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Bionic Biphasic Composite Scaffold with Osteochondrogenic Factors for Regeneration of Full-Thickness Osteochondral Defect

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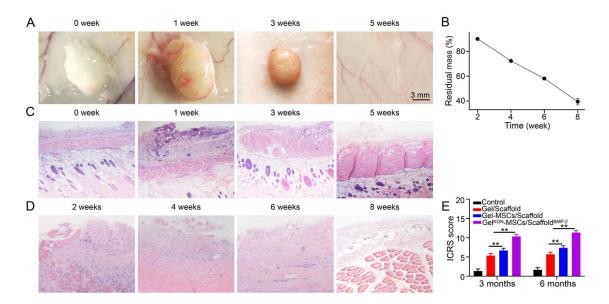


Fig. S1. In vivo degradation of mPEG-*b*-PLV thermogel **(A)** and PLGA/HA scaffold **(B)**. In vivo biocompatibility of mPEG-*b*-PLV thermogel **(C)** and PLGA/HA scaffold **(D)**. ICRS histological score **(E)**.

Table S1. ICRS Macroscopic Evaluation of Neocartilage.

Cartilage repair assessment ICRS	Points
Degree of defect repair	
In level with surrounding cartilage	4
75% repair of defect depth	3
50% repair of defect depth	2
25% repair of defect depth	1
0% repair of defect depth	0
Integration to border zone	
Complete integration with surrounding cartilage	4
Demarcating border <1 mm	3
3/4th of graft integrated, 1/4th with a notable border > 1 mm	2
width	
1/2 of graft integrated with surrounding cartilage, 1/2 with a	1
notable border >1 mm	
From no contact to 1/4th of graft integrated with surrounding	0
cartilage	
Macroscopic appearance	
Intact smooth surface	4
Fibrillated surface	3
Small, scattered fissures, or cracs	2
Several, small, or few but large fissures	1
Total degeneration of grafted area	0
Overall repair assessment	
Grade I: normal	12
Grade II: nearly normal	11-8
Grade III: abnormal	7–4
Grade IV: severely abnormal	3-1

Table S2. ICRS Histological Grading Scale for Cartilage Regeneration.

Description	Points	
Cell morphology		
Hyaline cartilage	0	
Mostly hyaline cartilage	1	
Mostly fibrocartilage	2	
Mostly non-cartilage	3	
Non-cartilage only	4	
Matrix staining (metachromasia)		
Normal (compared with host adjacent cartilage)	0	
Slightly reduced	1	
Markedly reduced	2	
No metachromatic stain	3	
Surface regularity		
Smooth (>3/4)	0	
Moderate (>1/2-3/4)	1	
Irregular (1/4–1/2)	2	
Severely irregular (<1/4)	3	
Thickness of cartilage		
>2/3	0	
1/3–2/3	1	
<1/3	2	
Integration of donor with host adjacent cartilage		
Both edges integrated	0	
One edge integrated	1	
Neither edge integrated	2	
Maximum total	14	

Table S3. Sequences of Primers Used for Real-time PCR.

Gene	Forward primer (5'–3')	Reverse primer (5'–3')
COL2A	TCCTGTGCGACGACATAATTCT	CCTTTGGTCCTGGTTTCC
1		
FAM	CCTCGTCATCTTTATCTTC	CTGCGTTGTGCCTATCGT
ACAN	TCCACCATTCGGCATAAC	TACGGATTCTACCAAAGTTCT
		G
SOX9	CTTCAACCTCCCGCACTAC	TTGCCTTCAGCTTCAATGTC
COL1A	CACGGACTTTGGAGATTTGT	CACTCGGGTCTTCAAGCA
1		
ALPL	AGGCTTCTTCCTGCTGGTG	GGGTTTCTTGTCCGTGTCG
SPP1	GCTAAACCCTGACCCATCT	CGTCGGATTCATTGGAGT
SPARC	AAACCGAAGAGGAAGTAGTGG	TTGTGGCAAAGAAGTGGC