

## Supporting Information

# Concentrated Polymer Brush-Modified Cellulose Nanofibers Promotes Chondrogenic Differentiation of Human Mesenchymal Stem Cells by Controlling Self-Assembly

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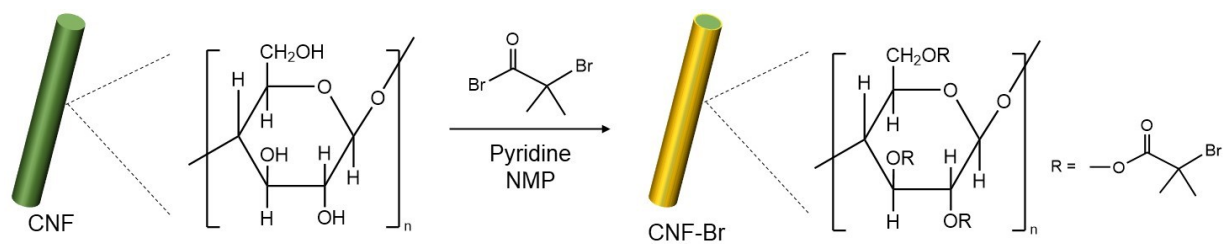
**Table S1.** Primers used in qPCR analysis

Gene	Forward primer (sequence 5' to 3')	Reverse primer (sequence 5' to 3')
GAPDH	AAGGTGAAGGTCGGAGTCAAC	GGGGTCATTGATGGCAACAATA
Collagen I	CGGCTCCTGCTCCTCTTAG	CACACGTCTCGGTCATGGTA
Collagen II	GGCAATAGCAGGTTACGTACA	CGATAACAGTCTTGCCCCACTT
Aggrecan	TCGAGGACAGCGAGGCC	TCGAGGGTGTAGCGTGTAGAGA
SOX 9	GTACCCGCACTTGCACAAC	GTAATCCGGGTGGTCCTTCT
COMP	AGCAGATGGAGCAAACGTATTG	ACAGCCTTGAGTTGGATGCC

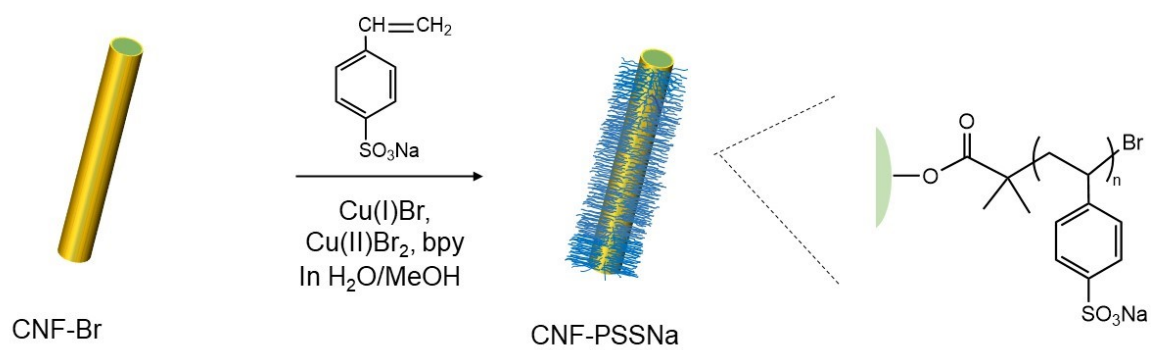
**Table S2.** Characteristics of CNF with PSSNa brush used in this work.

Entry	$M_n$	$M_w/M_n$	$M_{n,c}^a$	$\sigma^b$ (chains/nm <sup>2</sup> )	$\sigma^{*c}$	Brush Type
1	9200	1.1	11100	0.7	0.9	CPB
2	9800	1.2	14800	0.7	0.9	CPB

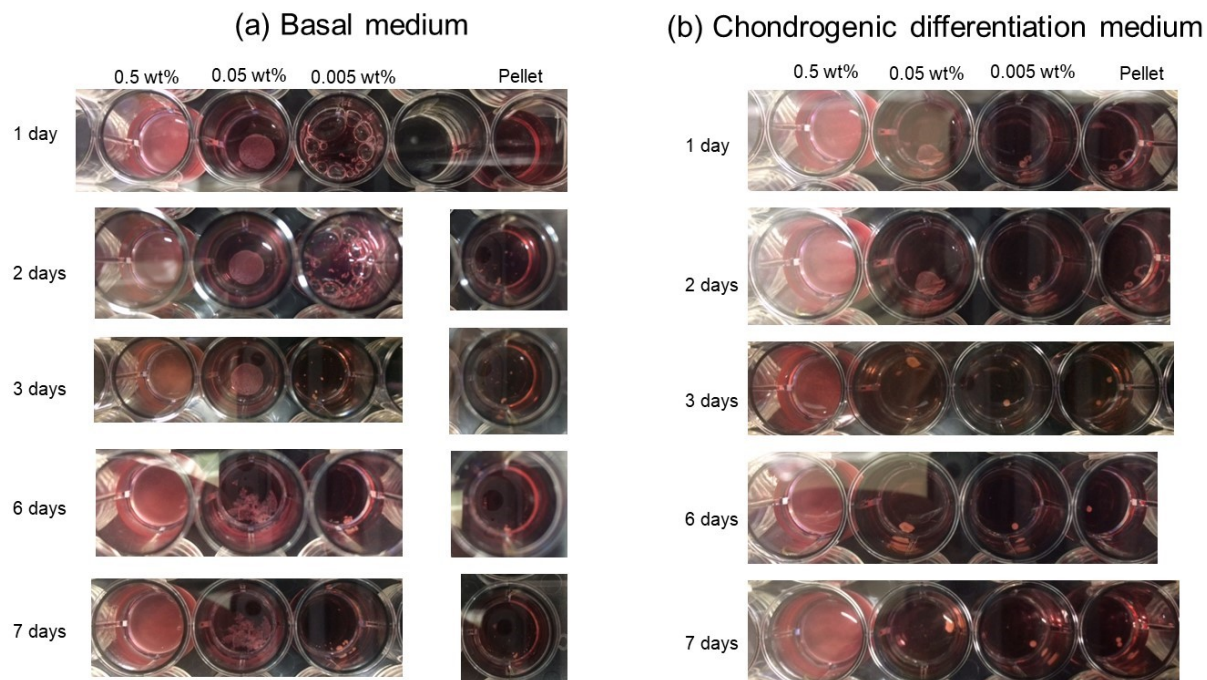
<sup>a</sup>Theoretical molecular weight calculated by the equation,  $M_{n,c} = [SSNa]_0/[EBIB]_0 \times \text{conversion} \times \text{molecular weight of SSNa}$ . <sup>b</sup>Graft density of PSSNa estimated by FT-IR measurement. <sup>c</sup>Dimensionless graft density. When  $\sigma^* > 0.1$ , the brush can be categorized into CPB.



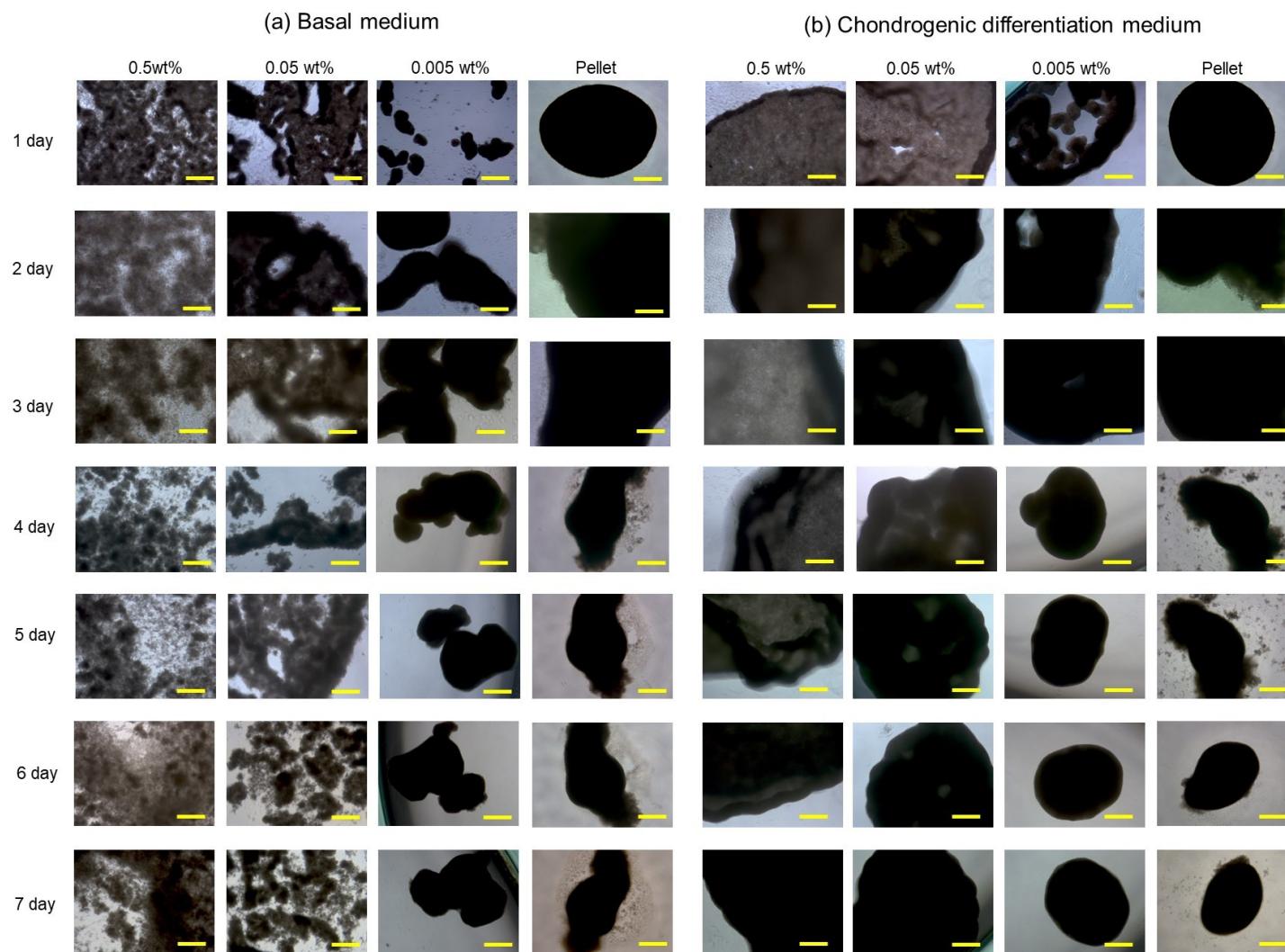
**Figure S1.** Preparation of CNF-Br by esterification.



**Figure S2.** SI-ATRP of SSNa from CNF-Br.

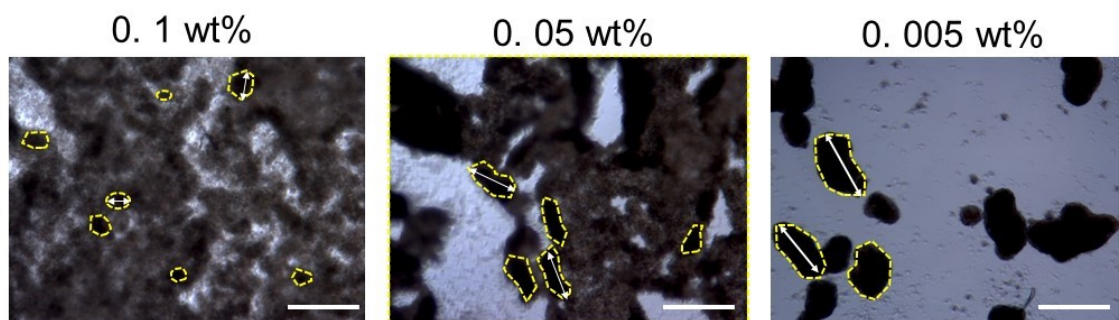


**Figure S3.** Photos of flocs and pellets (control).  $[\text{CNF-CPB}]_0 = 0.5, 0.05$  and  $0.005$  wt%.  $[\text{hMSC}]_0 = 5 \times 10^5$  cells/well. Cell culture using (a) basal medium and (b) chondrogenic differentiation medium. The scale bar =  $500 \mu\text{m}$ .

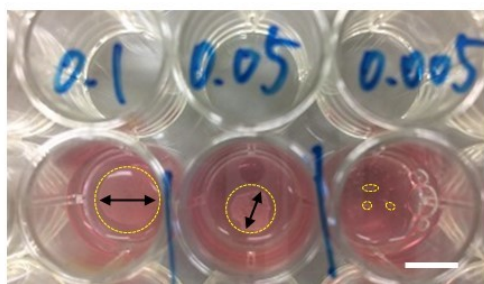


**Figure S4.** Phase contrast micrograms of flocs and pellets.  $[\text{CNF-CPB}]_0 = 0.5, 0.05$  and  $0.005$  wt%.  $[\text{hMSC}]_0 = 5 \times 10^5$  cells/well. Cell culture using (a) basal medium and (b) chondrogenic differentiation medium. The scale bar =  $500 \mu\text{m}$ .

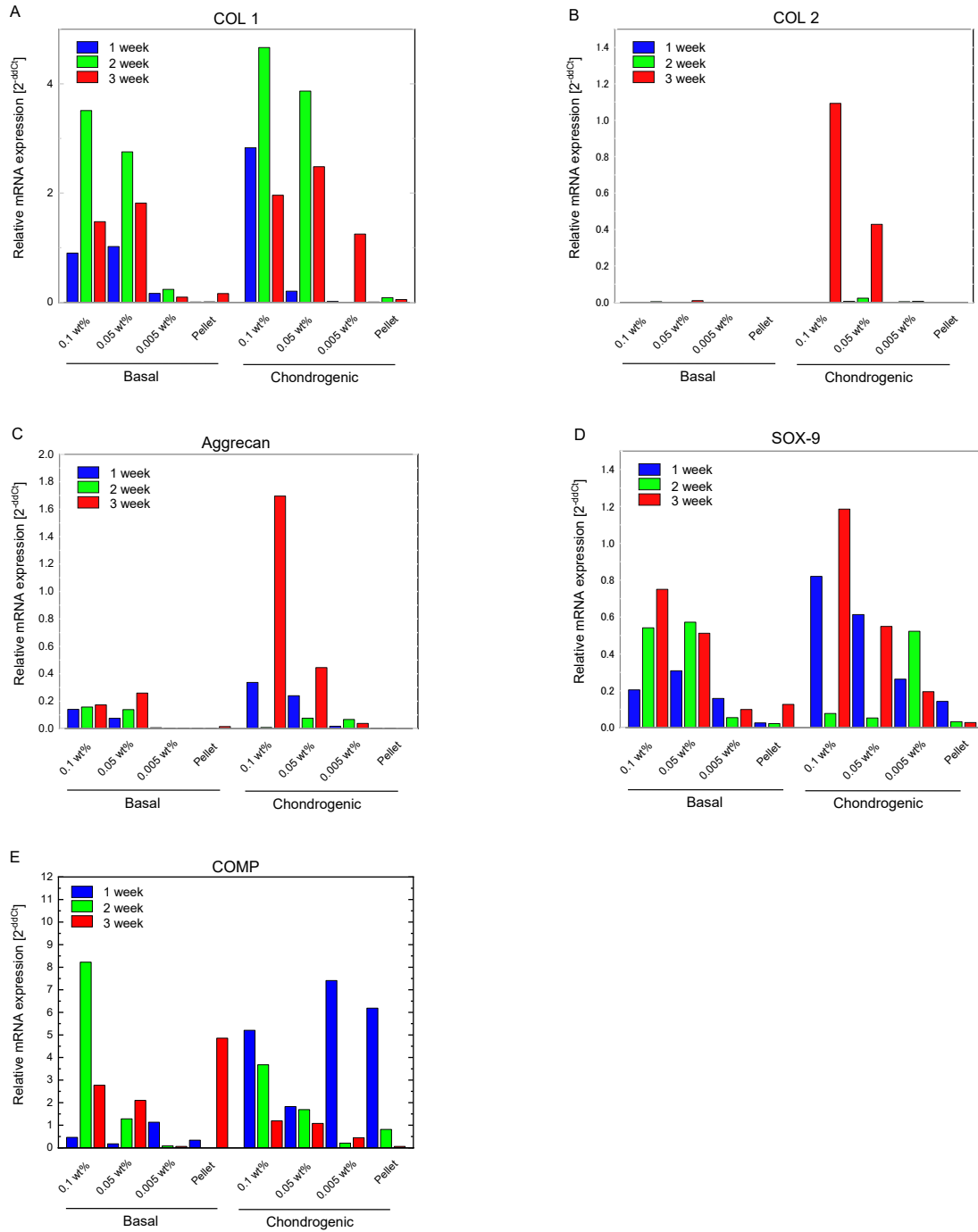
(a) Basal medium (Day 1)



(b) Chondrogenic differentiation medium (Day 1)



**Figure S5.** Examples of measuring floc sizes using the longest inner diameters (indicated by arrows). Representative flocs are indicated by dotted yellow circles. Scale bar = (a) 500  $\mu\text{m}$  and (b) 10 mm.



**Figure S6.** RT-qPCR measurements of chondrogenesis of hMSC/CNF-CPBs and pellet culture (control). (A)-(E) The fold-changes of target gene (COL 1, COL 2, Aggrecan, SOX-9, and COMP) expression.