

Supporting Information

A dual aperture (mesoporous and macroporous) system loaded with cell-free fat extract to optimize bone regeneration microenvironment

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1. Supplemental Table

Table S1. Primers used for real-time RT-PCR.

Genes	Sequence (5'-3')	
TNF- α	5' CCT CCC TCT CAT CAG TTC TA 3'	5' ACT TGG TGG TTT GCT ACG AC 3'
IL-1 β	5' CAG CTT TCG ACA GTG AGG AGA 3'	5' TTG TCG AGA TGC TGC TGT GA 3'
<i>ocn</i>	5' CTC ACA CTC CTC GCC CTA TT 3'	5' CCC AGC CAT TGA TAC AGG TAG 3'
<i>osx</i>	5' CAG TTG ATA GGG TTT CTC TTG TA3'	5' CAT AGG ACT TGA GGT TTC ACA G 3
<i>opn</i>	5' CAT TCC GAT GTG ATT GAT AGT C 3	5' CTT CCT TAC TTT TGG GGT CTA C 3
<i>alp</i>	5' TAC AAG CAC TCC CAC TTC ATC 3'	5' AGA CCC AAT AGG TAG TCC ACA T 3'
<i>vegf</i>	5' TGT GCC CAC TGA GGA GTC 3'	5' CAT TTG TTG TGC TGT AGG AAG 3
<i>pdgh-bb</i>	5' GCT GTT GAG GTG GCT GTA GAT 3	5' GCT GTT GAG GTG GCT GTA GAT 3
<i>gapdh</i>	5' GCT GTT GAG GTG GCT GTA GAT 3	5' GCT GTT GAG GTG GCT GTA GAT 3

2. Supplemental Figures

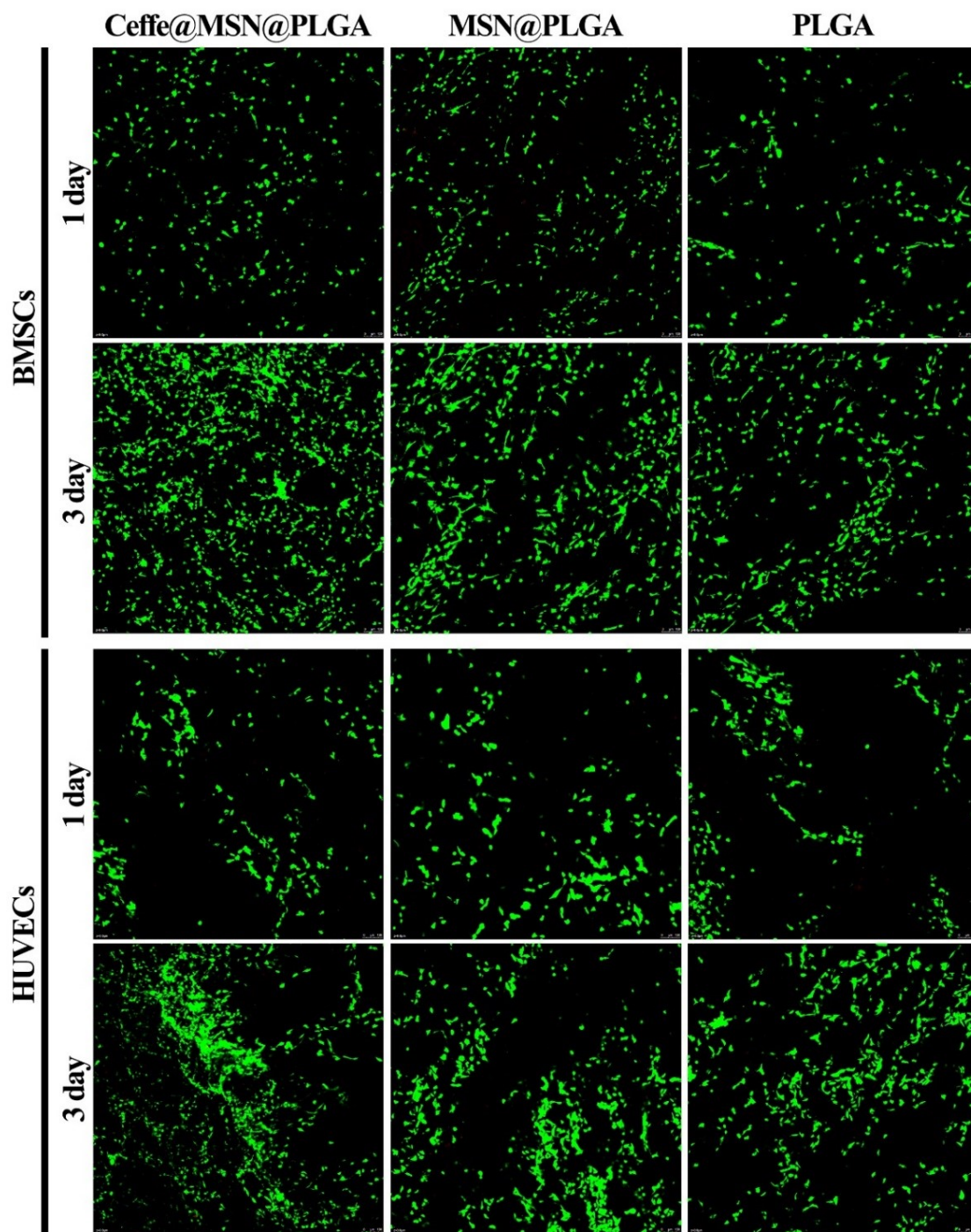


Figure S1. Confocal images of Living/Dead cell staining of HUVECs and BMSCs after cocultured with different scaffold for 1day and 3 day.

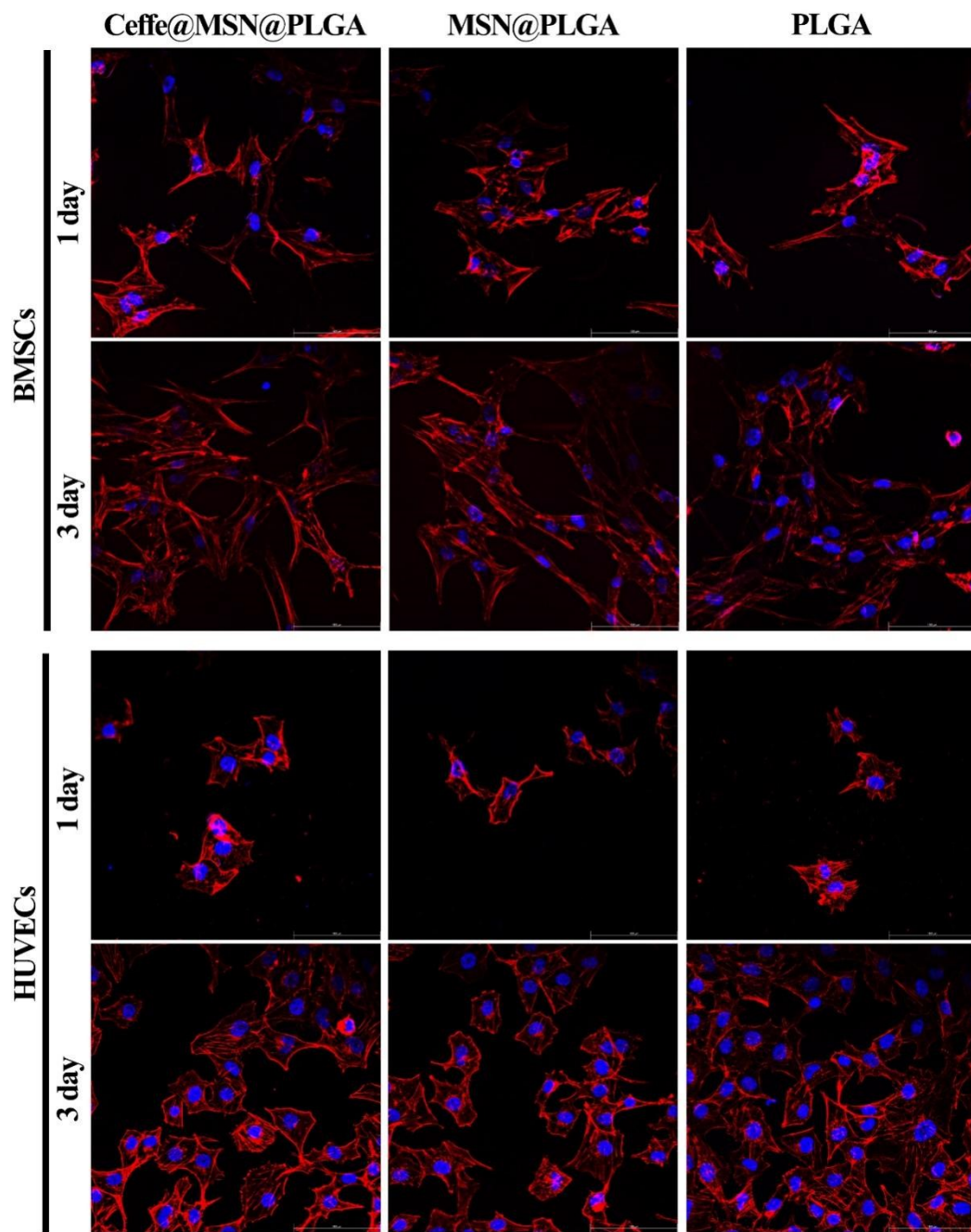


Figure S2. The images of cytoskeleton staining of HUVECs and BMSCs co-cultured with different scaffolds for 1 day and 3 day were taken by laser confocal microscopy.

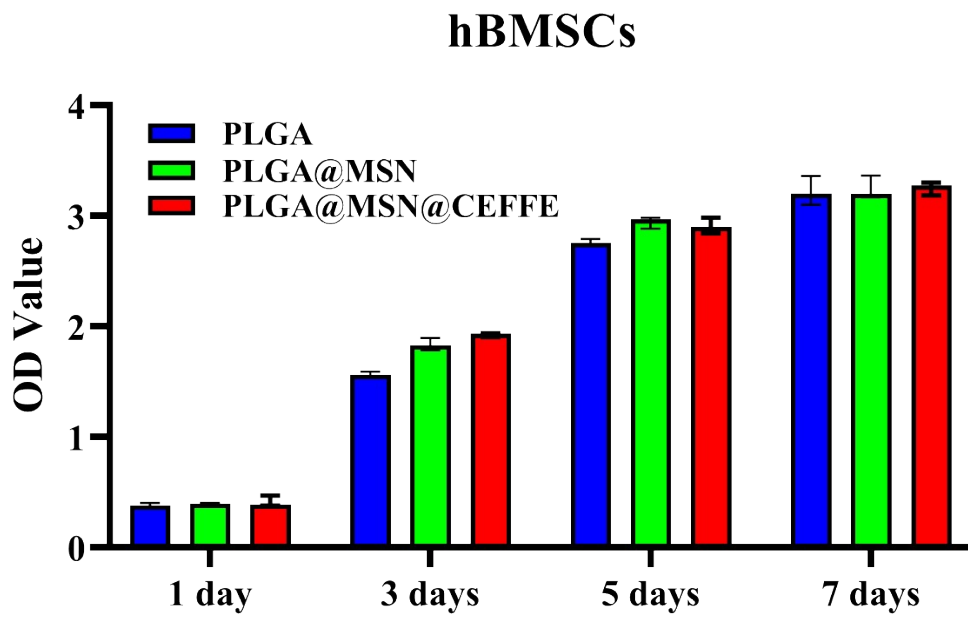


Figure S3. The Cck-8 assay kit was used to evaluate the cell proliferation of hBMSCs co-cultured with different scaffolds for 1, 3, 5, and 7 days, respectively.

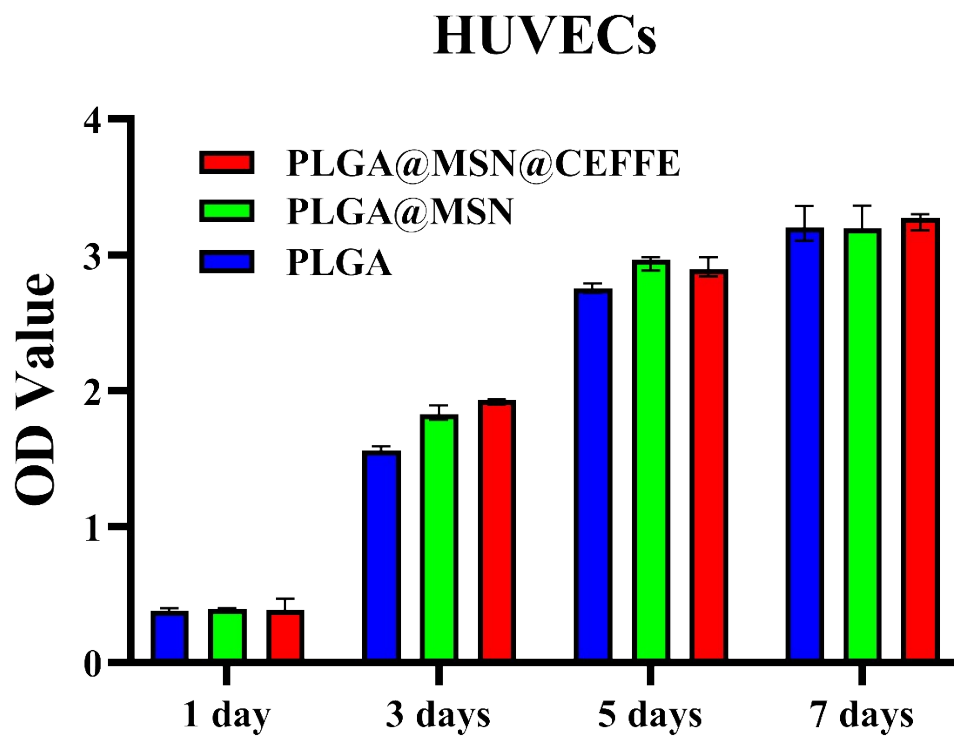


Figure S4. The Cck-8 assay kit was used to evaluate the cell proliferation of HUVECs co-cultured with different scaffolds for 1, 3, 5, and 7 days, respectively.