

## Supplementary Materials for

### **Biochemical and structural cues of 3D-printed matrix synergistically direct MSC differentiation for functional sweat gland regeneration**

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Published 4 March 2020, *Sci. Adv.* **6**, eaaz1094 (2020)

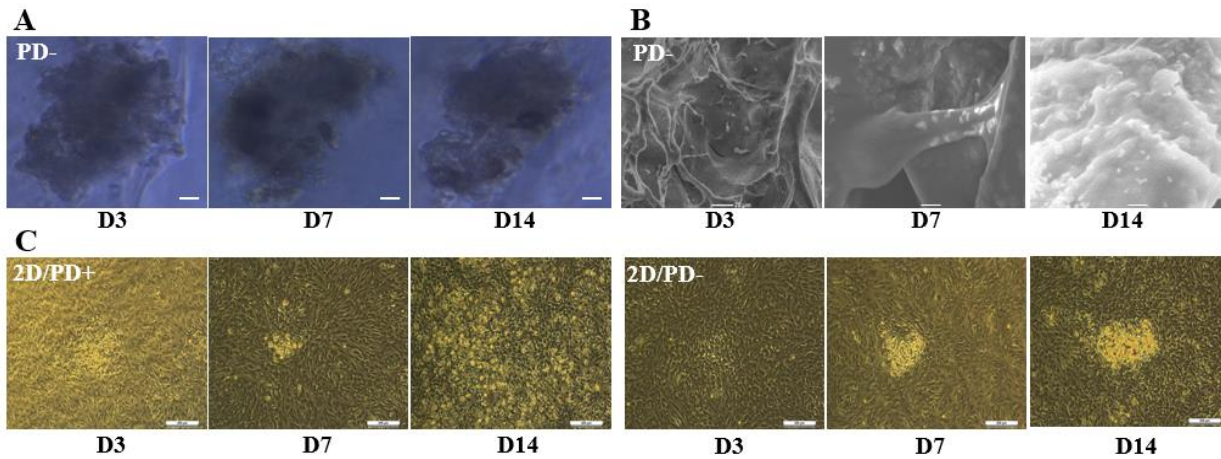
DOI: 10.1126/sciadv.aaz1094

#### **This PDF file includes:**

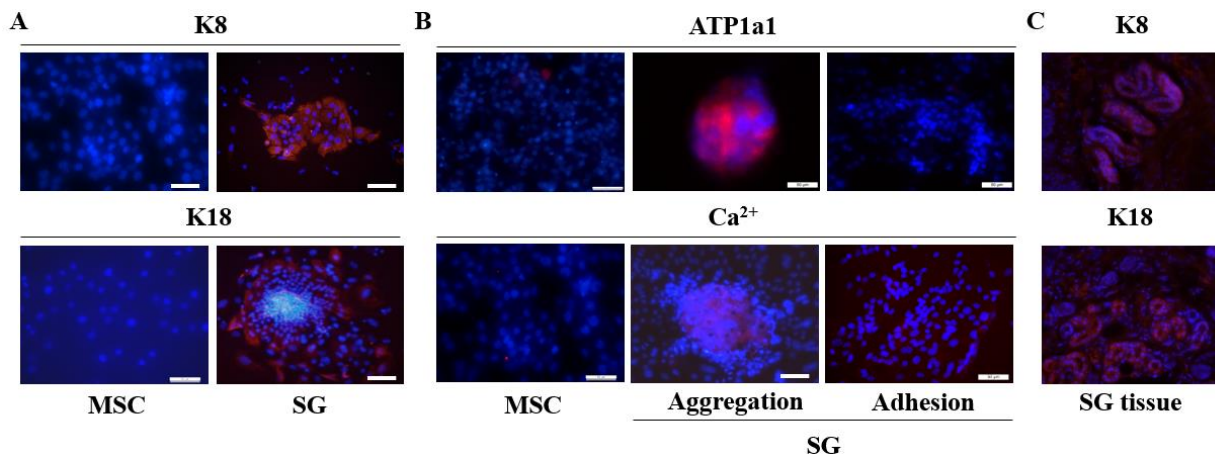
- Fig. S1. Biocompatibility of 3D-bioprinted construct and cellular morphology in 2D monolayer culture.
- Fig. S2. Expression of SG-specific and secretion-related markers in MSCs and SG cells in vitro.
- Fig. S3. Transcriptional and translational expression of epithelial and mesenchymal markers in 3D-bioprinted cells with and without PD.
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## Supplementary Materials

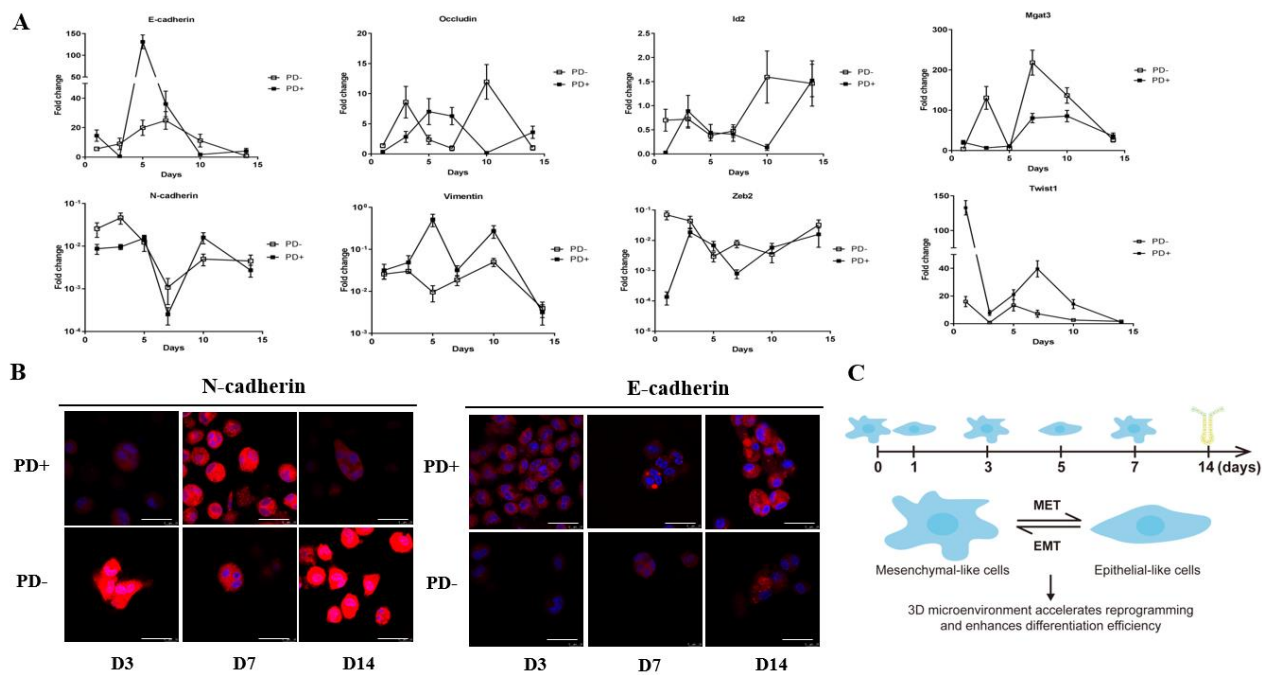
### Figure legends



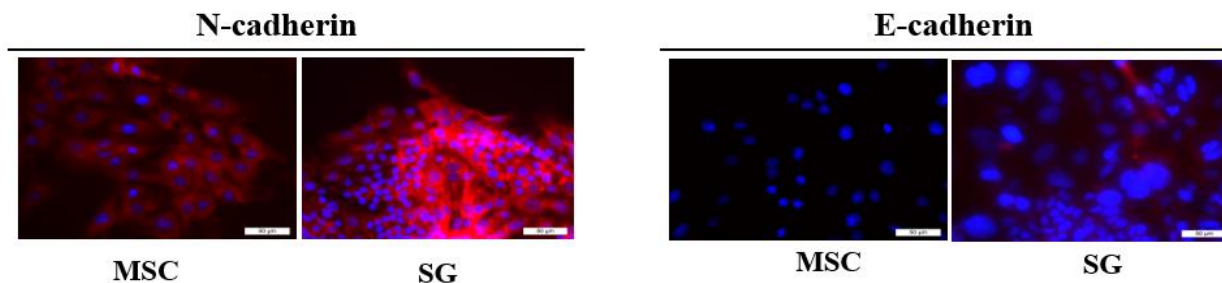
**Fig. S1. Biocompatibility of 3D-bioprinted construct and cellular morphology in 2D monolayer culture.** a Representative microscopy images of cell aggregates at different days (scale bars: 50  $\mu\text{m}$ ). b SEM images of 3D structure (scale bars: 20  $\mu\text{m}$ ). c Microscopic images of 2D monolayer culture cells with and without application of plantar region dermis (PD). (scale bars: 200  $\mu\text{m}$ ).



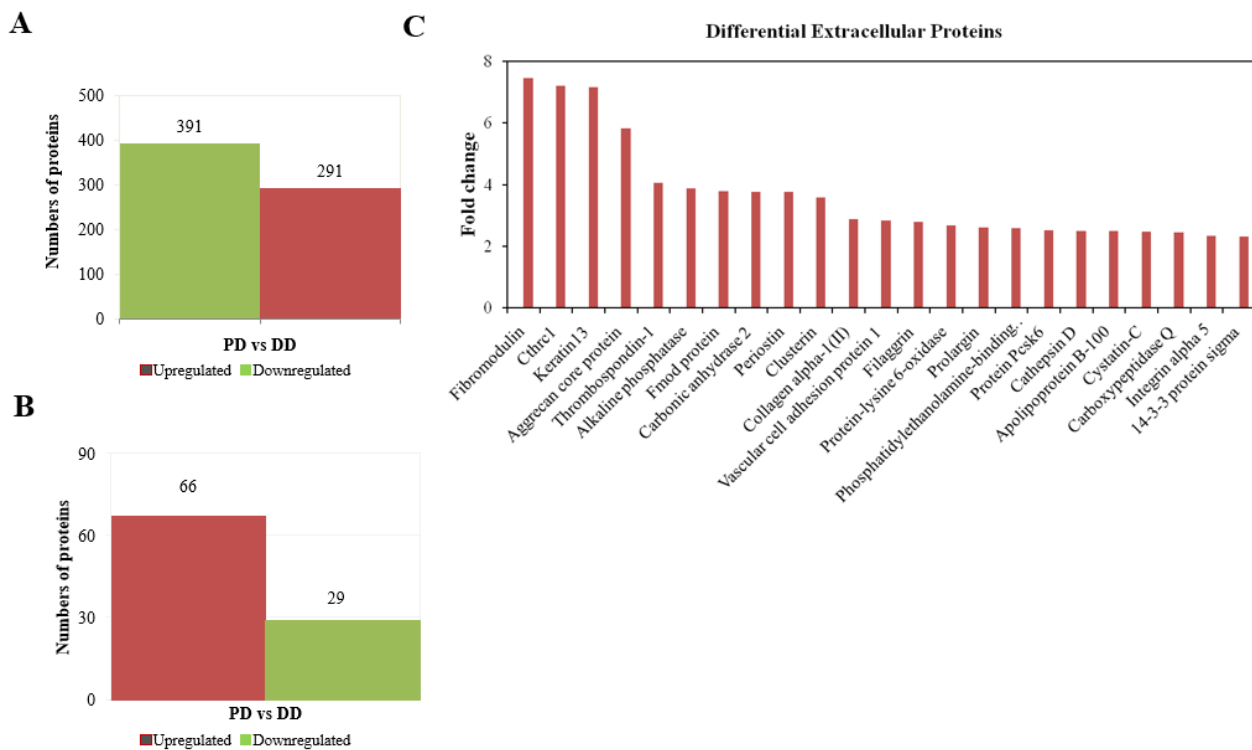
**Fig. S2. Expression of SG-specific and secretion-related markers in MSCs and SG cells in vitro.** a Expression of K8 and K18 in MSCs and SG cells in 2D monolayer culture. (K8, K18: red, DAPI: blue, scale bars: 200  $\mu\text{m}$ ) b Expression of ATP1a1 and Ca<sup>2+</sup> in MSCs and SG clusters and monolayer cells. (ATP1a1, Ca<sup>2+</sup>: red, DAPI: blue, scale bars: 50  $\mu\text{m}$ ).



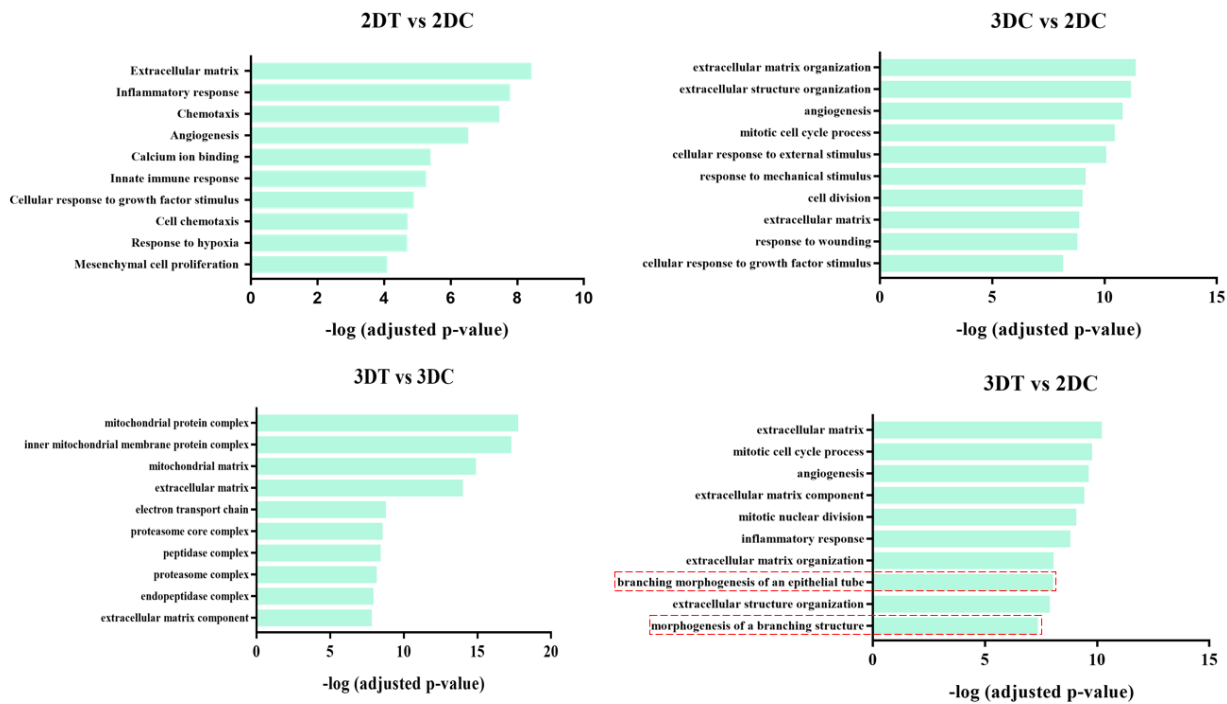
**Fig. S3. Transcriptional and translational expression of epithelial and mesenchymal markers in 3D-bioprinted cells with and without PD.** (A) Transcriptional expression of epithelial markers (E-cadherin, Occludin) and transcription factors of MET (Id2, Mgat3), mesenchymal markers (N-cadherin, Vimentin) and transcription factors of EMT (Twist1, Zeb2) in 3D-bioprinted cells with or without PD at day 1, 3, 5, 7, 10, and 14 culture by qRT-PCR. Data are mean±SEM. (B) Comparison of N-cadherin and E-cadherin expression in 3D-bioprinted cells with and without PD (N-cadherin, E-cadherin: red, DAPI: blue, scale bars: 50 μm). (C) Schematic illustration of the role of MET-EMT in cell differentiation.



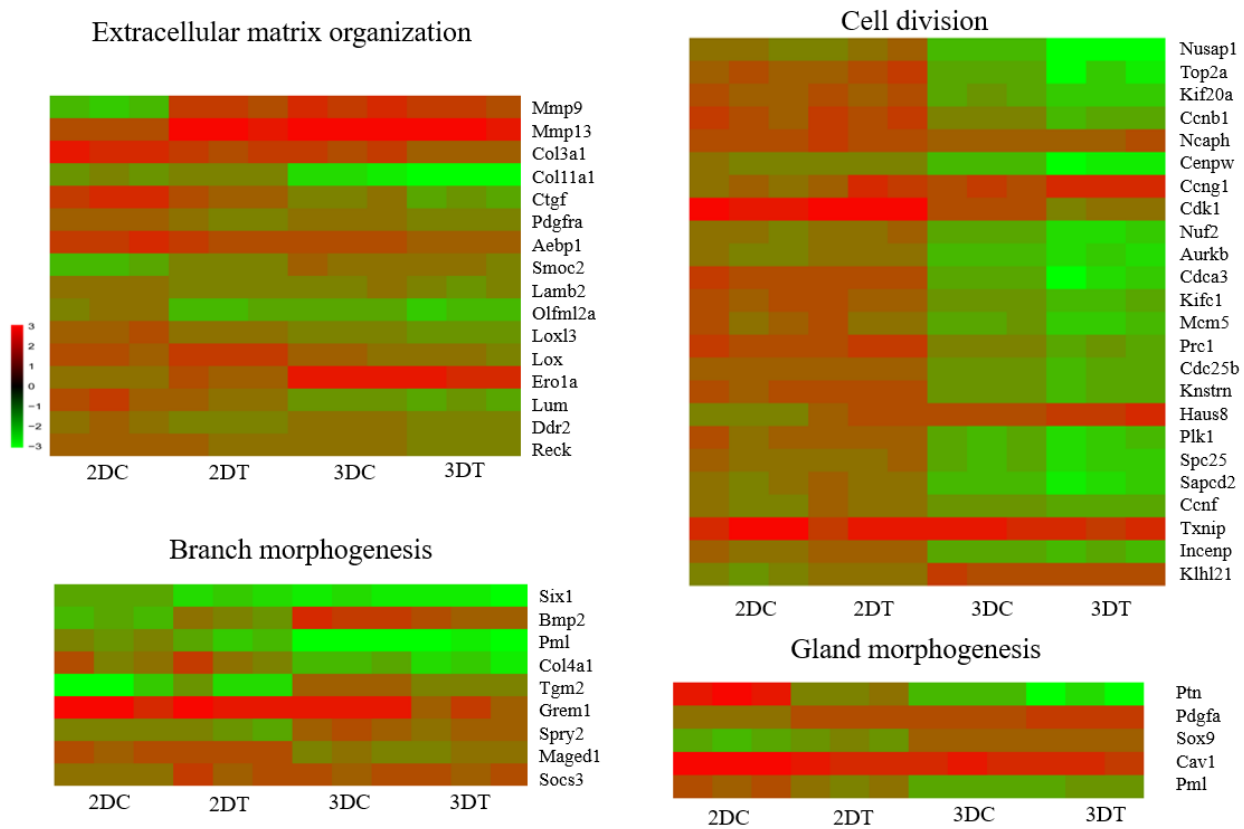
**Fig. S4. Expression of N- and E-cadherin in MSCs and SG cells in 2D monolayer culture.** (N-cadherin, E-cadherin: red, DAPI: blue, scale bars: 50 μm).



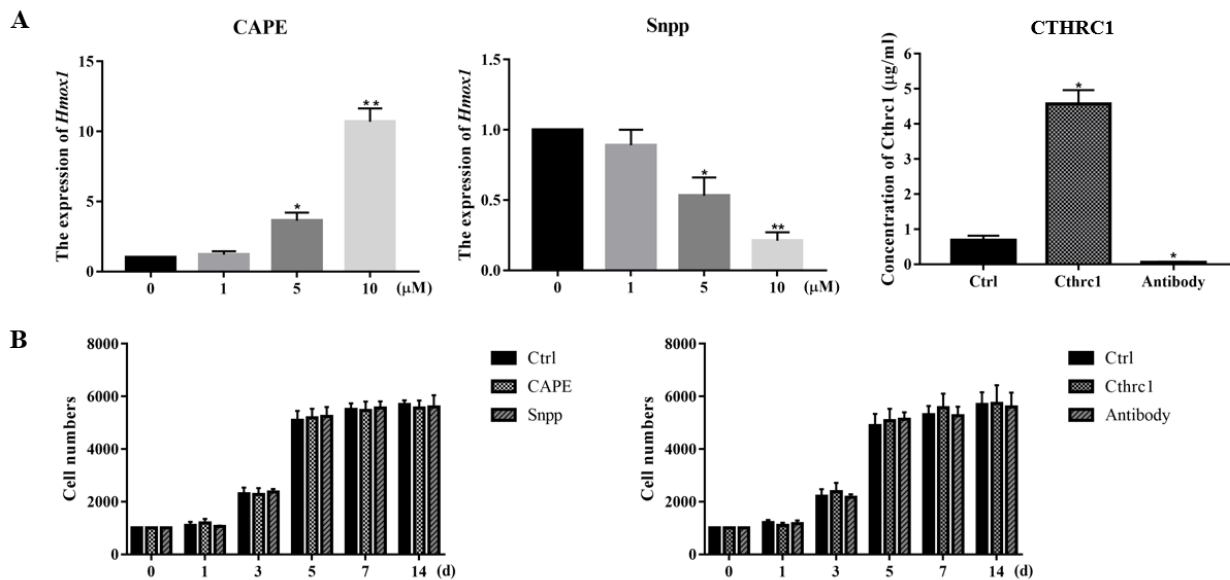
**Fig. S5. Proteomic microarray assay of differential gene expression between PD and DD ECM in postnatal mice. a** Comparison of proteins down- and upregulated between PD and DD ECM in postnatal mice. **b** Number of proteins in PD and DD ECM in postnatal mice. **c** Proteins with significant differential expression between PD and DD ECM in postnatal mice.



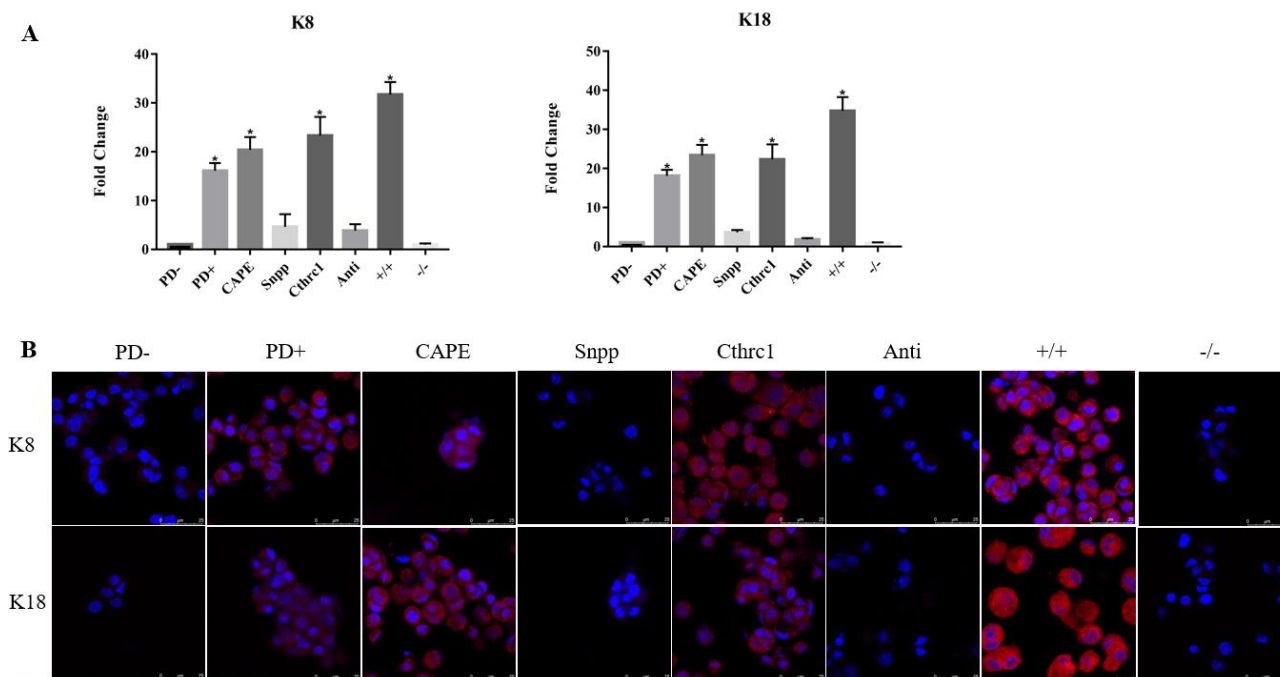
**Fig. S6. GO term analysis of differentially expressed pathways.**



**Fig. S7. Heat maps illustrating differential expression of genes implicated in ECM organization, cell division, and gland and branch morphogenesis.**



**Fig. S8. The expression of *Hmox1* and the concentration of CTHRC1 on treatment and the related effects on cell proliferation.** Effects of CAPE and Snpp on the expression of *Hmox1* and effects of recombinant CTHRC1 and CTHRC1 antibody on the concentration of CTHRC1 (A) and the related effects on cell proliferation (B) \* $p < 0.05$ .



**Fig. S9. The expression of K8 and K18 with *Hmox1* and *CTHRC1* regulation.** Expression of K8 and K18 at transcriptional (A) and translational (B) level with *Hmox1* and *CTHRC1* regulation (K8, K18: red, DAPI: blue, scale bars: 200  $\mu$ m) \* $p < 0.05$ .

**Table S1. Primers for qRT-PCR of all the genes.**

<b>Primers</b>	<b>Sequence(5'to3')</b>
<b>K8-F</b>	<b>GGAGGACTGAGTTCATCCTACGGG</b>
<b>K8-R</b>	<b>GGGTTTCAATCTTCTTCACGACCA</b>
<b>K18-F</b>	<b>CCGTCTTGCCGCTGATGACTTT</b>
<b>K18-R</b>	<b>TCCGCCATGATCTTGCTGAGGT</b>
<b>ATP1a1-F</b>	<b>GACGCCTTTCAGAATGCCTACC</b>
<b>ATP1a1-R</b>	<b>TGTGACCATGATGACCTTAATCCC</b>
<b>Fxyd2-F</b>	<b>ATGGACAGGTGGTACTTGGGTG</b>
<b>Fxyd2-R</b>	<b>CATTGACCTGCCTATGCTTCTTACT</b>
<b>Aqp5-F</b>	<b>GCCTTATCCATTGGCTTGTCTGTC</b>
<b>Aqp5-R</b>	<b>CCCAGTCCTCCTCCGGCTCATA</b>
<b>E-cadherin-F</b>	<b>CCATCGCCTACACCATCCTCAG</b>
<b>E-cadherin-R</b>	<b>TACGGGCACCGACCTCATTCTC</b>
<b>Occludin-F</b>	<b>GCTTGGGACAGAGCCTATGGAA</b>
<b>Occludin-R</b>	<b>CCCGATCTAATGACGCTGGTAA</b>
<b>Mgat3-F</b>	<b>CCGACTTGTTACGGACTCCACTCTA</b>
<b>Mgat3-R</b>	<b>TCAGCCACCTTGGTCTTCTCCTCT</b>
<b>Id2-F</b>	<b>TCCGGTGAGGTCCGTTAGGAAA</b>
<b>Id2-R</b>	<b>GGTGCAGGCTGACGATAGTGGG</b>
<b>Vimentin-F</b>	<b>TGACCGCTTCGCCAACTACATC</b>
<b>Vimentin-R</b>	<b>GCTTCCTCCCTCTGGAGCATCTC</b>
<b>N-cadherin-F</b>	<b>CCTCATGTTTCGTGGTATGGATG</b>
<b>N-cadherin-R</b>	<b>TATTGTGGCTCAGCGTGGATAG</b>
<b>Zeb2-F</b>	<b>GTTCGGCATGAACCCATTTAGT</b>
<b>Zeb2-R</b>	<b>ATCCTTGTTTCCGCTGGTATTT</b>
<b>Twist1-F</b>	<b>ACCCAGTCGCTGAACGAGGCATT</b>
<b>Twist1-R</b>	<b>TCCATCCTCCAGACGGAGAAGGC</b>
<b>GAPDH-F</b>	<b>CTCTGGAAAGCTGTGGCGTGAT</b>
<b>GAPDH-R</b>	<b>GGAGACAACCTGGTCCTCAGTGTAG</b>
<b>IL-10-F</b>	<b>GCTCTTACTGACTGGCATGAG</b>
<b>IL-10-R</b>	<b>CGCAGCTCTAGGAGCATGTG</b>
<b>Mmp9-F</b>	<b>CTGGACAGCCAGACACTAAAG</b>
<b>Mmp9-R</b>	<b>CTCGCGGCAAGTCTTCAGAG</b>
<b>Ptges-F</b>	<b>GGATGCGCTGAAACGTGGA</b>
<b>Ptges-R</b>	<b>CAGGAATGAGTACACGAAGCC</b>
<b>Vsig-F</b>	<b>TGCAGGTCAGGTCAGTATGGT</b>
<b>Vsig-R</b>	<b>CTCGGTGGTGGTATAGAGGCA</b>
<b>Vwa1-F</b>	<b>GGGGGACCTGTTGTTCTCTG</b>
<b>Vwa1-R</b>	<b>CATCCTGTATAGCCTGGCCT</b>
<b>Hmox1-F</b>	<b>AAGCCGAGAATGCTGAGTTCA</b>
<b>Hmox1-R</b>	<b>GCCGTGTAGATATGGTACAAGGA</b>

<b>Bmp2-F</b>	<b>GGGACCCGCTGTCTTCTAGT</b>
<b>Bmp2-R</b>	<b>TCAACTCAAATTCGCTGAGGAC</b>
<b>Sox9-F</b>	<b>GAGCCGGATCTGAAGAGGGA</b>
<b>Sox9-R</b>	<b>GCTTGACGTGTGGCTTGTTT</b>
<b>Tgm2-F</b>	<b>GACAATGTGGAGGAGGGATCT</b>
<b>Tgm2-R</b>	<b>CTCTAGGCTGAGACGGTACAG</b>