## Enrichment of CD146<sup>+</sup> adipose-derived stem cells in combination with articular cartilage extracellular matrix scaffold promotes

cartilage regeneration

Supplementary materials



Figure S1. Isotype control of flow cytometry



**Figure S2.** Quantification of mRNA levels of *CD40*, *CD80*, *HGF*, *ID01*, *IL10*, and *PGE2* in  $CD146^+$  cells and hADSCs by qRT-PCR. Data represent mean  $\pm$ SD from three independent experiments.



Figure S3. X-ray images of rabbit knees at 3 months and 6 months after surgery.



Figure S4. ICRS scoring system for macroscopic evaluation of cartilage repair after 3 and 6 months. The results represent the mean  $\pm$  standard error of the mean. \*P < 0.05, \*\*P < 0.01.



**Figure S5.** Histological analysis of cartilage defect after 3 and 6 months by H&E staining. The black solid arrows denote the repair interface. The red solid arrows denote the depth of the repaired cartilage. HC, host cartilage; RC, repaired cartilage. Scale bar: 200 µm.



**Figure S6.** Histological analysis of cartilage defect after 3 and 6 months by Safranin O staining. The black solid arrows denote the repair interface. The red solid arrows denote the depth of the repaired cartilage. HC, host cartilage; RC, repaired cartilage. Scale bar: 200 μm



**Figure S7.** Histological analysis of cartilage defect after 3 and 6 months by Toluidine Blue staining. The black solid arrows denote the repair interface. The red solid arrows denote the depth of the repaired cartilage. HC, host cartilage; RC, repaired cartilage. Scale bar: 200 µm.



**Figure S8.** Histological analysis of cartilage defect after 3 and 6 months by Sirius Red staining. The black solid arrows denote the repair interface. The red solid arrows denote the depth of the repaired cartilage. HC, host cartilage; RC, repaired cartilage. Scale bar: 200 μm.

Gene	Forward primer	Reverse primer
CD80	GCTGGCTGGTCTTTCTCACT	GTGCCAGCTCTTCAACAGAA
CD40	CTCACCTCGCTATGGTTCGT	GGCACAAAGAACAGCACTGA
IdO1	GATGAAGAAGTGGGCTTTGC	CAGGCAGATGTTTAGCAATGA
PGE2	CCTTCTTCGAAAGTTTTGCC	ATGTGCAGTTGCCCTCTGTA
HGF	CGCTGGGAGTACTGTGCAAT	CCCTGTAGCCTTCTCCTTGA
IL-10	GTGGAGCAGGTGAAGAATGC	GCCACCCTGATGTCTCAGTT
ACTIN	ACCTTCTACAATGAGCTGCG	CCTGGATAGCAACGTACATGG
GAPDH	CAAGAAGGTGGTGAAGCAGG	GGTGTCGCTGTTGAAGTCAG

## Table S1. List of Primers Used for RT-PCR Analysis

Criteria	Findings	Scores
Repaired cartilage signal	Isointense	2
	Hyperintense	1
	Hypointense	0
Repaired lesion morphologic features	Flush	2
	Proud	1
	Depressed	0
Repaired cartilage fill	Good (67%-100%)	2
	Moderate (34%-66%)	1
	Poor (0%-33%)	0
Peripheral repaired cartilage integration	No gap	2
	Small (gap of $\leq 2$ mm)	1
	Large (gap of >2 mm)	0
Subchondral edema	None	3
	Mild (<1 cm2)	2
	Moderate (1-3 cm2)	1
	Severe (>3 cm2)	0
Osseous overgrowth	No	1
	Yes	0

 Table S2. Score System for MRI Evaluation

Evaluation of Cartilage Repair			
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	4		
surrounding cartilage			
Demarcating border <1 mm	3		
3/4th of graft integrated, 1/4th	2		
with a notable border $>1$ mm width			
1/2 of graft integrated with surrounding	1		
cartilage, $1/2$ with a notable border >1 mm			
From no contact to 1/4th of graft integrated	0		
with surrounding cartilage			
Macroscopic appearance			
Intact smooth surface	4		
Fibrillated surface	3		
Small, scattered fissures or cracks	2		
Several, small or few but large fissures	1		
Total degeneration of grafted area	0		
Overall repair assessment			
Grade I: normal			
Grade II: nearly normal			
Grade III: abnormal			
Grade IV: severely abnormal	3–1		

 Table S3. International Cartilage Repair Society Macroscopic

ICRS, International Cartilage Repair Society.

Assessment Seale for Carthage Repair	
Feature	Points
I. Surface	
Smooth/continuous	3
Discontinuities/irregularities	0
II. Matrix	
Hyaline	3
Mixture: hyaline/fibrocartilage	2
Fibrocartilage	1
Fibrous tissue	0
III. Cell distribution	
Columnar	3
Mixed/columnar clusters	2
Clusters	1
Individual cells/disorganized	0
IV. Cell population viability	
Predominantly viable	3
Partially viable	1
<10% viable	0
V. Subchondral bone	
Normal	3
Increased remodeling	2
Bone necrosis/granulation tissue	1
Detached/fracture/callus at base	0
VI. Cartilage mineralization	
(calcified cartilage)	
Normal	3
Abnormal/inappropriate location	0

 Table S4. International Cartilage Repair Society Visual Histological

Assessment Scale for Cartilage Repair