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Supplemental Data Part 1: The preparation of MNCs or Placebo

As reported previously by our team and others [1-4], MNCs were prepared from the bone marrow aspirated from the posterior iliac crest under local anesthesia. A volume of approximately 80-100ml of bone marrow was harvested aseptically at the day of intracoronary infusion and treated with 10IU/ml of heparin.

Volume depletion and enrichment of bone marrow-derived MNCs were performed using apheresis (Cobe Spectra, Gambro BTC Inc., Lakewood, CO) to obtain a concentrated stem cell suspension. The bone marrow cell preparation was diluted (1:10) with acid citrate dextrose solution A to prevent coagulation. One hour prior to intracoronary infusion, the MNC suspension was centrifuged, and cell pellets were re-suspended in 10 ml heparinized saline. MNCs were examined microscopically and counted by hemocytometer. The same volume of heparinized normal saline was prepared as the placebo for intracoronary infusion.

Supplemental Data Part 2: Intracoronary infusion of autologous bone marrow MNC

Patients who underwent the primary percutaneous coronary intervention (PCI) were examined by coronary angiography and then MNCs infusion was performed. Otherwise, patients were first revascularized for the infarct related coronary artery left anterior descending (LAD) and then given by the MNC infusion following the index procedure. An over-the-wire balloon catheter (Marverick, Boston Scientific Co.) connected with pressure pump containing the MNC solutions was used for stem cell delivery. Intracoronary nitroglycerin was administered to prevent coronary spasms or abrupt closure caused by the high viscosity infusion of the cell solutions. During the infusion, the balloon was inflated for two minutes to prevent washout of MNCs. The procedure of intracoronary cell infusion was well tolerated, and did not show any severe adverse events or increases in the troponin levels. In the placebo, 10 ml heparinized saline solution was infused under the same procedure.

Supplemental Data Part3: Cardiac MRI

Cardiac MRI was performed using a 1.5-T clinical MRI scanner (Siemen Avanto, Germany) with a phase array radiofrequency receiver coil, electrocardiography gating, and respiratory triggering. Left ventricular (LV) volume and function were evaluated, including LV end-diastolic volume (EDV), end-systolic volume (ESV), and ejection fraction (LVEF). The endocardial and epicardial borders were traced in all end-diastolic and end-systolic short axis and long axis slices to determine EDV and ESV for calculation of global LVEF, with $LVEF = [(EDV - ESV) / EDV] \times 100\%$. Two experienced MRI experts blinded to all clinical data analyzed the images. All MRI was quantitatively analyzed according to the previous study of our team and others [2,3,5].

Supplemental Data Part 4: Two-dimensional echocardiography

Following the previous study protocol[6], 2DE was conducted in the left lateral decubitus position using a clinically available system (Vivid 7 and E9, General Electric-Vingmed, Horten, Norway). Data acquisition was performed with a 3.5-MHz transducer at a depth of 16 cm in the standard parasternal and apical views. During the breath hold, M-mode and two dimension (2D) images were obtained and saved in cine-loop format. Data analysis was performed offline (EchoPAC version 111.0.0; General Electric-Vingmed,). EDV, ESV, and LVEF were derived from the conventional 2D apical two and four-chamber view images, using the biplane Simpson's rule.

Supplemental Data Part 5: Analysis of myocardial perfusion and viability by SPECT and PET

According to previous studies [7-11], myocardial perfusion was assessed by a dual-head SPECT (Siemens Medical, GER) 90 min after intravenously injecting a dose of 740 MBq technetium-99m sestamibi ($^{99m}\text{Tc-MIBI}$). Thirty-two projection images per 40 s in a 64×64 matrix were achieved by 180° -rotation arc from the 45-right and anterior sector to the 45-left and anterior sector. SPECT was reconstructed with Butterworth cut-off frequency of 0.45, with an order of 5. Quantitative analysis of imaging was performed by using Cedars quantitative perfusion SPECT software (QPS). The left ventricle was divided into 17 segments and scored from 0 to 4 automatically (0 = normal perfusion, 1 = mild perfusion decrease, 2 = moderate decrease, 3 = severe decrease, 4 = perfusion defect) according to the American Heart Association/American College of Cardiology recommendations [12]. The improvement of myocardial perfusion after cell infusion was defined as >1 score perfusion improvement in the segments pre-cell infusion compared with the relative segments post-cell infusion.

PET scans were performed within 2 days after SPECT. After an overnight fasting for at least 8 hours, an oral glucose dose of 50 g was given to the patients. Insulin was intravenously administered according to the blood glucose level. Fluorodeoxyglucose (FDG, 111–185 MBq) was administered intravenously when the blood glucose level was appropriate. Images were acquired 1 hour after tracer injection using a Biograph 64 PET/CT scanner (Siemens Medical Solutions, Knoxville, TN) equipped with

high-performance LSO PET crystals and a 64-slice CT. Images were collected and reconstructed using attenuation weighted-OSEM iterative reconstruction (8 subsets, 4 iterations). A non-gated, noncontrast-enhanced CT transmission scanning (140 kV, 35 mA) was performed for attenuation correction and anatomical localization preceded the PET scan.

After reconstruction, FDG uptake in left ventricle was quantitatively analyzed by QPS software in the same workstation as perfusion analysis. Areas of scar (perfusion–metabolism match) or hibernated myocardium (perfusion–metabolism mismatch) were assessed by two experienced nuclear medicine experts blinded to the present trial. The segmental tracer activity was calculated as a percentage of uptake of the segment with maximal uptake (normalization to the 100% perfusion maximum) and categorized on a 4-point scale: 1 denoting tracer activity of >75%; 2 denoting tracer activity of 50% to 75%; 3 denoting tracer activity of 25% to 49%; and 4 denoting tracer activity of <25%. Metabolic improvement was defined as a significant fill-in (>10%) of FDG defects was observed on the post-cell infusion images compared with baseline [13-15].

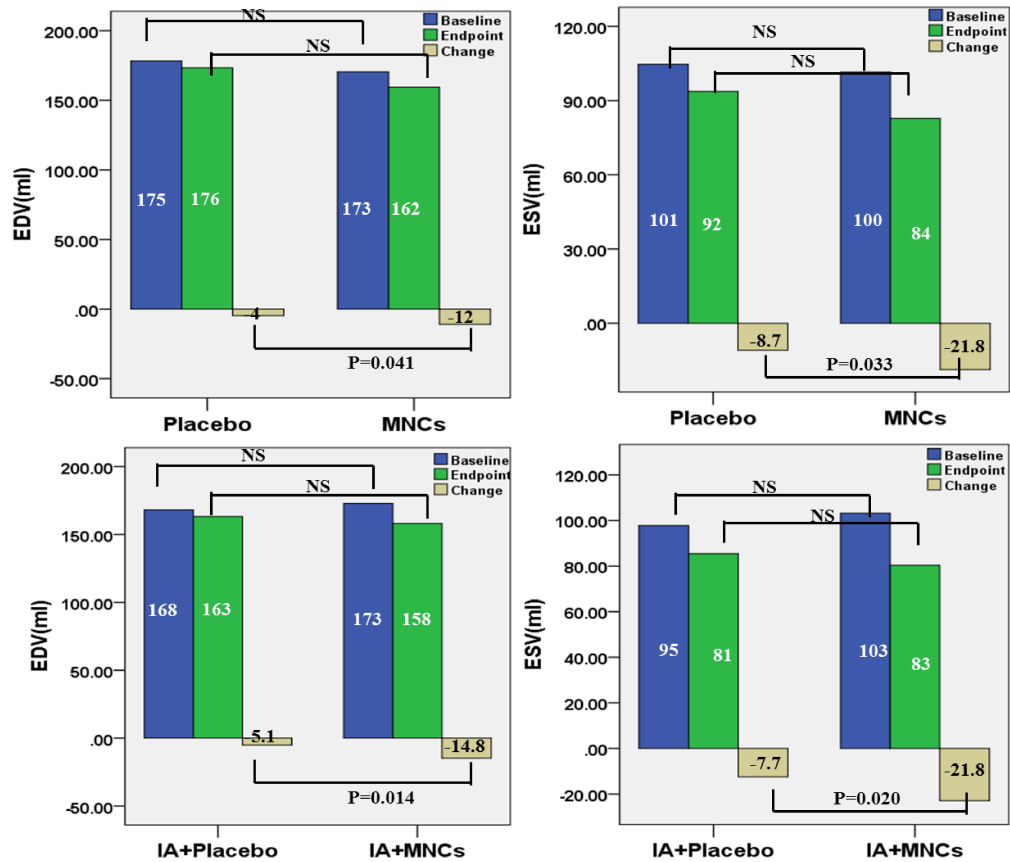
Supplemental Data Part 6: Sample size calculation

In a 2x2 factorial design trial, multiplicity issue should be considered to avoid the type I error inflation. Our detail considerations were as follow.

The present trial had three research hypotheses, the first one is to verify that bone marrow-mononuclear cells (MNC) can enhance LVEF of AMI patients; the second one is that intensive atorvastatin (IA) can increase LVEF of AMI patients than routine atorvastatin (RA); the third one is that IA can coordinate with MNC to further increase LVEF. We treated the above hypotheses in a hierarchical manner. The separate comparison of MNC and IA were tested at the first tier. In this tier, significance level was 0.025 (1-sided) for each comparison. By this way, the overall type I error could be controlled at 0.05. If any of the hypotheses in the first tier was significant, the last one assumption (IA + MNC vs. IA+Placebo) would be tested at a 2-sided 0.05 alpha level. Because the above sequential test process, it would have no impact on the overall type I error. These settings of significance level were incorporated into the sample size calculation procedure.

For the first hypothesis, according to the previous studies [Grajek S, et al. *Eur Heart J.* 2010; 31(6): 691-702. Traverse JH, et al. *JAMA.* 2011;306(19):2110-9.], LVEF at 6 months after AMI increases by $1\% \pm 3\%$ on average compared to the baseline without MNC implantation, however, MNC transplantation can increase LVEF by $4\% \pm 5\%$. When the statistical significance level is taken as 1-sided 0.025 and the power as 80%, the minimal sample size of each group is calculated as 30 cases. For the second

research hypothesis, according to the previous report [Kim EK, et al. *J Korean Med Sci*. 2015. 30(4): 435-41], IA can increase LVEF by $3\% \pm 3\%$ compared with RA. When the statistical significance level is 1-sided 0.025 and the power is 80%, the minimal sample size is 36 for each group. For the third hypothesis, according to our previous studies [Yang YJ, et al. *European Heart Journal*. 2008, 29(12):1578-90], it is estimated that the IA + MNC transplantation group can enhance LVEF of AMI patients by $9\% \pm 5\%$. When the statistical significance level is taken as 1-sided 0.025 and the power as 80%, the minimal sample size of each group is 16 cases. Combined with the sample size determined according to the statistical principle, and considering the randomization scheme and the possible maximum drop rate as 20%, this study plans to enroll 100 subjects who are allocated into group RA+Placebo, RA+MNCs, IA+Placebo, and IA+MNCs (25 cases in each group) according to the proportion of 1:1:1:1. In fact, according to the current design, this trial will provide more than 80% assurance to prove the expected research hypotheses.



Supplemental Figure 1. The reductions in LV volumes with 2DE after MNCs transplants denoting LV remodeling attenuation

2DE: two-dimensional echocardiography; EDV: end-diastolic volume; ESV: end-systolic volume; IA: intensive atorvastatin; LV: left ventricle; MNCs: mononuclear cells; RA: regular atorvastatin

Supplemental Table 1. Baseline comparison between 24 patients lost to follow-up and 76 ones completing endpoint assessments

| | Patients completing endpoint assessments (n=76) | Patients lost to follow-up (n=24) | <i>P</i> value |
|---|--|--|-----------------------|
| Male (%) | 68(89.5) | 23(95.8) | 0.649 |
| Age (years) | 52.9±11.5 | 56.3±11.9 | 0.612 |
| BMI (Kg/m ²) | 25.1±5.6 | 25.6±4.3 | 0.784 |
| Killip class | 1.87±1.32 | 1.65±0.94 | 0.311 |
| IABP (%) | 7(9.2) | 2(8.3) | 0.659 |
| Hypertension (%) | 38(50.0) | 14(58.3) | 0.225 |
| Diabetes (%) | 17(22.4) | 5(20.8) | 0.203 |
| Hyperlipidemia (%) | 26(34.2) | 9(37.5) | 0.192 |
| Stroke(%) | 1(1.3) | 1(4.1) | 0.103 |
| Smoking(%) | 53(69.7) | 18(75.0) | 0.076 |
| CHD family history(%) | 11(14.5) | 3(12.5) | 0.233 |
| Previous MI(%) | 6(7.9) | 2(8.3) | 0.655 |
| Previous coronary revascularization(%) | 3(3.9) | 1(4.2) | 0.719 |
| SBP(mmHg) | 115.3±19.2 | 120.1±15.4 | 0.628 |
| HR(bpm) | 79.5±18.6 | 79.1±16.5 | 0.733 |

| | | | |
|--------------------------|------------|------------|-------|
| ALT(IU/L) | 61.5±39.2 | 47.3±30.6 | 0.478 |
| Scr(umol/L) | 82.8±15.1 | 88.5±18.2 | 0.827 |
| HbA1c(%) | 6.2±1.1 | 6.6±1.3 | 0.895 |
| WBC(*10 ⁹ /L) | 8.8±2.9 | 9.3±1.8 | 0.859 |
| HCT(%) | 43.3±6.1 | 39.8±5.7 | 0.462 |
| HGB(g/L) | 143.6±17.1 | 139.5±15.8 | 0.489 |
| PLT(*10 ⁹ /L) | 258.1±98.6 | 221.4±94.3 | 0.724 |
| Medications | | | |
| Aspirin(%) | 76(100) | 24(100) | NS |
| Clopidogrel(%) | 76(100) | 24(100) | NS |
| Beta-blockers(%) | 70(92.1) | 21(87.5) | 0.867 |
| RAASi(%) | 56(73.7) | 20(83.3) | 0.459 |
| Spirolactone(%) | 50(65.8) | 16(66.7) | 0.872 |
| Diuretics(%) | 55(72.4) | 18(75.0) | 0.856 |

ALT: alanine aminotransferase; BMI: body mass index; HbA1c: glycosylated hemoglobin; HCT: Hematocrit; HGB: Hemoglobin; HR: heart rate; IA: intensive atorvastatin; IABP: intra-aortic balloon pump; MNCs: mononuclear cells; PLT: platelet; RA: regular atorvastatin; RAASi: renin angiotensin aldosterone system inhibitor; SBP: systolic blood pressure; Scr: Serum creatinine; WBC: white blood cell;

Supplemental Table 2. The properties of coronary lesions and intervention

| | RA+Placebo | RA+MNCs | IA+Placebo | IA+MNCs | <i>P value</i> |
|---|------------|-----------|------------|-----------|----------------|
| Emergent PCI or thrombolysis(%) | 5(26.3) | 5(25.0) | 8(47.1) | 10(50.0) | 0.225 |
| LAD patency at implant(%) | 8(42.1) | 9(45.0) | 10(58.8) | 12(60.0) | 0.272 |
| LAD total occlusion at implant (%) | 11(57.9) | 11(55.0) | 7(41.2) | 8(40.0) | 0.610 |
| LAD lesion only(%) | 7(36.8) | 10(50.0) | 6(35.3) | 12(60.0) | 0.386 |
| Dual vessel lesions(%) | 9(47.4) | 8(40.0) | 5(29.4) | 3(15.0) | 0.154 |
| Triple vessel lesions(%) | 3(15.8) | 1(5.0) | 5(29.4) | 5(25.0) | 0.215 |
| LM lesion(%) | 1(5.3) | 1(5.0) | 3(17.6) | 1(5.0) | 0.525 |
| Stent implant(%) | 18(94.7) | 18(90.0) | 14(82.4) | 18(90.0) | 0.688 |
| Complete revascularization(%) | 9(47.4) | 10(50.0) | 8(47.1) | 9(45.0) | 0.992 |
| Duration from symptom onset to MNC infusion (d) | 24.0±16.7 | 27.8±19.6 | 25.6±13.8 | 28.7±19.1 | 0.866 |

LAD: left anterior descending; LM: left main; MNCs: mononuclear cells;PCI: percutaneous coronary intervention;

Supplemental Table 3. Blood lipid profile at baseline and one-year endpoint

| | Total RA (n=39) | Total IA (n=37) | Total Placebo (n=36) | Total MNCs (n=40) |
|------------------------|------------------------------|----------------------------|---------------------------------|------------------------------|
| TG(mmol/L) | | | | |
| Baseline | 2.07±1.50 | 2.05±1.37 | 2.16±1.58 | 1.96±1.29 |
| Endpoint | 1.55±0.68 | 1.60±0.98 | 1.74±1.05 | 1.42±0.53 |
| Adjusted | -0.52 | -0.45 | -0.42 | -0.54 |
| Difference(95%CI) # | (-1.06, -0.14) | (-0.90, -0.07) | (-0.95, -0.01) | (-0.99, -0.19) |
| <i>P value</i> | 0.821 | | 0.695 | |
| TC(mmol/L) | | | | |
| Baseline | 4.45±1.00 | 4.75±1.26 | 4.87±1.19 | 4.35±1.04 |
| Endpoint | 3.72±0.81 | 3.72±1.02 | 3.91±0.80 | 3.55±0.99 |
| Adjusted | -0.72 | -1.03 | -0.96 | -0.80 |
| Difference(95%CI) # | (-1.00, -0.44) | (-1.44, -0.67) | (-1.27, -0.63) | (-1.16, -0.47) |
| <i>P value</i> | 0.203 | | 0.512 | |
| LDL-C(mmol/L) | | | | |
| Baseline | 2.41±0.78 | 2.85±0.90 | 2.78±0.87 | 2.44±0.83 |
| Endpoint | 2.09±0.78 | 2.08±0.75 | 2.19±0.71 | 2.00±0.80 |
| Adjusted | -0.32 | -0.78 | -0.59 | -0.45 |
| Difference(95%CI) # | (-0.51, -0.18) | (-0.97, -0.59) | (-0.73, -0.37) | (-0.69, -0.28) |
| <i>P value</i> | 0.013 | | 0.422 | |
| HDL-C(mmol/L) | | | | |
| Baseline | 1.01±0.25 | 0.95±0.29 | 0.97±0.26 | 1.00±0.28 |
| Endpoint | 1.03±0.22 | 1.06±0.25 | 1.05±0.19 | 1.04±0.27 |
| Adjusted | 0.02 | 0.05 | 0.06 | 0.04 |
| Difference(95%CI) # | (-0.15, 0.11) | (-0.20, 0.23) | (0.41, 1.02) | (0.08, 0.71) |
| <i>P value</i> | 0.145 | | 0.378 | |
| Total RA (n=39) | | | | |
| | RA+Placebo (n=19) | RA+MNCs (n=20) | Total IA (n=37) | |
| | | | IA+Placebo (n=17) | IA+MNCs (n=20) |

| TG(mmol/L) | | | | |
|----------------------|---------------|---------------|---------------|---------------|
| Baseline | 2.11±1.68 | 2.02±1.35 | 2.22±1.51 | 1.90±1.25 |
| Endpoint | 1.71±0.82 | 1.39±0.47 | 1.78±1.28 | 1.44±0.60 |
| Adjusted | -0.40 | -0.63 | -0.44 | -0.46 |
| Difference(95%CI) # | (-1.17,0.14) | (-1.23,-0.17) | (-1.07,0.10) | (-1.12,0.04) |
| <i>P value</i> | 0.634 | | 0.967 | |
| TC(mmol/L) | | | | |
| Baseline | 4.69±1.01 | 4.22±0.97 | 5.06±1.36 | 4.49±1.12 |
| Endpoint | 3.80±0.79 | 3.66±0.86 | 4.04±0.82 | 3.45±1.12 |
| Adjusted | -0.90 | -0.56 | -1.03 | -1.04 |
| Difference(95%CI) # | (-1.33,-0.46) | (-0.93,-0.20) | (-1.50,-0.59) | (-1.64,-0.51) |
| <i>P value</i> | 0.269 | | 0.985 | |
| LDL-C(mmol/L) | | | | |
| Baseline | 2.61±0.75 | 2.22±0.78 | 3.04±0.99 | 2.71±0.83 |
| Endpoint | 2.13±0.70 | 2.06±0.86 | 2.28±0.73 | 1.92±0.75 |
| Adjusted | -0.48 | -0.16 | -0.76 | -0.79 |
| Difference(95%CI) # | (-0.76,-0.18) | (-0.33,-0.03) | (-0.54,-0.25) | (-1.12,-0.33) |
| <i>P value</i> | 0.179 | | 0.896 | |
| HDL-C(mmol/L) | | | | |
| Baseline | 0.99±0.23 | 1.03±0.27 | 0.93±0.29 | 0.96±0.29 |
| Endpoint | 1.00±0.15 | 1.06±0.28 | 1.13±0.23 | 1.01±0.26 |
| Adjusted | 0.11 | 0.03 | 0.20 | 0.05 |
| Difference(95%CI) # | (0.05,0.21) | (0,0.17) | (0.09,0.31) | (0,0.25) |
| <i>P value</i> | 0.857 | | 0.209 | |

#values are median (95%CI), and adjusted for values at baseline and the other intervention

HDL-C:high density lipoprotein-cholesterol; IA: intensive atorvastatin; LDL-C:low density lipoprotein-cholesterol; MNCs: mononuclear cells;RA: regular atorvastatin; TC: total cholesterol; TG: triglyceride

**Supplemental Table 4 A. Comparison of LVEF by MRI
after adding “missing values” of 24 patients lost to
follow-up**

| | Total RA (n=50) | Total IA (n=50) | Total Placebo (n=50) | Total MNCs (n=50) |
|------------------------------------|----------------------------|----------------------------|---------------------------------|------------------------------|
| LVEF | | | | |
| Baseline | 35.3(32.1,40.5) | 31.7(30.8,37.2) | 32.4(31.2,34.8) | 33.0(31.5,37.4) |
| Endpoint | 42.4(40.1,45.3) | 41.6(39.0,46.0) | 36.5(37.0,43.3) | 43.3(41.5,48.1)* |
| Adjusted Difference# (95%CI) | 5.4(3.6,8.5) | 8.2(6.4,10.7) | 3.5(2.1,8.0) | 8.6(6.4,11.2) |
| <i>P value</i> | 0.085 | | 0.026 | |

#values are median (95%CI), and adjusted for values at baseline and the other intervention;

* endpoint values in total MNCs group vs total Placebo group(P=0.013);

**Table 4 B. Comparison of LVEF between ATV groups after
adding “missing values” of 24 patients lost to follow-up**

| | Total RA (n=50) | | Total IA (n=50) | |
|------------------------------------|------------------------------|---------------------------|------------------------------|---------------------------|
| | RA+Placebo (n=25) | RA+MNCs (n=25) | IA+Placebo (n=25) | IA+MNCs (n=25) |
| LVEF | | | | |
| Baseline | 35.2(27.5,38.1) | 34.8(32.2,36.9) | 33.1(29.9,35.0) | 31.8(29.2,36.5) |
| Endpoint | 40.3(34.4,43.2) | 43.5(39.2,46.4) | 36.9(33.5,40.7) | 43.4(40.5,50.0)* |
| Adjusted Difference# (95%CI) | 2.9(0.3,8.1) | 5.8(4.1,8.9) | 3.3(2.0,8.1) | 9.5(7.7,16.5) |
| <i>P value</i> | 0.751 | | 0.005 | |

#values are median (95%CI), and adjusted for values at baseline and the other intervention;

* the comparison of endpoint values in IA+ MNCs group vs. IA+Placebo group(P=0.001);

ATV: atorvastatin; IA: intensive atorvastatin; LVEF: left ventricular ejection fraction; MNCs: mononuclear cells; RA: regular atorvastatin.

Supplemental Table 5. The structural and functional changes of left ventricle with 2DE after MNCs therapy

| | Total RA (n=39) | Total IA (n=37) | Total Placebo (n=36) | Total MNCs (n=40) |
|--|------------------------------|---------------------------|------------------------------|-----------------------|
| LVEF(%) | | | | |
| Baseline | 43(40,45) | 40(40,45) | 41(40,45) | 40(40,45) |
| Endpoint | 45(43,50) | 50(46,53) | 47(44,50) | 49(45,54) |
| Adjusted Difference [#] (95%CI) | 5(2,7) | 8(5,10) | 4(2,6) | 9(5,10) |
| <i>P value</i> | | 0.156 | | 0.022 |
| EDV(ml) | | | | |
| Baseline | 174(169,180) | 175(167,180) | 175(170,181) | 173(164,180) |
| Endpoint | 170(160,177) | 167(153,175) | 176(164,184) | 162(153,170) |
| Adjusted Difference [#] (95%CI) | -6.0(-10.0,-3.0) | -11.0(-14.0,-5.0) | -4.0(-7.0,-3.0) | -12.0(-17.0,-8.0) |
| <i>P value</i> | | 0.147 | | 0.041 |
| ESV(ml) | | | | |
| Baseline | 100(95,110) | 100(98,105) | 101(98,105) | 100(94,108) |
| Endpoint | 92(86,99) | 83(68,90) | 92(85,99) | 84(69,90) |
| Adjusted Difference [#] (95%CI) | -11.0(-20.2,-6.7) | -19.8(-24.4,-10.4) | -8.7(-13.8,-6.4) | -21.8(-25.3,-14.5) |
| <i>P value</i> | | 0.113 | | 0.033 |
| Total RA (n=39) | | | | |
| | RA+Placebo (n=19) | RA+MNCs (n=20) | Total IA (n=37) | |
| | | | IA+Placebo (n=17) | IA+MNCs (n=20) |
| LVEF(%) | | | | |
| Baseline | 41(40,45) | 42(37,45) | 42(37,45) | 40(39,45) |
| Endpoint | 45(43,50) | 46(42,55) | 49(43,53) | 50(46,54) |
| Adjusted Difference [#] (95%CI) | 3(2,7) | 6(0,14) | 4(0,10) | 10(8,12) |
| <i>P value</i> | | 0.188 | | 0.045 |
| EDV(ml) | | | | |
| Baseline | 187(176,201) | 167(157,179) | 168(155,180) | 173(163,184) |
| Endpoint | 182(171,197) | 161(147,175)§ | 163(147,177) | 158(145,171) |
| Adjusted Difference [#] | -4.7(-7.3,-1.8) | -7.2(-14.5,0.6) | -5.1(-9.4,-0.9) | -14.8(-20.2,-8.9) |

| (95%CI) | | | | |
|---------------------------------|-------------------|------------------|------------------|--------------------|
| <i>P value</i> | | 0.578 | | 0.014 |
| ESV(ml) | | | | |
| Baseline | 103(99,104) | 95(90,111) | 95(90,111) | 103(99,114) |
| Endpoint | 92(90,100) | 85(65,102) | 81(67,105) | 83 (68,89) |
| Adjusted | -10.0(-18.5,-6.7) | -19.4(-25.3,3.3) | -7.7(-24.4,-1.8) | -21.8(-29.4,-18.7) |
| Difference [#] (95%CI) | | | | |
| <i>P value</i> | | 0.381 | | 0.020 |

#values are median (95%CI) , and adjusted for values at baseline and the other intervention;

§the comparison of endpoint values in RA+ MNCs group vs RA+Placebo group(P=0.033)

2DE:two-dimensional echocardiography; EDV: end-diastolic volume; ESV: end-systolic volum;

IA: intensive atorvastatin; LVEF: left ventricular ejection fraction; MNCs: mononuclear cells;RA:

regular atorvastatin;

**Supplemental Table 6. Myocardial perfusion with SPECT
after MNCs transplantation**

| | Total RA | Total IA | Total Placebo | Total MNCs |
|------------------------------------|-------------------|-----------------|----------------------|-------------------|
| | (n=39) | (n=37) | (n=36) | (n=40) |
| Perfusion Defect(%) | | | | |
| Baseline | 34.2(31,37.3) | 32(29,35) | 33.5(30.3,36.7) | 32.8(30.1,35.5) |
| Endpoint | 32.1(29,35.1) | 29.5(26.7,32.5) | 31.4(28.3,34.5) | 30.3(27.7,32.9) |
| Adjusted | -2.1(-2.4,-1.8) | -2.5(-2.8,-2.1) | -2.1(-2.3,-1.8) | -2.5(-2.9,-2.1) |
| Difference [#] (95%CI) | | | | |
| <i>P value</i> | | 0.541 | | 0.772 |
| SRS | | | | |
| Baseline | 22.2±6.9 | 20.9±6.5 | 21.7±7.3 | 21.4±6.3 |
| Endpoint | 21.7±6.9 | 20.1±6.5 | 21.3±7.4 | 20.6±6.0 |
| Adjusted | -0.40 | -0.41 | -0.24 | -0.57 |
| Difference [#] (95%CI) | (-0.67,-0.17) | (-0.70,-0.18) | (-0.50,-0.07) | (-0.88,-0.31) |
| <i>P value</i> | | 0.810 | | 0.078 |
| Total RA (n=39) | | | | |
| Total IA (n=37) | | | | |
| | RA+Placebo | RA+MNCs | IA+Placebo | IA+MNCs |
| | (n=19) | (n=20) | (n=17) | (n=20) |
| Perfusion Defect(%) | | | | |
| Baseline | 34.7(29.9,40.3) | 33.8(29.9,37.9) | 32.3(27.8,37.1) | 31.8(28.1,35.8) |
| Endpoint | 32.8(27.9,38.2) | 31.5(27.7,35.4) | 30.1(25.6,34.8) | 29.1(25.6,32.8) |
| Adjusted | -1.9(-2.4,-1.6) | -2.3(-2.8,-1.8) | -2.2(-2.5,-1.9) | -2.7(-3.3,-2.0) |
| Difference [#] (95%CI) | | | | |
| <i>P value</i> | | 0.285 | | 0.288 |
| SRS | | | | |
| Baseline | 22.4±7.5 | 22.1±6.6 | 21.0±7.2 | 20.8±6.1 |
| Endpoint | 22.2±7.5 | 21.4±6.4 | 20.5±7.5 | 19.9±5.6 |
| Adjusted | -0.19 | -0.64 | -0.31 | -0.50 |
| Difference [#] (95%CI) | (-0.56,0) | (-1.15,-0.20) | (-0.73,0) | (-0.93,-0.21) |
| <i>P value</i> | | 0.101 | | 0.390 |

#values are median (95%CI) , and adjusted for values at baseline and the other intervention;

IA: intensive atorvastatin; MNCs: mononuclear cells;RA: regular atorvastatin; SRS: summed rest score

Supplemental Table 7. Biomarkers at baseline and endpoint

| | Total RA (n=39) | Total IA (n=37) | Total Placebo (n=36) | Total MNCs (n=40) |
|---------------------------------|-----------------------|-----------------------|-------------------------|-----------------------|
| NT-proBNP(pg/ml) | | | | |
| Baseline | 1716.7±135.5 | 1801.5±177.0 | 1703.8±132.8 | 1801.8±170.7 |
| Endpoint | 754.2±82.2 | 590.0±124.3 | 914.2±127.8 | 669.9±55.6 |
| Adjusted difference# (95%CI) | -962.4±155.8 | -1211.5±179.8 | -789.6±143.3 | -1340.2±175.5 |
| <i>P value</i> | 0.316 | | 0.023 | |
| hs-CRP(mg/L) | | | | |
| Baseline | 7.7±5.2 | 7.8±4.6 | 8.3±4.7 | 7.2±5.1 |
| Endpoint | 2.5±2.2 | 1.3±1.0§ | 1.9±1.6 | 2.5±1.8 |
| Adjusted difference# (95%CI) | 5.9(5.0,6.8) | 6.8(5.9,7.7) | 6.1(5.2,7.1) | 6.7(5.8,7.6) |
| <i>P value</i> | 0.369 | | 0.392 | |
| Endothelin(pmol/L) | | | | |
| Baseline | 0.74 (0.55, 0.86) | 0.67 (0.52, 0.89) | 0.65 (0.54, 0.84) | 0.82 (0.54, 1.20) |
| Endpoint | 0.50 (0.43, 0.72) | 0.54 (0.40, 0.74) | 0.54 (0.39, 0.81) | 0.51 (0.43, 0.73) |
| Adjusted difference# (95%CI) | 0.09 (-0.09, 0.40) | 0.11 (-0.14, 0.55) | 0.06 (-0.10, 0.33) | 0.10 (-0.13, 0.63) |
| <i>P value</i> | 0.950 | | 0.799 | |
| | | | | |
| | Total RA (n=39) | | Total IA (n=37) | |
| | RA+Placebo (n=19) | RA+MNCs (n=20) | IA+Placebo (n=17) | IA+MNCs (n=20) |
| NT-proBNP(pg/ml) | | | | |
| Baseline | 1644.3±163.0 | 1776.2±201.4 | 1767.8±222.3 | 1841.3±288.5 |
| Endpoint | 921.2±170.5 | 616.7±68.5 | 906.6±190.1 | 215.8±18.0* |
| Difference | -723.1±238.3 | -1159.5±206.6 | -861.2±172.3 | -1652.5±289.3 |
| <i>P value</i> | 0.190 | | 0.034 | |
| hs-CRP(mg/L) | | | | |

| | | | | |
|------------------------------|-------------------------|-------------------------|------------------------|------------------------|
| Baseline | 8.13±4.81 | 7.19±5.67 | 9.10±4.66 | 7.36±4.61 |
| Endpoint | 2.17±2.13 | 2.24±2.06 | 1.00±0.89 | 1.24±0.62 |
| Adjusted Difference# (95%CI) | -6.07 (-5.47,-7.08) | -4.86 (-4.03,-5.68) | -8.06 (-7.17,-8.98) | -6.13 (-5.43,-7.01) |
| <i>P</i> value | 0.563 | | 0.370 | |
| Endothelin(pmol/L) | | | | |
| Baseline | 0.73 (0.51, 0.80) | 0.71 (0.57, 0.99) | 0.68 (0.59, 0.86) | 0.81 (0.62, 1.00) |
| Endpoint | 0.53 (0.41, 0.68) | 0.50 (0.38, 0.65) | 0.52 (0.37, 0.76) | 0.49 (0.33, 0.61) |
| Adjusted Difference(95%CI) # | -0.09 (-0.06, -0.31) | -0.13 (-0.16, -0.45) | 0.07 (-0.45,0.19) | -0.20 (-0.61,0.22) |
| <i>P</i> value | 0.180 | | 0.046 | |

#values are median (95%CI) , and adjusted for values at baseline and the other intervention;

§ the endpoint values in total RA vs total IA group (P=0.001);

*the endpoint values in total MNCs group vs total Placebo group (P=0.005)

hs-CRP: high sensitivity C reactive protein; IA: intensive atorvastatin; MNCs: mononuclear cells;NT-proBNP:amino-terminal pro-B-type natriuretic peptide; RA: regular atorvastatin

Supplemental Table 8. MACEs after MNC therapy

| | Total RA (n=39) | Total IA (n=37) | Total Placebo (n=36) | Total MNCs (n=40) |
|-----------------------|-----------------------------|---------------------------|--------------------------------|-----------------------------|
| Death | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| Recurrent MI | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| TVR | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| New arrhythmia | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| | Total RA (n=39) | | Total IA (n=37) | |
| | RA+Placebo (n=19) | RA+MNCs (n=20) | IA+Placebo (n=17) | IA+MNCs (n=20) |
| Death | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| Recurrent MI | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| TVR | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |
| New arrhythmia | 0 | 0 | 0 | 0 |
| <i>P value</i> | | NS | | NS |

IA: intensive atorvastatin; MI: myocardial infarction; MNCs: mononuclear cells; RA: regular atorvastatin; TVR: target vascular revascularization

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