

**Directed differentiation of human pluripotent stem cells into diverse organ-specific mesenchyme of the digestive and respiratory systems.**

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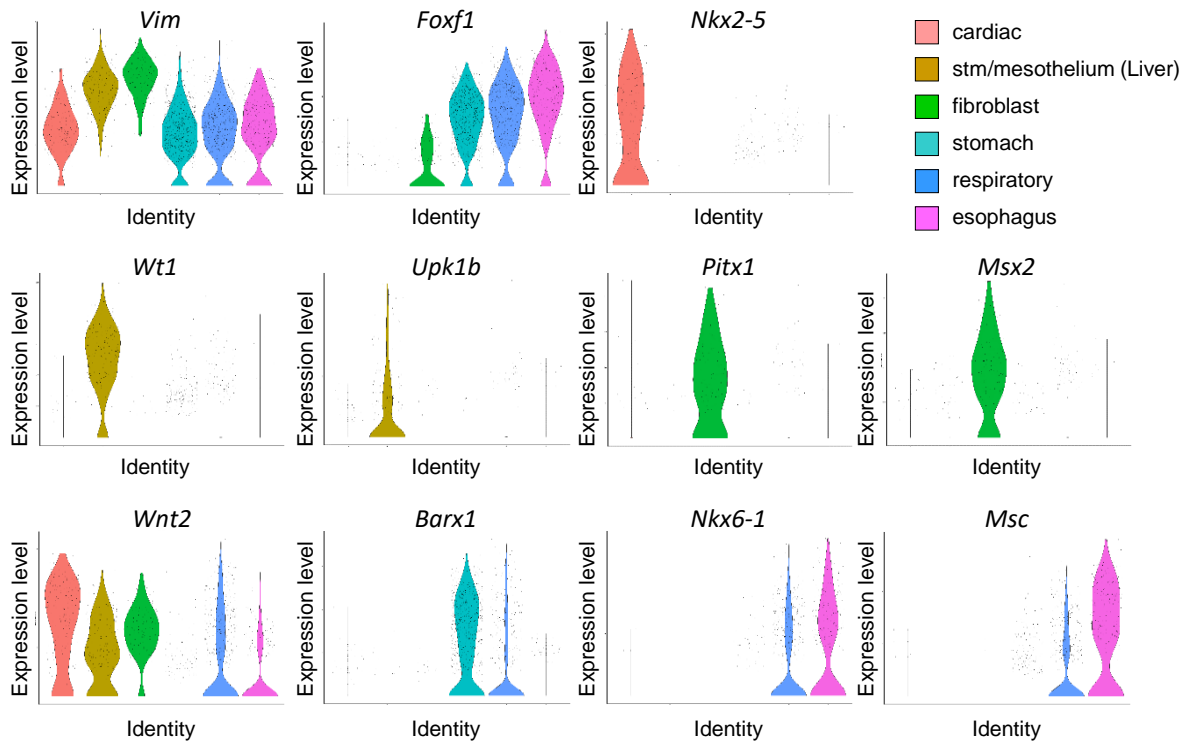
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**Contents:** Supplementary Figures 1-3

# Supplementary Figure 1

**a**

Marker gene expression in mesoderm at mouse E9.5

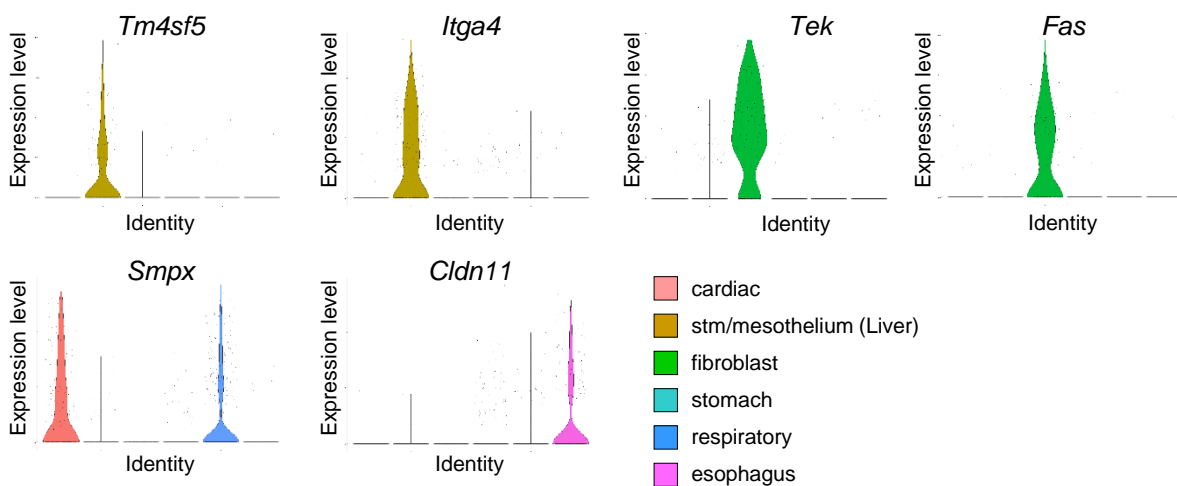


**b**

Organ-specific mesoderm	Marker genes
Liver STM/Mesothelium	<i>Wt1, Upk1b, Gata4, Tbx18</i>
Liver fibroblasts	<i>Pitx1, Krt19, Msx1, Msx2</i>
Gastric mesoderm	<i>Barx1, Nkx3-2, Foxf1</i>
Respiratory mesoderm	<i>Nkx6-1, Tbx5, Wnt2, Foxf1</i>
Esophageal mesoderm	<i>Nkx6-1, Msc, Wnt4, Foxf1</i>

**c**

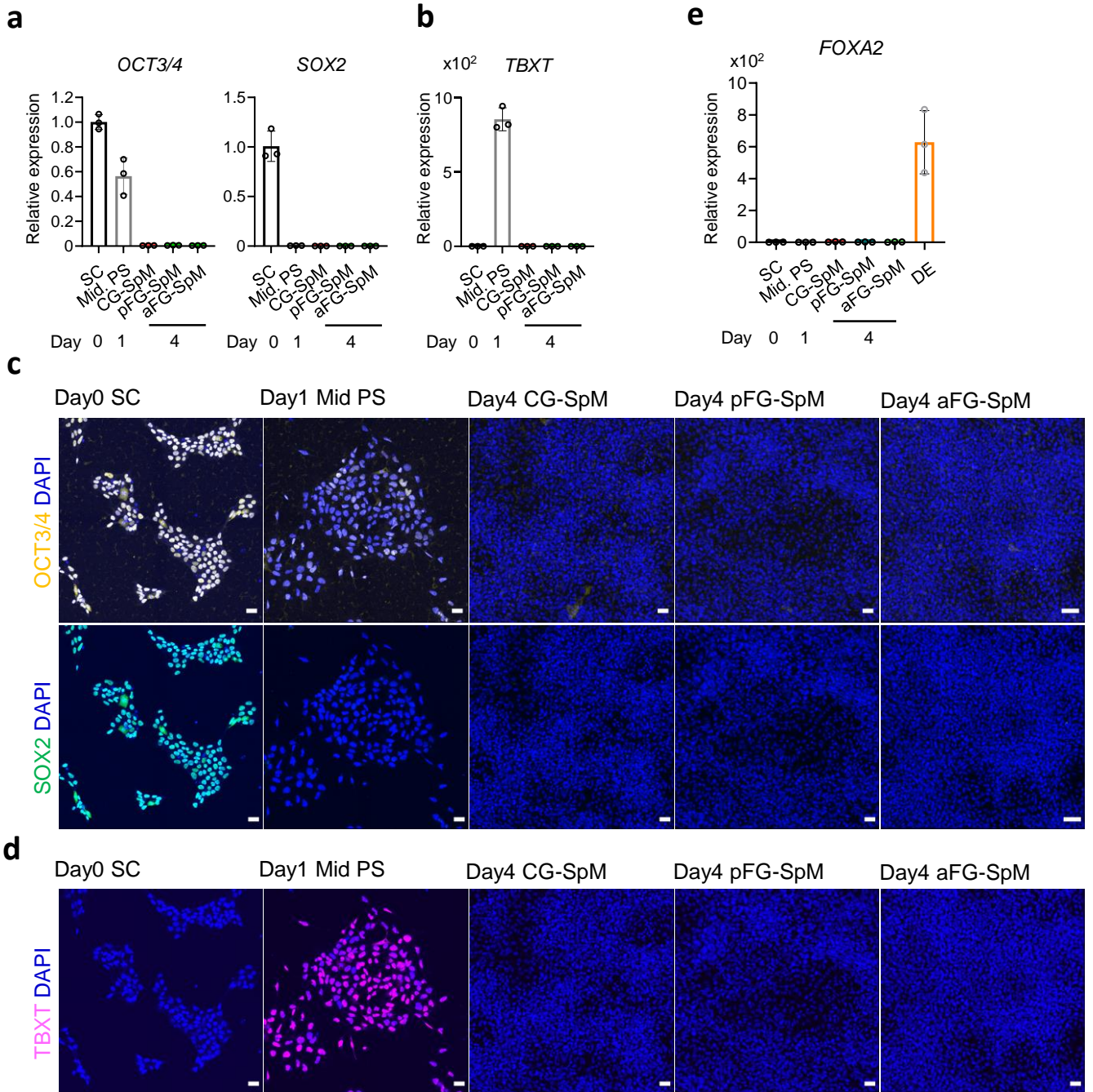
Surface marker gene expression in mesoderm at mouse E9.5



## Supplementary Fig. 1

**a**, The violin plots display the sc-RNA-seq expression of organ-specific mesenchyme markers in different mouse embryonic mesoderm cell populations at E9.5. **b**, Summary of combinatorial marker gene expression from the mouse foregut that were used to define the different hPSC-derived cell types. **c**, The violin plots display the expression of organ-specific mesenchymal surface markers in mouse embryonic mesoderm at E9.5. The plasma membrane genes (1,917 genes) were extracted from the human protein atlas (<https://www.proteinatlas.org/>). Among these genes, 390 genes were overlapped with differentially expressed genes in mesoderm at E9.5. Representative organ-specific mesoderm markers are shown.

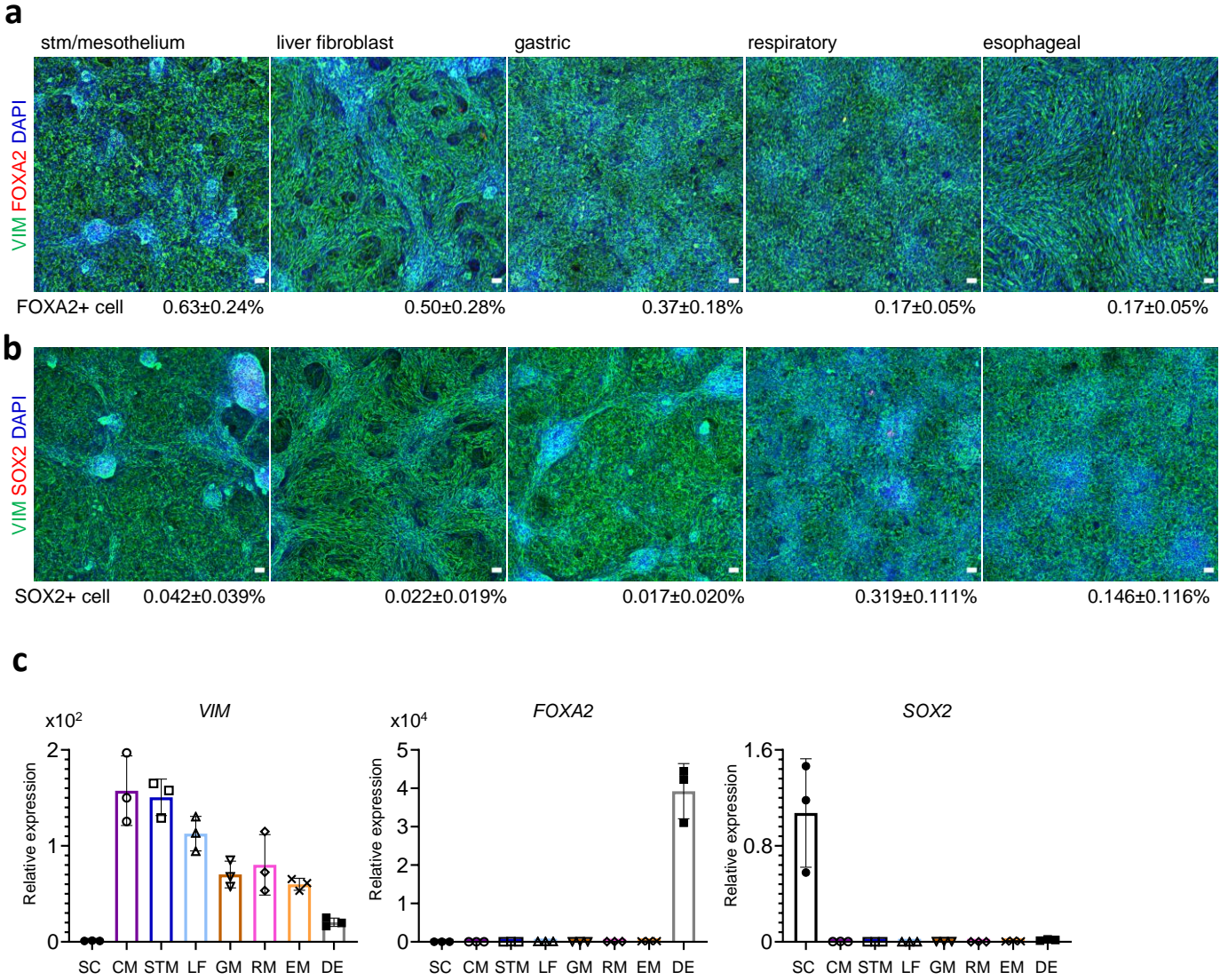
# Supplementary Figure 2



## Supplementary Fig. 2

**a**, Relative mRNA expression of pluripotent marker genes, *OCT3/4* and *SOX2* by quantitative RT-PCR from day0 to day4. **b**, Relative mRNA expression of *TBXT* as an early mesoderm marker by quantitative RT-PCR from day0 to day4. **c**, Immunostaining for *OCT3/4* (yellow) and *SOX2* (green). **d**, Immunostaining for *TBXT* (red). The images in **c** and **d** are maximum intensity projection of confocal stacks. **e**, Relative mRNA expression of *FOXA2* as an endoderm marker by quantitative RT-PCR from day0 to day4. Each bar indicates the average from the 3 independent wells with standard deviation. Scale bar; 50 $\mu$ m

# Supplementary Figure 3



## Supplementary Fig. 3

**a**, Immunostaining for VIM (green), FOXA2 (red), and DAPI (blue) at day7 mesenchyme. **b**, Immunostaining for VIM (green), SOX2 (red), and DAPI (blue) at day7 mesenchyme. The images in a and b are maximum intensity projection of confocal stacks. **c**, Relative mRNA expression of pan-mesoderm marker, VIM, endoderm marker, FOXA2, and ectoderm marker, SOX2, by quantitative RT-PCR. Each bar indicates the average from the 3 independent wells with standard deviation. Scale bar; 50µm (a, b).