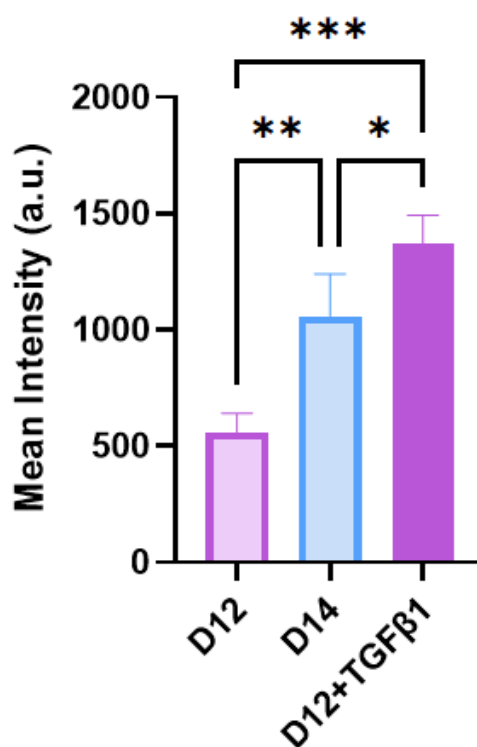


RESEARCH ARTICLE

Digital light processing-printed macroencapsulated human liver organoids preserve hepatic stellate cell quiescence for transplantation in immunocompetent mice

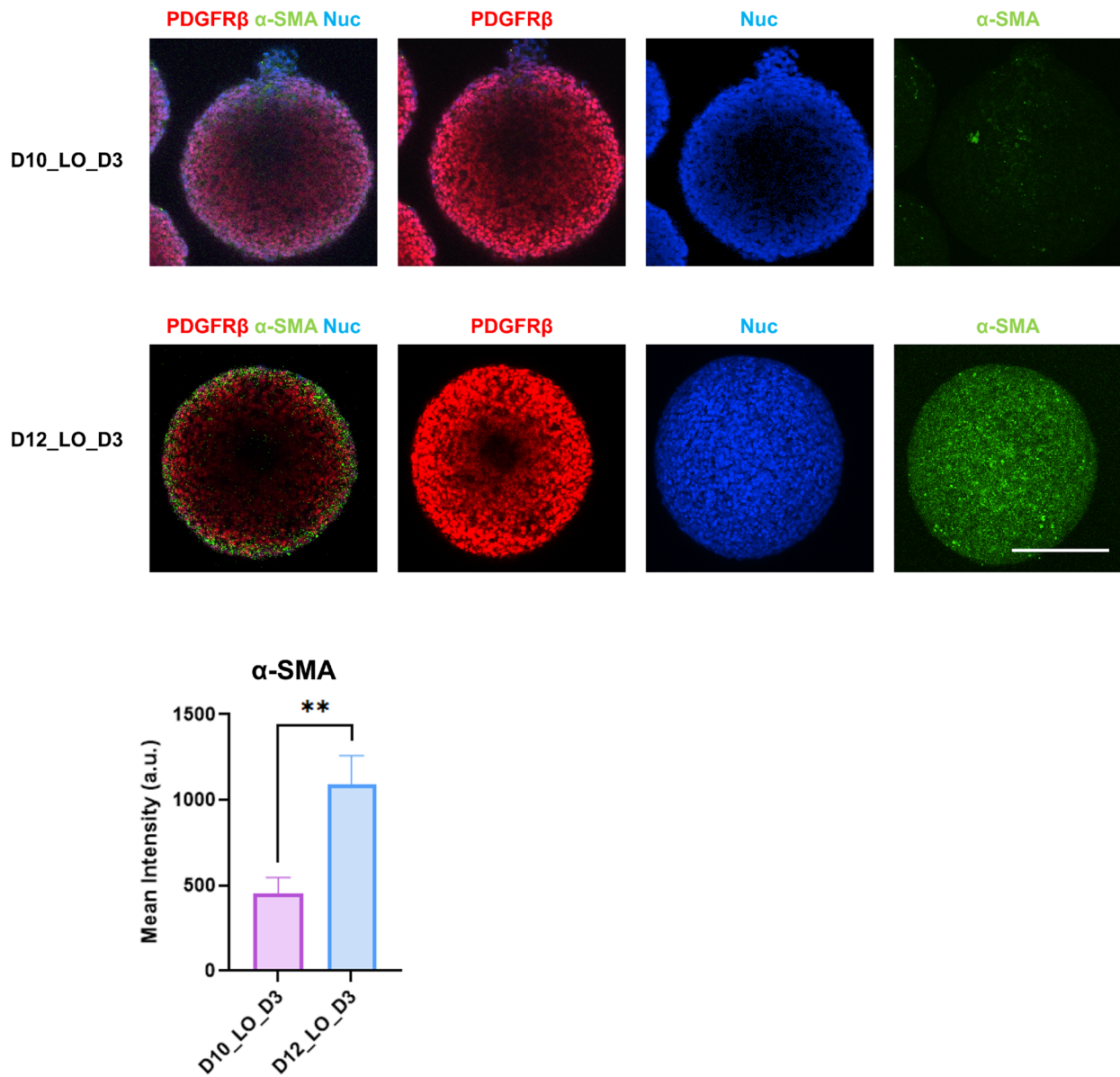
Supplementary Files



**Figure S1.** Quantitative analysis of immunofluorescence intensity of  $\alpha$ -SMA staining for HSCs at different days

Notes: Data are mean  $\pm$  SD. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

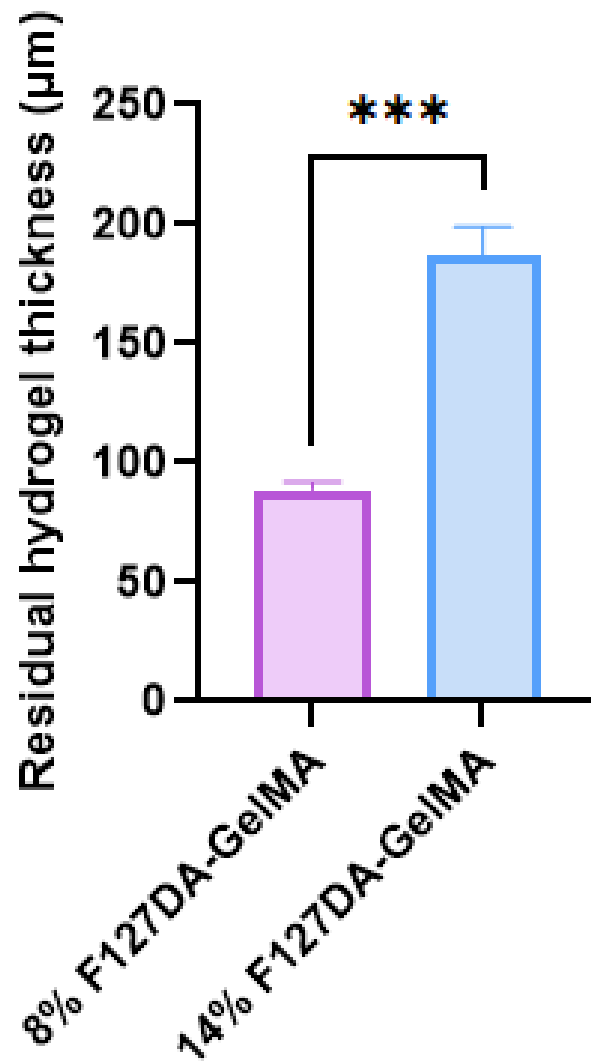
Abbreviations:  $\alpha$ -SMA: Alpha-smooth muscle actin; HSC: Hepatic stellate cell; TGF- $\beta$ 1: Transforming growth factor-beta 1.



**Figure S2.** Quantitative analysis of immunofluorescence intensity of α-SMA staining for HSCs at different days. (A) Single immunofluorescence images of Figure 2B. Scale bars: 50 μm; magnifications: 40×. (B) α-SMA fluorescence quantification.

Notes: Data are mean ± SD. \*\*p < 0.01.

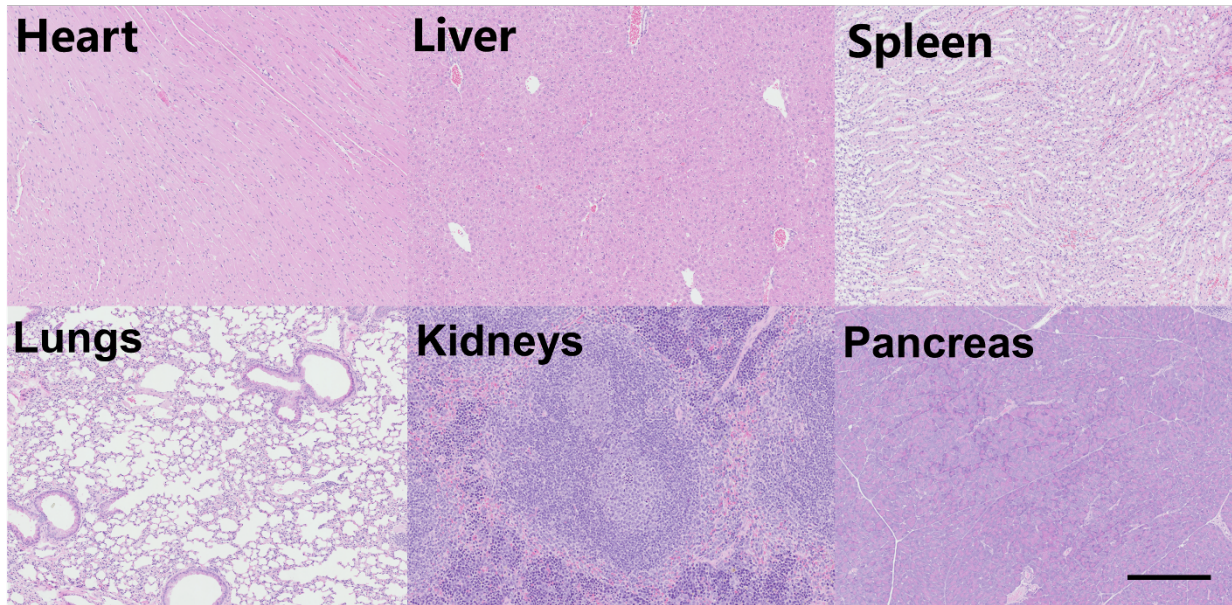
Abbreviations: α-SMA: Alpha-smooth muscle actin; HSC: Hepatic stellate cell; LO: Liver organoid; Nuc: Nucleus; PDGFRβ: Platelet-derived growth factor receptor beta.



**Figure S3.** Quantitative analysis of the residual hydrogel scaffolds after in vivo implantation

Notes: Data are mean  $\pm$  SD. \*\*\* $p < 0.001$ .

Abbreviations: GelMA: Gelatin methacryloyl; F127DA: Pluronic F127 diacrylate.



**Figure S4.** Images of the representative histological HE staining of major tissues after in vivo implantation of F127DA. Scale bars: 200  $\mu\text{m}$ ; magnifications: 4 $\times$ .

**Table S1. Primer list**

Symbol	Forward sequence (5'-3')	Reverse sequence (5'-3')
<i>KDR</i>	GGAACCTCACTATCCGCAGAGT	CCAAGTTCGTCTTTTCTGGGG
<i>ACTA2</i>	CTATGCCTCTGGACGCACAAC	CAGATCCAGACGCATGATGGCA
<i>VIM</i>	AGGCAAAGCAGGAGTCCACTGA	ATCTGGCGTTCCAGGGACTCAT
<i>PDGFRB</i>	TGCAGACATCGAGTCCTCCAAC	GCTTAGCACTGGAGACTCGTTG
<i>NGFR</i>	CCTCATCCCTGTCTATTGCTCC	GTTGGCTCCTTGCTTGTCTGC
<i>ALCAM</i>	TCCAGAACACGATGAGGCAGAC	GTAGACGACACCAGCAACAAGG
<i>RELN</i>	GTCTACCTTCCACTCTCCACCA	GTCCAGCATCACAATCCCTCG
<i>HGF</i>	GAGAGTTGGGTTCTTACTGCACG	CTCATCTCCTCTCCGTGGACA
<i>COL1A1</i>	GATTCCCTGGACCTAAAGGTGC	AGCCTCTCCATCTTTGCCAGCA
<i>PPARG</i>	AGCCTGCGAAAGCCTTTTGGTG	GGCTTCACATTCAGCAAACCTGG
<i>LOXL2</i>	TGACTGCAAGCACACGGAGGAT	TCCGAATGTCCTCCACCTGGAT
<i>NCAM1</i>	CATCACCTGGAGGACTTCTACC	CAGTGTACTGGATGCTCTTCAGG
<i>TIMP1</i>	GGAGAGTGTCTGCGGATACTTC	GCAGGTAGTGATGTGCAAGAGTC
<i>GFAP</i>	CTGGAGAGGAAGATTGAGTCGC	ACGTCAAGCTCCACATGGACCT
<i>PCDH7</i>	GAGGAGTCAGAAACACCAAGCAG	TCAGGGTACATCTGGAAGAGG
<i>GAPDH</i>	GTCTCCTCTGACTTCAACAGCG	ACCACCTGTTGCTGTAGCCAA
<i>P75NTR</i>	CCTCATCCCTGTCTATTGCTCC	GTTGGCTCCTTGCTTGTCTGC