Title

Direct conversion of human fibroblasts into hepatocyte-like cells by ATF5, PROX1, FOXA2, FOXA3, and HNF4A transduction

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Figure S1 Induction of hepatocyte-like cells from human fetal fibroblasts (A) The procedure for conversion of human fetal fibroblasts (MRC5) cells into hiHeps is presented schematically. MRC5 cells were transduced with hepatic transcription factor-expressing lentivirus vectors (LV-TFs) for 12 hr, and cultured until day 28. (**B**) MRC5 cells were transduced with 0, 100, 500, or 1,000 VP/cell of LV-Venus (modified

green fluorescent protein) for 12 hr, and cultured until day 3. The percentage of Venus

(green)-positive cells was examined by FACS.



Figure S2 Generation of hiHeps by HNF4A transduction

MRC5 cells were transduced with LV-5TFs or LV-HNF4A for 12 hr, and cultured until day 28. (A) The ALB secretion capacity in LV-5TFs- and LV-HNF4A-transduced cells, PHH 48hr, and PHH 4hr was examined by ELISA. **p < 0.01 (vs LV-5TFs). (B) The CYP1A2 and CYP3A4 activities were examined in LV-5TFs- and LV-HNF4A-transduced cells, PHH 48hr, and PHH 4hr. The CYP1A2 and CYP3A4 activity levels in PHH 48hr were taken as 1.0. (C) The percentage of ASGR1-positive cells in LV-5TFs- or LV-HNF4A-transduced cells was examined by FACS. *p < 0.05 (vs LV-5TFs). All data are represented as means ± SD (n=3).



Figure S3 The expression ratios of *ALB/AFP* and *CYP3A4/CYP3A7*

MRC5 cells were transduced with LV-5TFs for 12 hr, and cultured until day 28. The hepatic gene (*ALB* and *CYP3A4*) and fetal-specific hepatic gene (*AFP* and *CYP3A7*) expression levels in iPS-Hepa, hiHeps, PHH 48hr, and PHH 0hr were then measured by real-time RT-PCR. The *ALB/AFP* and *CYP3A4/CYP3A7* ratios were calculated. **p < 0.01 (vs hiHeps).



Figure S4 Immunostaining analysis of fibroblast and hepatic markers MRC5 cells were transduced with LV-5TFs or LV-control for 12 hr, and cultured until day 28. (**A**, **B**) The percentages of fibroblast marker (THY-1)- and ASGR1-positive cells were examined by FACS analysis. Representative 2D plot data are shown in **figure S4A**. Quantitative data of FACS analysis are shown in **figure S4B**. All data are represented as means \pm SD (*n*=3).



Figure S5 Exogenous and total *ATF5*, *PROX1*, *FOXA2*, *FOXA3*, and *HNF4A* expression profiles during the hepatic reprogramming

MRC5 cells were transduced with LV-5TFs for 12 hr, and cultured until day 28. (**A**) The exogenous gene (*ATF5*, *PROX1*, *FOXA2*, *FOXA3*, and *HNF4A*) expression levels were

measured by qRT-PCR. The gene expression levels in LV-5TF-transduced cells (day 4) were taken as 1.0. (**B**) The total (endogenous and exogenous) gene (*ATF5*, *PROX1*, *FOXA2*, *FOXA3*, and *HNF4A*) expression levels were measured by qRT-PCR. The gene expression levels in LV-5TF-transduced cells (day 0) were taken as 1.0. All data are represented as means \pm SD (*n*=3).



Figure S6 Comparison of hepatic direct reprogramming protocols

The hiHeps were generated by transducing MRC-5 cells with LV-5TFs (ATF5, PROX1, FOXA2, FOXA3, and HNF4A). The hiHeps were also generated by transducing MRC-5 cells with LV-3TFs (FOXA3, HNF1A, and HNF4A). The percentage of ASGR1-positive cells in hiHeps was examined by FACS. *p < 0.05 (vs LV-3TFs). All data are represented as means ± SD (n=3).

Gene	Genebank Number	Primers (forward/reverse; 5' to 3')	
ATF5	NM_001193646	TGCTCGAGCCACCATGTCACTCCTGGCGACCCTG	
		AAGCGGCCGCCTAGCAGCTACGGGTCCTCTG	
PROX1	NM_ 001270616	AACGAATTCGTGATGCCTGACCATGACAGCA	
		TGTGCGGCCGCCTACTCATGAAGCAGCTCTTGTAGGCA	
CEBPA	NM_004364	TCTCTCGAGCCCATGGAGTCGGCCGACTT	
		AGCGGATCCTCACGCGCAGTTGCCCAT	
FOXA2	NM_153675	GCGGAATTCAGTATGCTGGGAGCGGTGAA	
		GTCGGATCCTTAAGAGGAGTTCATAATGGGCCGGGA	
FOXA3	NM_004497	TAACTCGAGGGGATGCTGGGCTCAGTGAA	
		CCCGCGGCCGCCTAGGATGCATTAAGCAAAGAGCGGG	
HNF1A	NM_000545	GGCGAATTCGCCATGGTTTCTAAACTGAGCCAG	
		AGGGCGGCCGCTTACTGGGAGGAAGAGGCCATCT	
HNF4A	NM_178849	GAGGAATTCAGAATGCGACTCTCCAAAACCCTC	
		CCAGCGGCCGCCTAGATAACTTCCTGCTTGGTGATGGT	
HNF6	NM_004498	GCCCTCGAGACGATGAACGCGCAGCTGACCAT	
		GTTGCGGCCGCTCATGCTTTGGTACAAGTGCTTGATGAAGAAG	
GATA4	NM_002052	CTCCTCGAGACCATGTATCAGAGCTTGGCCATGG	
		GAGGAATTCTTACGCAGTGATTATGTCCCCGTGAC	

Table S1Primers for cloning

Primers (forward/reverse; 5' to 3')		
GCACAGAATCCTTGGTGAACAG/ATGGAAGGTGAATGTTTTCAGCA		
ACTGTCAACTTCGGGGGACAC/CATGCCTAAACGCTTCATCA		
AAGTCGCCTCGAAGATACACA/AAGGAGAGAACACTGCTCGTG		
GAGGGCCAAGACGAAGACATC/CAGATCACGTCATCGCACAAC		
CTAGTGGACCAGAGCCTTCG/TGGAGTGCACACGTGTAGGT		
AGAAGGCCTCAGCACCTAC/GGCCTGATTGTTCCAGGATT		
CTTGAATCCCGAATGGAAAGGG/GTGTATATCCCAGGGTGATCCTC		
GGCAGCTACAGCATGATGATGCAGGAGC/CTGGTCATGGAGTTGTACTGCAGG		
TGGGACCCGAACTTTCCA/GGCCACATCCAGGACTAGTTTC		
AAGGTCGCCTCAAAGAGACA/TGCACTTTCTGCTGGACATC		
GACCGCAAGCAAAAGAAGAG/GGCCTTGTAAACCTCGATGA		
TTGACATTGGAGTGAAAAGGACG/TGCTCAGAACCTTGGGGGATTC		
CCAAGAAGTCGGTGGACAAG/AGGCGGTCATTGTCACTGGT		
GCGACCCCAAGACCTACAG/GGTTCTGCCGGTAGAAGGG		
TCATGTAGGAGTTGAGGGGGG/GAAGATGGAGGCCCATGAC		
TACACCACTCTGGCAGCCACACT/CGGTGGGTACATTGGTGACAGAAC		
CGTCATCGTTGCCAACACAAT/GGGCCACTCACACATCTGTC		
AACCCTGGAGCAAACTCAAA/AAAAAGCCCAGGTTGGTCTT		
CATCAAGACGGAGCCTGGCC/TGACTGTCGGCCAAGACCAG		
TCATCGTCTGCTCCTCT/AGGTGTCATCAGCAGCCTTT		
CAATCAGGTGGTGGTGTCAG/GCTCCTGGACTGTTTTCTGC		
GGACAGAGACGACAAGCACA/CATCTGTGTAGGGCATGTGG		
ACTTGGAGCTGGGACAGAGA/CATCTGTGTAGGGCATGTGG		
CTTTCGCCCCAACGGTCTC/TTTTGGAAGCGTAGGACCTTG		
ACCCGAGACACCATTTTCAG/TCCAGCACACACTCGTTTTC		
ATCTTGGTCTGTGGCTGCTC/AGAAGGTGGAGCAGGTGGT		
TAAGTGGCTACCCCAAAACG/GCTTTGCATTGTCCATCTGA		
GGTGGTCTCCTCTGACTTCAACA/GTGGTCGTTGAGGGCAATG		

 Table S2
 Primers for real-time RT-PCR analysis

Antigen	Туре	Company
ALB	goat	Bethyl Laboratories
AAT	rabbit	Dako
Alexa Fluor 488 anti-goat IgG	donkey	Thermo Fisher Scientific
Alexa Fluor 594 anti-rabbit IgG	donkey	Thermo Fisher Scientific

 Table S3
 Antibodies for immunohistochemistry