

Corresponding author(s):	Sang Jin Lee
Last updated by author(s):	Jan 25, 2020

Reporting Summary

X Life sciences

Nature Research wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Research policies, see <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

Statistics			
For all statistical analyse	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.		
n/a Confirmed			
The exact sam	ple size (n) for each experimental group/condition, given as a discrete number and unit of measurement		
A statement o	n whether measurements were taken from distinct samples or whether the same sample was measured repeatedly		
The statistical Only common to	test(s) used AND whether they are one- or two-sided sets should be described solely by name; describe more complex techniques in the Methods section.		
A description	of all covariates tested		
A description	of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons		
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)		
For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.			
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings			
For hierarchic	al and complex designs, identification of the appropriate level for tests and full reporting of outcomes		
Estimates of e	ffect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated		
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.		
Software and c	ode		
Policy information abou	ıt <u>availability of computer code</u>		
Data collection	FV10-ASW 04.02., Image Pro 6.3., LAS-AF 3.1.8976.3, cellSens Dimension 1.18, MI-RAT 2.74		
Data analysis	Image J 1.8.0_112 (https://imagej.nih.gov/ij/download.html), SPSS software 19, Origin pro 8.5, GraphPad Prism 5		
	om algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.		
Data			
- Accession codes, uni - A list of figures that l	It <u>availability of data</u> nclude a <u>data availability statement</u> . This statement should provide the following information, where applicable: que identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability		
The data had been includ corresponding author on	ed in the manuscript or its supplementary material. The additional data generated or analysed during this study are available from the request.		
Field-speci	fic reporting		

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Ecological, evolutionary & environmental sciences

Behavioural & social sciences

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sam	n	C	17	c
Jaili	v	0	14	ζ

No sample size calculation was performed in this study. Sample size was determined to be adequate based on the magnitude and consistency of measurable differences between groups. In our previous study (Kim, J.H. et al. 3D Bioprinted Human Skeletal Muscle Constructs for Muscle Function Restoration. Scientific reports 8, 12307 (2018) - https://www.nature.com/articles/s41598-018-29968-5) and this study, the statistically significant differences among groups had showed with the determined sample size. Detail of sample size of all experiments were provided in the text and figure legends.

Data exclusions

On principle, data were only excluded for failed experiments, reasons for microbial contamination of in vitro experiments.

Replication

The experimental findings (Fig. 3h-3i, Supplementary Fig. S2, S4 and S5) were qualitatively reproduced three times. Replicate experiments were successful.

Randomization

For the in vivo study, recipient rats were allocated randomly for the engineered tissue implantation of Printed-MPC, Printed-MPC+NSC, or non-treated (defect only without implantation).

Blinding

The investigators were blinded to group allocation during data collection and analysis.

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials &	experimental	systems
-------------	--------------	---------

___ Methods

n/a	Involved in the study	
	Antibodies	
	Eukaryotic cell lines	
\boxtimes	Palaeontology	
	Animals and other organisms	
\boxtimes	Human research participants	
\boxtimes	Clinical data	

n/a	Involved in the study

\boxtimes	ChIP-seq
X	Flow cytor

MRI-based neuroimaging

Antibodies

Antibodies used

mouse anti-MF-20 antibody (MF-20, 1 μ g/ml, Developmental Studies Hybridoma Bank), mouse anti-myoD (MA1-41017, 1:200 dilution, Thermo Scientific Inc.), rabbit anti-myogenin (ab124800, 1:200 dilution, Abcam), rabbit anti- β IIIT (ab18207, 1:100 dilution, Abcam), rabbit anti-NF (N4142, 1:80 dilution, MilliporeSigma), rabbit anti-GFAP (ab7260, 1:100 dilution, Abcam), rat anti-AChR antibody (ab24719, 1:100 dilution, Abcam), rabbit anti-VWF (A0082, 1:400 dilution, Dako) and mouse anti- α -SMA (sc-53015, 1:50 dilution, Santa Cruz Biotechnology), chicken anti-NF (ab4680, 1:1000 dilution, Abcam), rabbit anti-HLA (ab52922, 1:100 dilution, EMD Millipore), mouse anti-HNA (clone 3E1.3, MAB4383, 1:100 dilution, EMD Millipore), Texas Red-conjugated anti-mouse IgG (TI-2000, 1:200 dilution, Vector Labs), Texas Red-conjugated anti-rabbit IgG (TI-1000, 1:200 dilution, Vector Labs) Texas Red-conjugated anti-rabbit IgG (Al1008, 1:200 dilution, Invitrogen), Alexa 488-conjugated anti-rab IgG (Al1008, 1:200 dilution, Invitrogen), Cy5-conjugated anti-rabbit IgG (Al10523, 1:200 dilution, Invitrogen), Cy5-conjugated anti-rabbit IgG (Al10523, 1:200 dilution, Invitrogen), Cy5-conjugated anti-rabbit IgG (Al10523, 1:200 dilution, Invitrogen).

All the antibodies used for these studies are listed in the manuscript as well.

Validation

The specificity of each antibody was validated by the manufacturer, and provided on the website. mouse anti-MF-20 antibody (MF-20, https://dshb.biology.uiowa.edu/MF-20), mouse anti-myoD (MA1-41017, https://www.thermofisher.com/antibody/product/MYOD-Antibody-clone-5-8A-Monoclonal/MA1-41017), rabbit anti-myogenin (ab124800, https://www.abcam.com/myogenin-antibody-epr4789-ab124800.html), rabbit anti-βIIIT (ab18207, https://www.abcam.com/beta-iii-tubulin-antibody-ab18207.html), rabbit anti-NF (N4142, https://www.sigmaaldrich.com/catalog/product/sigma/n4142?lang=en®ion=US), rabbit anti-GFAP (ab7260, https://www.abcam.com/gfap-antibody-ab7260.html), rat anti-AChR antibody (ab24719, https://www.abcam.com/nicotinic-acetylcholine-receptor-alpha-1--3--5-antibody-210-ab24719.html), rabbit anti-VWF (A0082, https://www.agilent.com/cs/library/packageinsert/public/SSA0082IVD-US_01.pdf), mouse anti-α-SMA (sc-53015, https://datasheets.scbt.com/sc-53015.pdf), chicken anti-NF (ab4680, https://www.abcam.com/neurofilamen-heavy-polypeptide-antibody-ab4680.html), rabbit anti-HLA (ab52922, https://www.nature.com/articles/s41598-018-29968-5), mouse anti-HNA (clone 3E1.3, MAB4383, http://www.emdmillipore.com/US/en/product/Anti-Nuclei-Antibody-clone-3E1.3,MM_NF-MAB4383?bd=1), Texas Red-conjugated anti-mouse lgG (TI-2000, https://vectorlabs.com/texas-red-poat-anti-rabbit-igg-antibody.html), Texas Red-conjugated anti-rabbit lgG (TI-1000, https://www.thermofisher.com/texas-red-goat-anti-rab-igg-antibody), Alexa 488-conjugated anti-rabbit lgG (A11008, https://www.thermofisher.com/

antibody/product/Goat-anti-Rabbit-IgG-H-L-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A-11008), Alexa 488-conjugated anti-rat IgG (A11006, https://www.thermofisher.com/antibody/product/Goat-anti-Rat-IgG-H-L-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A-11006), Alexa 488-conjugated anti-chick IgG (A11039, https://www.thermofisher.com/antibody/product/ Goat-anti-Chicken-IgY-H-L-Secondary-Antibody-Polyclonal/A-11039), Cy5-conjugated anti-mouse IgG (A10524, https:// www.thermofisher.com/antibody/product/Goat-anti-Mouse-IgG-H-L-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A10524), Cy5-conjugated anti-rabbit IgG (A10523, https://www.thermofisher.com/antibody/product/Goat-anti-Rabbit-IgG-H-L-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A10523).

Eukaryotic cell lines

Policy information about cell lines

ReNCell VM were obtained from Millipore. Human muscle progenitor cells were obtained from biopsies of human gracilis Cell line source(s)

muscle (from 51- ad 62-year-old women, de-identified).

Authentication Cell lines were not additionally authenticated.

Cell lines were not additionally tested for mycoplasma. Mycoplasma contamination

Commonly misidentified lines (See ICLAC register)

No commonly misidentified cell lines were used.

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

RNU rats (male, 10-12 weeks old, Charles River Laboratory, Wilmington, MA) were used in this study. Laboratory animals

Wild animals This study did not involve wild animals.

Field-collected samples No field-collected samples were used in the study.

All animal experiments were performed in accordance with a protocol approved by the Institutional Animal Care and Use Ethics oversight

Committee (IACUC) at Wake Forest University.

Note that full information on the approval of the study protocol must also be provided in the manuscript.