

## Supplemental Material

**Supplemental Table 1.** Forward and reverse primer sequences used for RT-PCR.

<b>Gene Name</b>	<b>Gene ID</b>	<b>Accession #</b>	<b>Sequence</b>
<b>IFN-γ</b>	15978	NM_008337	<b>F:</b> TTGGCTTGAGCTTCC <b>R:</b> TGACTGTGCCGTGGCAGTA
<b>IL-12Rβ2</b>	16162	NM_008354	<b>F:</b> CAAGCATTGCATCGCTATCA <b>R:</b> TTGCCGGAAGTAACGAATTGA
<b>CCR5</b>	12774	NM_009917	<b>F:</b> GTCAAACGCTTTGCAAACG <b>R:</b> TGAGCTTGCACGATCAGGATT
<b>IL-18</b>	16173	NM_008360	<b>F:</b> CACATGCGCCTTGTGATGAC <b>R:</b> TGCAGCCTCGGGTATTCTGT
<b>IL-17a</b>	16171	NM_010552	<b>F:</b> CCTGGCGGCTACAGTGAAG <b>R:</b> TTTGGACACCGCTGAGCTTTG
<b>IL-2</b>	16183	NM_008366	<b>F:</b> AAACTAAAGGGCTCTGACAACACA <b>R:</b> CACCACAGTTGCTGACTCATCA
<b>IL-4</b>	16189	NM_021283	<b>F:</b> GGTCTCAACCCCCAGCTAGT <b>R:</b> GCCGATGATCTCTCAAGTGAT
<b>IL-13</b>	16163	NM_008355	<b>F:</b> TTGAGGAGCTGAGCAACATCAC <b>R:</b> CCATGCTGCCGTTGCA
<b>IL-5</b>	16191	NM_010558	<b>F:</b> TGCACTTGAGTGTCTGACTCTCA <b>R:</b> TGTGCTCATGGAAATCTCCAT
<b>IL-33</b>	77125	NM_001164724	<b>F:</b> ACCAGGTGCTACTACGCTACTATGAG <b>R:</b> CACACCGTCGCCTGATTG
<b>TGF-β</b>	21803	NM_011577	<b>F:</b> GCAGTGGCTGAACCAAGGA <b>R:</b> AGCAGTGAGCGCTGAATCG
<b>IL-10</b>	16153	NM_010548	<b>F:</b> GATGCCCCAGGCAGAGAA <b>R:</b> CACCCAGGGAATTCAAATGC

**Supplemental Figure Legends**

**Figure S1.** **(A)** Representative M-mode echocardiograms and quantitative group data for LV ejection fraction (EF), end-diastolic volume (EDV), and end-systolic volume (ESV) from heart failure (HF) and sham mice 8 w after permanent coronary ligation or sham surgery (n = 5-6 per group). **(B)** Representative gross images of a sham-operated and failing heart, and group gravimetric data for heart, and wet lung weights. **(C)** Representative gross image of spleens isolated from a HF and sham mouse and group quantitation of organ weight, and total cell counts for CD45<sup>+</sup> leukocytes and CD3<sup>+</sup> T-cells per spleen. All weights normalized to tibia length (n = 4-6 per group).

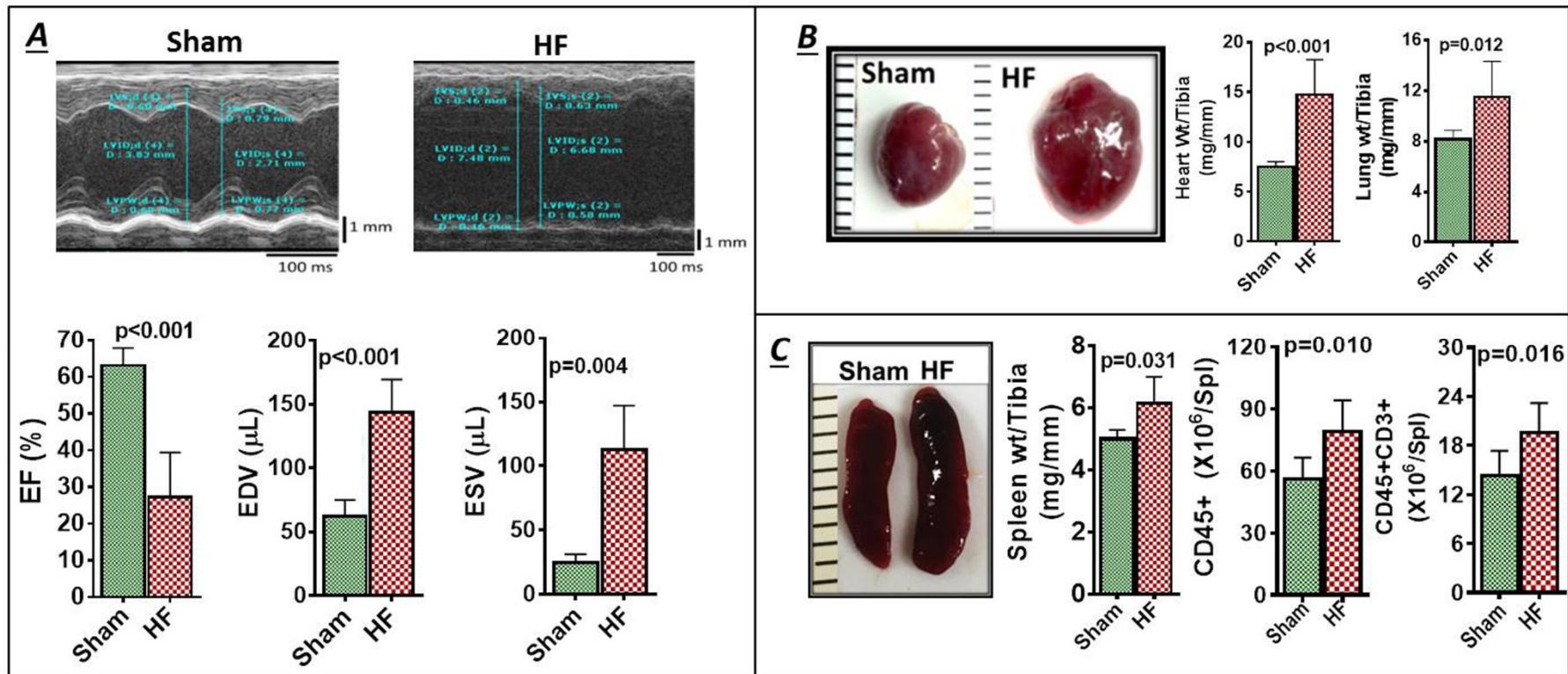
**Figure S2.** **(A)** Representative flow cytometry scatter plots for CD3<sup>+</sup>CD4<sup>+</sup> helper and CD3<sup>+</sup>CD8<sup>+</sup> cytotoxic T-cells and group quantitation of CD8<sup>+</sup> T-cell frequency in the circulation at different time intervals post-MI, and absolute cell counts at 8 w post-MI (HF) or sham surgery. **(B)** Representative confocal microscopy images of CD8<sup>+</sup> T-cells (green) in sham-operated and failing hearts (8 w post-MI) and group quantitation. **(C)** Representative flow cytometry density plots and group quantitation of CD3<sup>+</sup>CD8<sup>+</sup> T-cells in hearts, **(D)** spleens, and **(E)** lymph nodes (LNs) isolated from sham-operated and HF mice (n = 5-10).

**Figure S3.** **(A)** Example gross images of heart-draining mediastinal lymph nodes (LNs) from a sham-operated and HF mouse (*Inset*), along with flow cytometry scatter plots for live cell SSC/FSC gate and CD3<sup>+</sup>CD4<sup>+</sup> helper and CD3<sup>+</sup>CD4<sup>+</sup>Foxp3<sup>+</sup> regulatory T-cells in mediastinal LNs and **(B)** corresponding group quantitation. **(C)** Group quantitation of the frequency and absolute number of CD4<sup>+</sup> T-cell LN subsets: Th1 (IFN- $\gamma$ <sup>+</sup>), Th2 (IL-4<sup>+</sup>), and Th17 (IL-17<sup>+</sup>) (n = 4-6 per group).

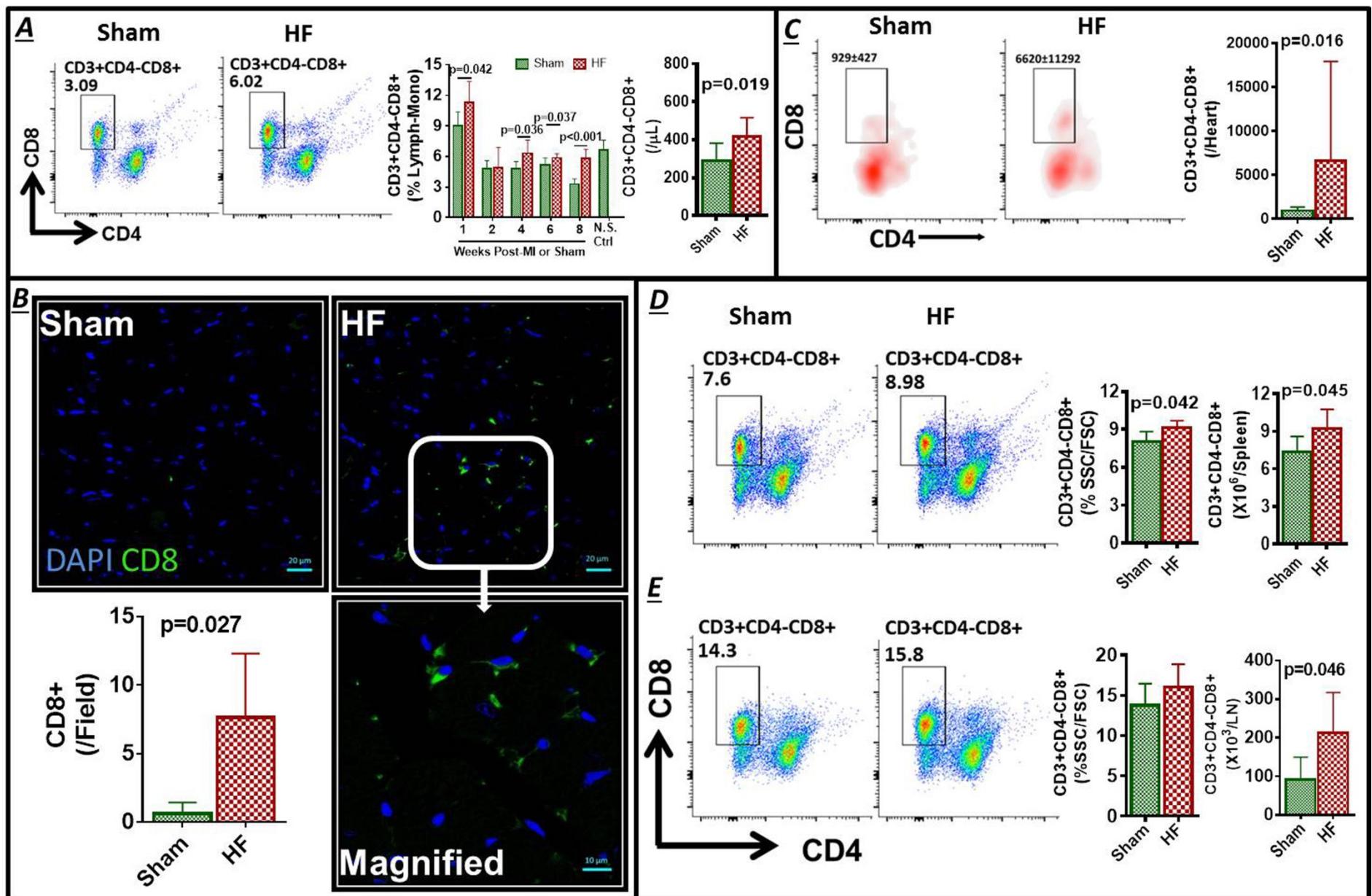
**Figure S4.** Group quantitation of baseline end-diastolic and end-systolic volume (EDV and ESV), and ejection fraction (EF) in HF mice 4 w post-MI prior to randomization to treatment with either isotype IgG or anti-CD4 antibody (n = 6-10 per group).

**Figure S5.** Naïve mice were adoptively transferred with cardiac CD3<sup>+</sup> T-cells sorted from either naïve control or HF mice and followed for 8 w, as outlined in Figure 6A. **(A)** Quantitative group data for changes in LVEF, EDV, and ESV, in recipient mice 4 and 8 w after cardiac CD3<sup>+</sup> T-cell transfer, along with measured septal and posterior wall (PW) thickness at 8 w. **(B)** Gravimetric data for the same experimental groups at 8 w after transfer. Representative Masson's trichrome stains and quantitation of fibrosis **(C)**, and wheat germ-agglutinin stains (cell membranes green, nuclei blue with DAPI counterstain; scale bar 200 µm) and quantitation of myocyte size **(D)** in mouse hearts 8 w after transfer of naïve control or HF-activated cardiac CD3<sup>+</sup> T-cells. Arrows indicate representative myocytes used for measurement of cross-sectional area.

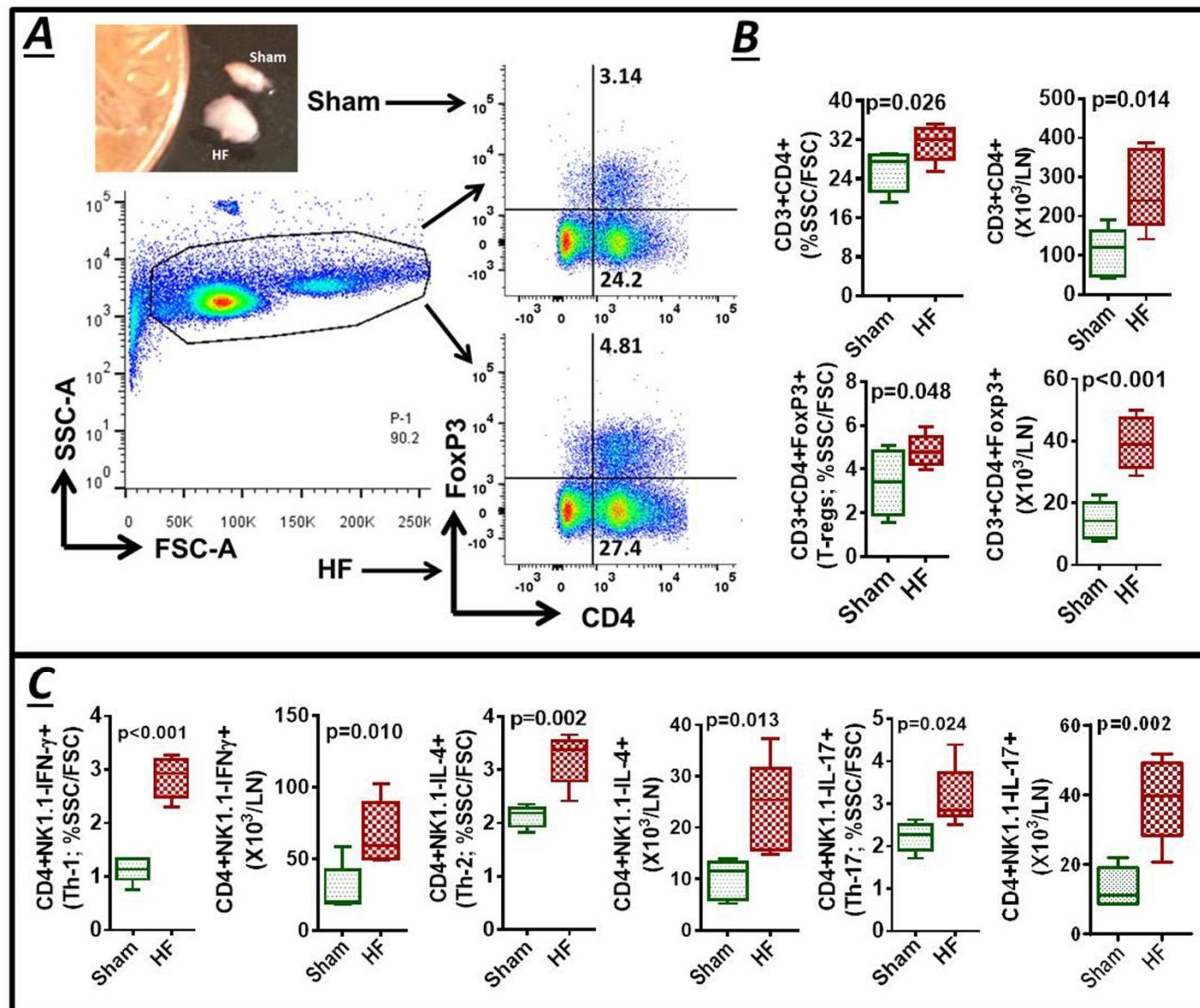
**Figure S1**



**Figure S2**



**Figure S3**



**Figure S4**

Baseline LV parameters prior to antibody treatment

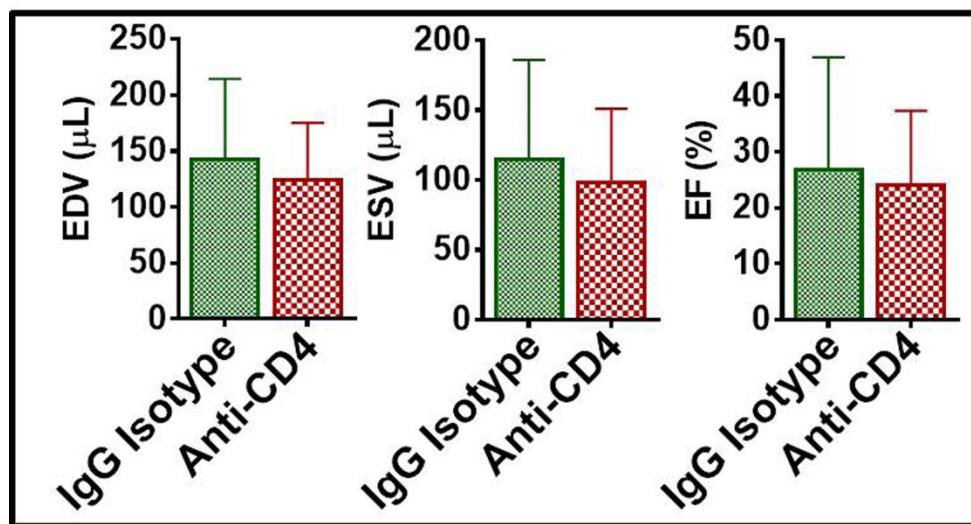


Figure S5

