Control MI (n = 11) | AdA-I MI (n = 11) |
|----------------------|---------------|---------------------|-------------------|
| **Left ventricle**   |               |                     |                   |
| Peak systolic pressure (mm Hg) | 103 ± 2       | 84.9 ± 2.6[I]       | 95.9 ± 3.8[I]     |
| End-diastolic pressure (mm Hg) | 2.19 ± 0.68   | 4.81 ± 1.75         | 1.92 ± 0.92       |
| Peak dP/dt max (mm Hg ms⁻¹) | 12.1 ± 0.6    | 6.60 ± 0.27[I]      | 9.01 ± 0.54[II]   |
| Peak dP/dt min (mm Hg ms⁻¹) | − 8.7 ± 0.38  | − 5.57 ± 0.23[I]    | − 7.32 ± 0.48[II] |
| Tau (ms)             | 5.19 ± 0.29   | 6.90 ± 0.35[I]      | 5.75 ± 0.31[I]    |
| Heart rate (bpm)     | 601 ± 15      | 554 ± 23            | 573 ± 26          |
| **Aorta**            |               |                     |                   |
| Mean pressure (mm Hg) | 78.8 ± 2.1    | 64.3 ± 2.6[I]       | 78.4 ± 4.2[II]    |
| Peak systolic pressure (mm Hg) | 99.5 ± 2.0    | 81.0 ± 2.3[I]       | 96.1 ± 4.4[II]    |
| Peak diastolic pressure (mm Hg) | 60.2 ± 2.7    | 50.6 ± 3.3          | 63.2 ± 3.5        |

Abbreviations: LDLr, low-density lipoprotein receptor; MI, myocardial infarction. Data are expressed as means ± s.e.m. *P < 0.05; **P < 0.01 for comparison versus control MI. ¹P < 0.05; ²P < 0.01; ³P < 0.001 versus Sham.

Figure 5. Comparison Kaplan–Meier survival curves after myocardial infarction in mice without (grey line) and with (black line) prior bone marrow transplantation during a 28-day follow-up period. Kaplan–Meier survival curves in panel (a) concern control MI mice. At the start of the study, 64 control MI mice without prior bone marrow transplantation and 17 control MI mice with prior bone marrow transplantation were included. Data in panel (b) are based on inclusion of 29 AdA-I MI mice without prior bone marrow transplantation and 15 AdA-I mice with prior bone marrow transplantation. Survival analysis was performed by log-rank test.

Table 5. Comparison of infarct parameters and of left ventricular remodelling at day 28 after myocardial infarction in mice without and with prior irradiation and bone marrow transplantation

|                      | No prior bone marrow transplantation | Prior bone marrow transplantation |
|----------------------|-------------------------------------|----------------------------------|
|                      | Control MI (n = 16) | AdA-I MI (n = 16) | Control MI (n = 16) | AdA-I MI (n = 14) |
| Infarct area (mm²)   | 3.06 ± 0.16           | 2.84 ± 0.26            | 2.44 ± 0.17[I]     | 2.05 ± 0.17[I]   |
| Infarct length (µm)  | 9750 ± 580            | 7430 ± 800            | 5000 ± 280[I]      | 4220 ± 250[I]    |
| Infarct thickness (µm) | 270 ± 12             | 379 ± 25²             | 590 ± 24[I]        | 613 ± 17[I]      |
| Whole left ventricular area (mm²) | 263.2 ± 3.15         | 230.0 ± 0.9           | 19.2 ± 0.5[I]      | 17.5 ± 0.5[I]    |
| Left ventricular cavity area (mm²) | 16.2 ± 1.5           | 11.7 ± 1.3¹           | 7.30 ± 0.58[II]   | 5.98 ± 0.40[II]  |
| Left ventricular muscle area (mm²) | 7.08 ± 0.60          | 8.49 ± 0.65           | 9.44 ± 0.36[I]     | 9.46 ± 0.32     |
| Septal wall thickness (µm) | 1060 ± 30            | 1240 ± 60             | 1100 ± 30          | 1150 ± 40       |
| Cardiomyocyte cross-sectional area (µm²) | 405 ± 16             | 399 ± 17              | 293 ± 12[I]        | 282 ± 12[I]    |
| Capillary density (number mm⁻²) | 4140 ± 270           | 4620 ± 290            | 3380 ± 180[I]      | 3550 ± 170[I]   |

Abbreviation: MI, myocardial infarction. Data are expressed as means ± s.e.m. *P < 0.05; **P < 0.01 for control MI versus AdA-I MI. ¹P < 0.05; ²P < 0.01; ³P < 0.001; ⁴P < 0.0001 for prior bone marrow transplantation groups versus respective no prior bone marrow transplantation groups.

significantly reduced, and infarct thickness was significantly increased in control MI mice and in AdA-I MI mice with prior irradiation and bone marrow transplantation compared with the respective MI mice groups without prior bone marrow transplantation. Furthermore, pronounced reductions in LV cavity area were observed in mice with prior irradiation and bone marrow transplantation compared with mice without prior bone marrow transplantation. Selective HDL-raising gene transfer
reduced infarct length by 16% \( (P < 0.05) \) and decreased LV cavity area by 18% \( (p = 0.079) \) in mice with prior irradiation and bone marrow transplantation (Table 5), indicating the robustness of the effects of selective HDL-raising gene transfer on cardiac structure and function. Comparison of representative Sirius Red-stained cross-sections on day 28 after MI in mice without or with prior bone marrow transplantation is shown in Figure 6.

Hemodynamic parameters in the left ventricle and in the aorta on day 28 after MI in control MI mice and AdA-I MI mice with prior irradiation and bone marrow transplantation are summarised in Table 6. Similar as in mice without prior bone marrow transplantation, the peak rate of isovolumetric contraction \((\text{dP/dt max})\) was markedly higher in AdA-I MI mice compared with the control MI mice (Table 6). The higher peak rate of isovolumetric relaxation \((\text{dP/dt min})\) \( (P < 0.05) \) and the lower time constant of LV relaxation \((\tau)\) \( (P < 0.05) \) in AdA-I MI mice with prior bone marrow transplantation than in control MI mice with prior bone marrow transplantation confirm the robustness of the effects of human apo A-I gene transfer. In further agreement with hemodynamic data in Table 4, systolic peripheral blood pressure in mice with prior bone marrow transplantation was higher \((P < 0.05) \) in AdA-I MI mice than in control MI mice (Table 6).

**DISCUSSION**

The current study shows that hepatocyte-directed human apo A-I gene transfer, resulting in a selective increase of HDL, exerts direct cardiac effects, leading to improved survival, attenuated infarct expansion, reduced LV dilatation, and enhanced LV systolic and diastolic function post MI. As the objective of this study was to

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**Figure 6.** Representative Sirius Red-stained cross-sections of hearts on day 28 after MI in control and AdA-I-treated C57BL/6 LDLr −/− mice without and with prior bone marrow transplantation. Control MI and AdA-I MI mice without prior bone marrow transplantation are illustrated in panel (a) and panel (b), respectively. Cross-sections of control MI mice and AdA-I MI mice with prior bone marrow transplantation are shown in panel (c) and panel (d), respectively. Sirius Red stains fibrotic infarct tissue red and healthy tissue orange. Scale bar represents 1 mm.

**Table 6.** Hemodynamic parameters in the left ventricle and the aorta at day 28 after myocardial infarction in bone marrow transplantation C57BL/6 LDLr −/− mice

| Parameter                          | Control MI \((n = 14)\) | AdA-I MI \((n = 10)\) |
|-----------------------------------|-------------------------|-----------------------|
| **Left ventricle**                |                         |                       |
| Peak systolic pressure \((\text{mm Hg})\) | 87.9 ± 3.9              | 96.7 ± 2.4            |
| End-diastolic pressure \((\text{mm Hg})\) | 2.49 ± 1.40             | 0.565 ± 0.818         |
| \(\text{dP/dt max } (\text{mm Hg ms}^{-1})\) | 7.07 ± 0.75             | 9.90 ± 0.54*          |
| \(\text{dP/dt min } (\text{mm Hg ms}^{-1})\) | –6.17 ± 0.50            | –7.69 ± 0.39*         |
| Tau \((\text{ms})\)               | 5.84 ± 0.32             | 4.93 ± 0.17*          |
| Heart rate \((\text{bpm})\)      | 577 ± 25                | 616 ± 21              |
| **Aorta**                         |                         |                       |
| Mean pressure \((\text{mm Hg})\)  | 66.9 ± 5.0              | 75.4 ± 3.2            |
| Systolic pressure \((\text{mm Hg})\) | 86.3 ± 4.6              | 98.4 ± 1.8*           |
| Diastolic pressure \((\text{mm Hg})\) | 55.1 ± 5.4              | 59.1 ± 5.3            |

Abbreviations: LDLr, low-density lipoprotein receptor; MI, myocardial infarction. Data are expressed as means ± s.e.m. *\(P < 0.05\) versus control MI.
investigate the effect of HDL on cardiac remodelling and heart failure development, a model of permanent ligation of the LAD was used, which excludes salutary coronary effects of HDL on myocardial salvage, as observed in a model of ischaemia/reperfusion injury.14,15 As expected, the infarct area and the area at risk were nearly identical at 24 h after ligation, implying that the initial increase in loading conditions was not different between human apo A-I gene transfer MI mice and control MI mice. Infarct expansion is the disproportionate thinning and dilatation of the infarct segment, and predisposes to myocardial rupture and congestive heart failure.16–19 The decreased incidence of ventricular rupture in AdA-I MI mice, mainly occurring between days 3 and 14 post MI, can be viewed as manifestation of attenuated infarct expansion.20 Our gene expression analysis data in the infarct zone on day 3 indicate that human apo A-I gene transfer resulted in immunomodulatory effects, and in accelerated fibrogenesis that may have attenuated infarct expansion and may have reduced the incidence of ventricular rupture. Furthermore, direct effects of apo A-I on the function of monocyte-derived macrophages have been described,21 which may have contributed to reduced infarct expansion.

Survival bias should be taken into account in the comparison of the morphometric data between days 3 and 14. In contrast, as mortality between days 14 and 28 was very low in both control MI and AdA-I MI mice, the morphometric comparison between these two time points, demonstrating suppression of infarct expansion and of LV dilatation in AdA-I MI mice, is not affected by survival bias. Neovascularisation has a role in scar formation and scar tissue remodelling.22 Increased neovascularisation was observed in the infarcts of AdA-I MI mice on day 28 after gene transfer. HDL is known to exert potent effects on the endothelium via enhanced endothelial survival,23 endothelial cell migration24 and EPC-mediated repair.25,26 We did not observe enhanced incorporation of bone marrow-derived EPCs into the infarct zone. Nevertheless, increased EPC number and function following human apo A-I gene transfer27 may have improved neovascularisation in a paracrine way by releasing angiogenic factors and prostaglandins to stimulate sprouting of local vessels.28

Selective HDL-raising gene transfer significantly improved systolic and diastolic cardiac function after MI and led to a preservation of peripheral blood pressure compared with sham mice, in contrast to the lower blood pressure in control MI mice. In agreement, human apo A-I gene transfer has been shown to improve cardiac function in a rat model of diabetic cardiomyopathy.13 Furthermore, HDL has also been demonstrated to increase cardiomyocyte contractility ex vivo.15 Overall, the effect of human apo A-I gene transfer on cardiac function post MI may reflect both the effect of HDL on LV remodelling and direct potentiation of cardiomyocyte contractility by HDL.

Mortality was strongly reduced, and LV dilatation was significantly attenuated in control MI mice with prior irradiation and bone marrow transplantation compared with control MI mice without prior bone marrow transplantation. Notwithstanding this, the significantly milder phenotype, the effect of AdA-I transfer was confirmed. Infarct size at risk was shorter and infarct size was attenuated and LV function was improved in AdA-I MI mice compared with the control MI mice under conditions of prior bone marrow transplantation and irradiation. This independent set of experiments conducted in a different setting strongly corroborates our findings in animals without prior irradiation and bone marrow transplantation. The milder phenotype in bone marrow transplantation mice might be related to immune reset. CD4+ T helper cells and regulatory T cells have been demonstrated to have a role in infarct healing and myocardial remodelling.29,30 Autologous bone marrow transplantation has been shown to reverse autoimmune arthritis in mice via induction of regulatory T cells.31 In addition, clinical hematopoietic stem cell transplantation with either myeloablative or nonmyeloablative regimens has been used for treatment of patients with severe autoimmune diseases.32,33 Similarly, bone marrow transplantation in the current study may have altered T cell biology. Regulatory T cells may suppress inflammatory activity of monocytes by secreting inhibitory signals such as IL-10 and transforming growth factor-β, and through contact-mediated actions.34

In conclusion, this murine experimental study supports the hypothesis that HDL may be an important target for treatment of myocardial remodelling post MI, and for prevention and treatment of heart failure.

MATERIALS AND METHODS

Construction of E1E3E4-deleted adeno viral gene transfer vectors for hepatocyte restricted overexpression of human apo A-I

Construction of the E1E3E4-deleted adeno viral vector Ada-I has been described previously.35 This vector contains the 1.2-kb DC172 promoter, consisting of an 890-bp human α1-antitrypsin promoter and two copies of the 160-bp α1-microglobulin enhancer, upstream of the genomic human apo A-I sequence and two copies of the hepatic control region-1. The E1E3E4-deleted control vector Adnull66 does not contain an expression cassette. Vector production was performed as described previously.37

Animal experiments

All animal procedures were approved by the ethical committee for animal experimentation of the Catholic University of Leuven, Leuven, Belgium. Male C57BL/6 LDLr−/− mice, originally purchased from Jackson Laboratories (Bar Harbor, ME, USA), received a hypercholesterolemic diet containing 0.2% (w/w) cholesterol and 10% (v/v) coconut oil ad libitum starting from the age of 12 weeks. Gene transfer in C57BL/6 LDLr−/− mice was performed 3 weeks after the start of the hypercholesterolemic diet by tail vein injection of 5 × 1010 adeno viral particles of Ada-I. This dose was chosen based on a prior study on the effect of this vector on HDL levels and on atherosclerosis in C57BL/6 LDLr−/− mice.38 Control mice were injected with the same dose of the control vector Adnull or with saline. As no differences were observed between the Adnull and saline-injected mice with regard to different end points, data of both the control groups were consistently pooled. The experimental diet was maintained throughout the entire duration of the experiments.

Two weeks after gene transfer or saline injection, MI was induced by permanent ligation of the LAD, as described.39 In selected experiments, C57BL/6 LDLr−/− mice were lethally irradiated with 9.5 Gy at the age of 8 weeks. Transplantation of 8 × 106 bone marrow cells obtained from male C57BL/6 j-actin GFP-transgenic mice was performed via tail vein injection 24 h after irradiation. Feeding of the hypercholesterolemic diet was started 4 weeks after bone marrow transplantation. Gene transfer and MI were performed in chimeric mice at the age of 15 and 17 weeks, respectively.

In vivo hemodynamic measurements

Invasive hemodynamic measurements were performed 28 days after MI, following anaesthesia induced by intraperitoneal administration of 1.4 g kg−1 urethane (Sigma, Steinheim, Germany). Measurements were performed with a 1.1 French Millar pressure catheter (SPR-67/NR; Millar Instruments, Houston, TX, USA), as described before.9 Data were registered with Powerlab Bridge Amplifier and Chart Software (sampling rate 2000 Hz; Fysicon, Oss, the Netherlands).

Plasma lipoprotein and apo analysis

Mouse lipoproteins were separated by density gradient ultracentrifugation in a swing-out rotor.40 Fractions were stored at −20 °C until analysis. Total cholesterol in plasma and lipoprotein fractions was determined with commercially available enzymes (Roche Diagnostics, Basel, Switzerland). Precipitation (Roche Diagnostics) was used as a standard. Human apo A-I levels in plasma were determined by sandwich ELISA.31

Area at risk and infarct size assessment 1 day after MI

Twenty-four hours after completion of the MI protocol, 2 ml of a blue tissue-marking dye (Polysciences, Warrington, PA, USA) was injected as a bolus into the aorta until most of the heart turned blue. Hearts were arrested in diastole by CdCl2 (100 μl; 0.1 N) and flushed with saline to wash out excess blue dye. After embedding the hearts in 5% low gelling temperature agarose (Sigma), 500-μm thick cross-sections were made
using a HM 650V Vibration Microtome (MICROM International GmbH, Walldorf, Germany). The slices were then incubated in a 1.5% trypthohyl tetrazolium chloride (Alfa Aesar, Ward Hill, MA, USA) solution in an isotonic phosphate buffer (pH 7.4) for 30 min at 37 °C. Images were made with a SteREO Lumar V.12 microscope (Zeiss, Zaventem, Belgium) and the areas of infarcted tissue, the non-infarcted area at risk, and the LV wall area were determined using Image J software (Wayne Rasband, National Institutes of Health, Bethesda, MD, USA).

Tissue preparation for histological and morphometric analysis
Mice were perfused via the abdominal aorta with phosphate-buffered saline and hearts were arrested in diastole with CdCl (100 µl; 0.1% Na), followed by perfusion fixation with 1% paraformaldehyde in phosphate-buffered saline. Hearts and lungs were dissected and weighed. Hearts were post fixed overnight in 1% paraformaldehyde, embedded in paraffin and 6-µm thick cross-sections at 130-µm spaced intervals were made, extending from the apex to the basal part of the left ventricle.

Morphometric analysis of infarct expansion and LV remodelling
Infarct expansion and LV remodelling were assessed by morphometric analysis on mosaics images of Sirius Red-stained heart cross-sections using Axiovision 4.6 software (Zeiss). Longitudinal comparisons of parameters were based on data obtained on day 0 or on days 3, 14 or 28 after MI. Midline infarct length was defined as the midline length of infarct that included >50% of the whole thickness of the myocardial wall. Whole LV area (µm²), LV cavity area (µm²), LV remote muscle area (µm²; including the septum), sepal wall thickness (µm) and infarct area (µm²) were analysed. Infarct wall thickness (µm) was measured at equidistant points over the infarct area perpendicular to the infarcted wall. All geometric measurements were computed in a blinded fashion from representative tissue sections of four separate regions, and the average value was used to represent that animal for statistical purposes.

Analysis of collagen deposition
To measure collagen content in the infarct zone and in the interstitium, Sirius Red staining was performed, as previously described by Junqueira et al.4 Sirius Red polarisation microscopy on a Leica DM RBE microscope (Leica Microsystems, Wetzlar, Germany) with KS300 software (Zeiss) was used to quantitively analyse bone marrow-derived collagen fibres as orange–red birefringent, and loosely packed less crosslinked and immature collagen fibres as yellow–green birefringent. Collagen-positive area was normalised to infarct area or LV remote area and was expressed as percentage. Two midventricular sections were studied per animal.

Immunohistochemistry
Cardiomyocyte hypertrophy was analysed on paraffin sections stained with rabbit anti-mouse laminin (Sigma; 1/50) by measuring the cardiomyocyte cross-sectional area (µm²) of at least 200 randomly selected cardiomyocytes in the non-infarcted LV myocardium. Capillary density in the infarct area and in the non-infarcted myocardium was determined on CD31-stained sections using rat anti-mouse CD31 antibodies (BD Biosciences, Erembodegem, Belgium; 1/500). Two midventricular cross-sections were analysed per mouse.

To evaluate EPC incorporation in the infarct area, heart sections from the infracted LV myocardium using TRIzol reagent (Invitrogen) and the Purelink RNA Mini Kit (Invitrogen). On-column DNase treatment was performed using Purelink DNase (Invitrogen), according to the manufacturer’s protocol. Total RNA (1 µg) was reverse transcribed using the QuantiTect Reverse Transcription kit (Qiagen, Hamburg, Germany). qRT-PCR was performed on a 7500 FAST real-time PCR system (Applied Biosystems, Carlsbad, CA, USA) using the TaqMan Fast Universal PCR Master Mix (Applied Biosystems), and a premade mix containing primers and MGB probes (Taqman gene expression assay, Applied Biosystems; see Supplementary Table 3 for details) to quantify IL-1β, IL-10, transforming growth factor β1, collagen type Iα1 and collagen type IIIα1 complementary cDNA levels (n = 8 per group). The Gapdh housekeeping gene was used as endogenous control. Data analysis was performed using ∆∆CT-based fold-change calculations.

Statistical analysis
All data are expressed as means ± s.e.m. Parameters between three groups were compared by one-way analysis of variance, followed by Tukey multiple comparison post-test using GraphPad Instat (GraphPad Software, San Diego, CA, USA). When indicated, a logarithmic transformation or a square root transformation or a non-parametric test was performed. Parameters between two groups were compared using Student’s t-test. When indicated, a logarithmic transformation, a square root transformation or a non-parametric Mann–Whitney test was performed. The assumption of Gaussian distribution was tested using the method Kolmogorov and Smirnov. Kaplan–Meier survival curves were analysed by log-rank test using Prism4 (GraphPad Software). A two-sided P-value of < 0.05 was considered statistically significant.

CONFLICT OF INTEREST
The authors declare no conflict of interest.

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