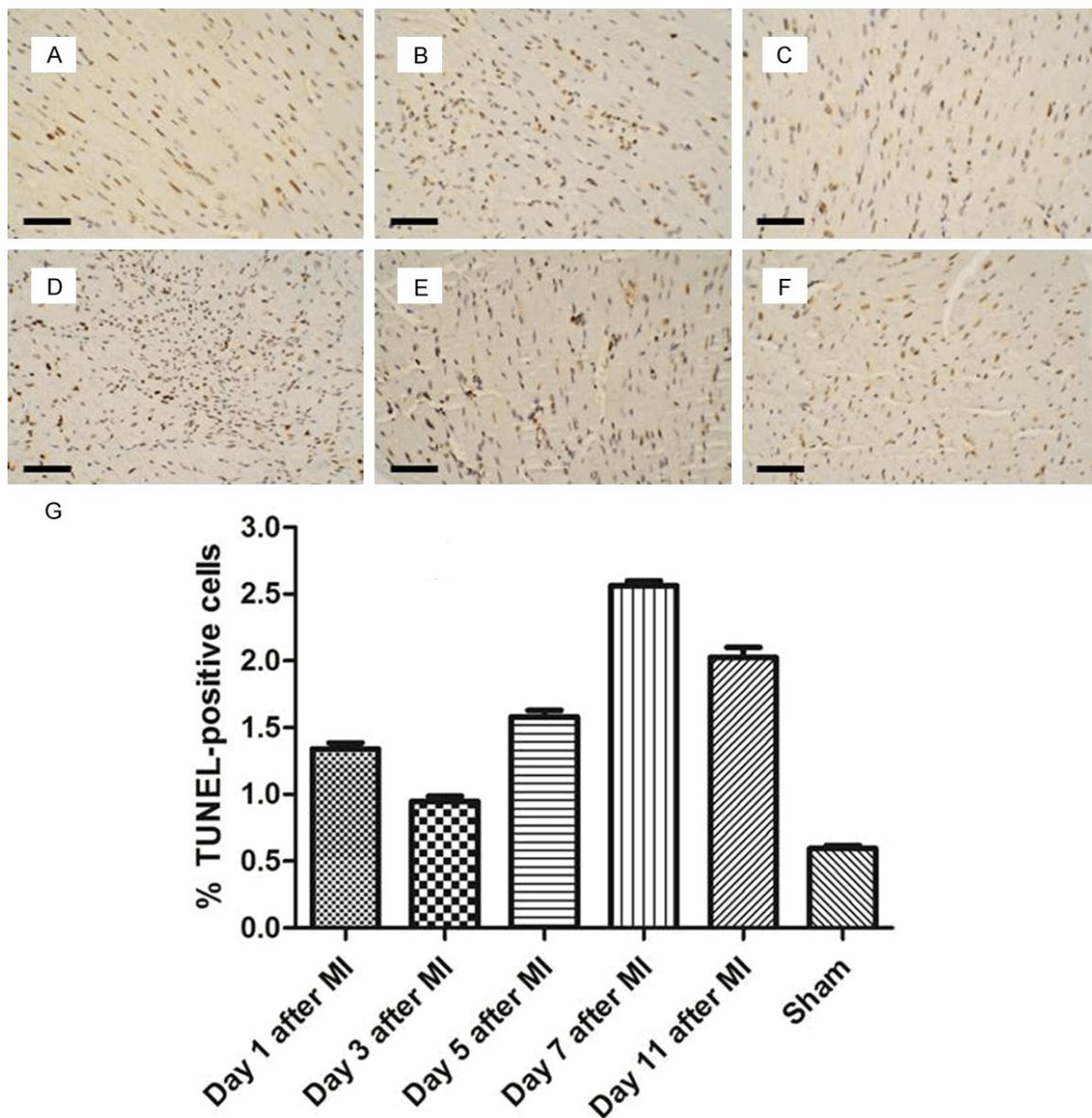


mERa modulating cardiac repair by BM c-kit⁺ cells

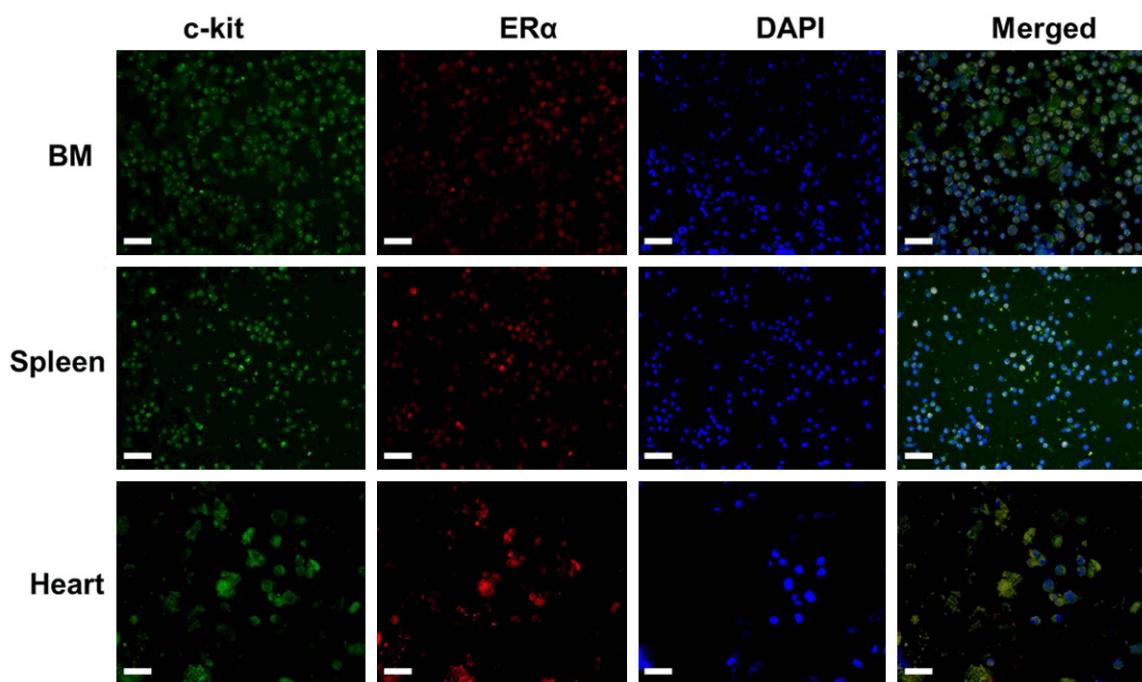
Supplementary Table 1. Oligonucleotide primers used for gene expression analysis by real-time PCR

Gene	Primer sequences (5'-3')	Location
Gapdh	Fwd: TGTGTCGTCGTGGATCTGA Rev: TTGCTGTTGAAGTCGCAGGAG	761-910
ERα	Fwd: ACATGTTGCTGGCTACGTCA Rev: GGGCACAAACGTTCTTGCATT	1470-1798
ERβ	Fwd: AATGTGCTAGTGAGCCGTCC Rev: CCTCATCCCTGTCCAGAACG	1239-1503
Gata4	Fwd: CGGAAGCCCAAGAACCTGAATA Rev: TTGCTGGAGTTACCGCTGGAG	1562 -1659
Tbx5	Fwd: AAGTTCCGAAGTGGGGTCAC Rev: GGAAGACGTGGGTGCAAAC	10-433
Ki67	Fwd: CCTGCCGACCCCTACAAAAT Rev: TCCGCCGTCTTAAGGTAGGA	2248-2560
Tbx3	Fwd: GAGGCCAAGGAACTTGGGA Rev: GCTTGGGAAGGCCAAAGTAA	1669-2022
Akt	Fwd: CTGGAGGACAACGACTACGG Rev: CTCACTGTCCACACACTCCAT	1331-1762
Notch	Fwd: GCTCCGAGGAGATCAACGAG Rev: TTGACATCACCCCTCACACCG	3803-4070



Supplementary Figure 1. TUNEL staining for heart sections from mouse in different days of AMI and sham mouse (A-E) TUNEL staining for heart sections from mouse in day 2, 3, 5, 7 and 11 after MI; (F) TUNEL staining for heart sections from Sham-operated mouse; (G) Analysis of TUNEL positive cells in heart sections after AMI. Scale bar =50 μ m.

mERa modulating cardiac repair by BM c-kit⁺ cells



Supplementary Figure 2. Immunofluorescence staining images of c-kit and ER α in BM, Spleen and heart after 7 days of AMI. After MI, ER α expressions were detected in BM, spleen and heart, especially in c-kit⁺ cells. Scale bar =25 μ m.