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Fig. S5 Transcriptome analysis of MSC and MSC chondrogenesis (3 days) groups. MSC group:
MSC cultured for 3 days. MSC chondrogenesis (3 days) group: MSC cultured for 3 days with
chondrogenic induction medium. (A) Heat map analysis of differential genes in MSC and MSC
chondrogenesis (3 days) groups. (B) KEGG enrichment analysis of differential genes in MSC and
MSC chondrogenesis (3 days) groups.

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Fig. S6 Transcriptome analysis of MSC and MSC+CyA chondrogenesis (3 days) groups. MSC group: MSC cultured for 3 days. MSC+CyA chondrogenesis (3 days) group: MSC cultured for 3 days with chondrogenic induction medium and 0.5 mg/mL CyA. (A) Heat map analysis of differential genes in MSC and MSC+CyA chondrogenesis (3 days) groups. (B) KEGG enrichment analysis of differential genes in MSC and MSC+CyA chondrogenesis (3 days) groups.



11 The significance is represented as p < 0.05 (*), p < 0.01 (**), p < 0.001 (***) and no significant





2 Fig. S9 Alcian Blue staining images of hBMSCs after 14 days in different groups.



9 Fig. S10 Relative mRNA expression of Acan, COL2A1, and SOX9 in hBMSCs after 14 days. The

10 significance is represented as p < 0.05 (*), p < 0.01 (**), p < 0.001 (***) and no significant

- 11 difference (ns). Error bar represents for SD.





Fig. S11 Western blot bands and quantitative analysis of chondrogenic marker expression in hBMSCs after 14 days. The significance is represented as p < 0.05 (*), p < 0.01 (**), p < 0.001(***) and no significant difference (ns). Error bar represents for SD.

Table S1 qPCR primer sequences

Gene name	Primers
Human-GAPDH-F	CAGGAGGCATTGCTGATGAT
Human-GAPDH-R	GAAGGCTGGGGGCTCATTT
Human-Acan-F	GTGCCTATCAGGACAAGGTCT
Human-Acan-R	GATGCCTTTCACCACGACTTC
Human-COL2A-F	CCTGGCAAAGATGGTGAGACAG
Human-COL2A-R	CCTGGTTTTCCACCTTCACCTG
Human-SOX9-F	AGGAAGCTCGCGGACCAGTAC
Human-SOX9-R	GGTGGTCCTTCTTGTGCTGCAC
Human-MMP13-F	CCTTGATGCCATTACCAGTCTCC
Human-MMP13-R	AAACAGCTCCGCATCAACCTGC