

1 **Endometrial organoids: a reservoir of functional mitochondria for uterine repair**

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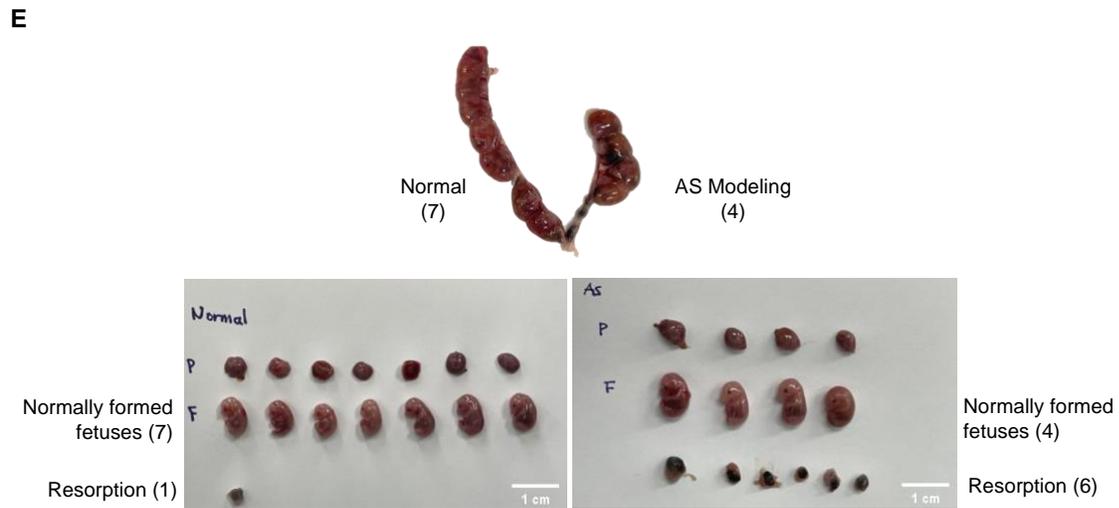
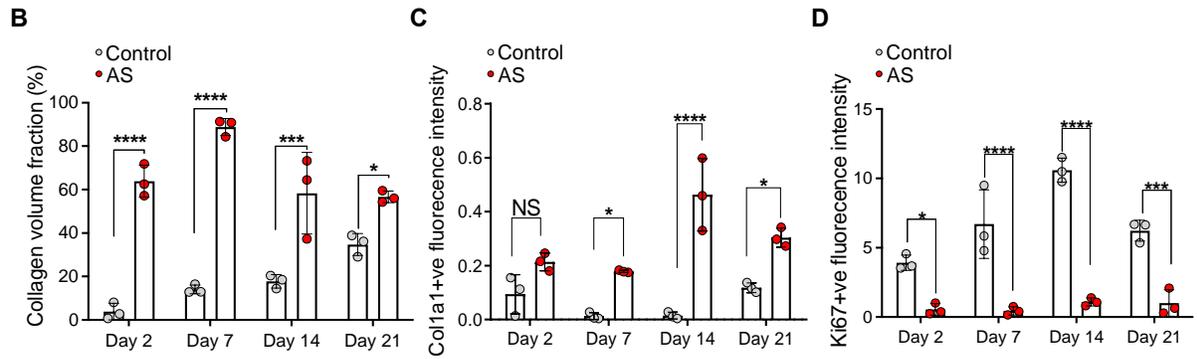
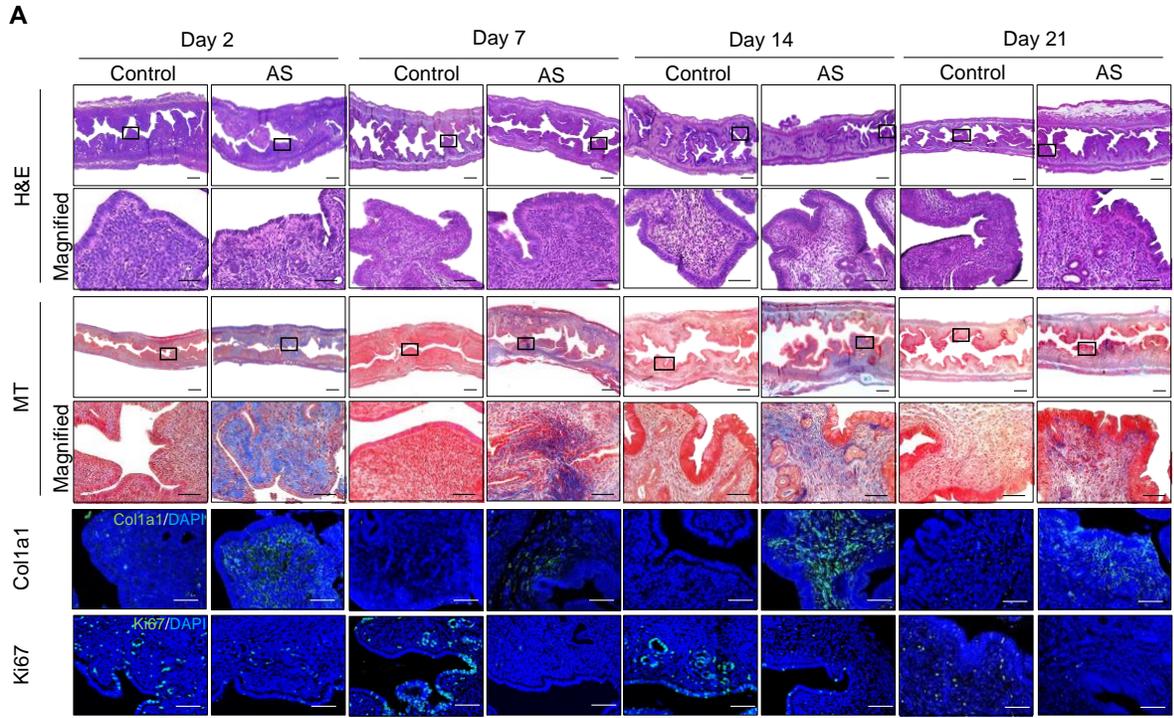
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22 **Supplementary Figure 1. Successful generation of a murine model of AS**

23 **(A)** Representative immunostaining images of H&E, MT, Col1a1 (green), and Ki67 (green) and DAPI (blue)
24 of normal and AS-induced endometrium at indicated days (Day 7, 14 and 21). Scale bar; 25um. Collagen
25 volume fraction **(B)**, Col1a1 intensity **(C)**, or Ki67⁺ fluorescence intensity **(D)** shown in **(A)** was quantified
26 (Total number of mice=48; 3 mice per group; triplicates). Data were expressed as mean \pm SD, analyzed
27 using the ordinary two-way ANOVA with Turkey's multiple comparisons test including P-values (*<0.05,
28 **<0.01, ***<0.001, ****<0.0001, NS; not significant). **(E)** Representative images of fetus and placenta from
29 normal and AS-induced horn at pregnancy day 14. Scale bar; 1cm.

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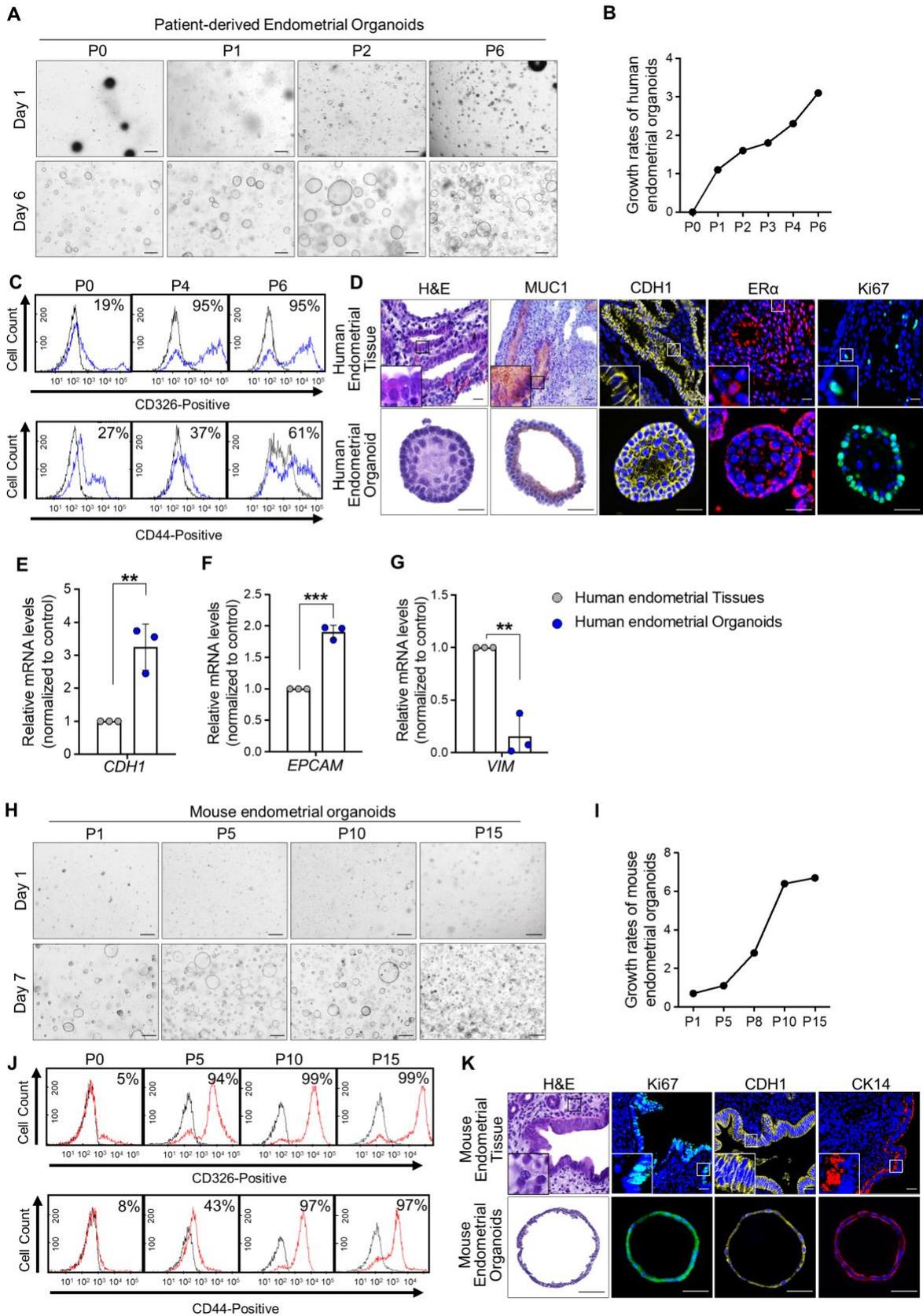
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47 **Supplementary Figure 2. Generation of human or mouse endometrial tissue-derived organoids**

48 **(A)** Expansion and morphological observation of human endometrial organoids from each passage at day
49 1 and 6. Scale bar; 25um. **(B)** Growth curve of human endometrial organoids up to passage 6. **(C)** Flow
50 cytometry analyses of CD326 and CD44 (blue) versus isotype controls (grey) in human endometrial
51 organoids from each passage. **(D)** Immunostaining images of H&E, MUC1, CDH1, ER α , and Ki67 in human
52 endometrial organoids and its parental tissues. Scale bar; Tissue-100um (H&E, MUC1), 20um (CDH1, ER α
53 and Ki67), Organoid-100um (H&E, MUC1), 30um (CDH1, ER α and Ki67). QRT-PCR analyses of epithelial
54 markers (*CDH1* **(E)** and *EPCAM* **(F)**) and stromal marker (*VIM* **(G)**) in human endometrial organoids
55 compared with its parental tissues (3 human endometrial organoids were used in characterization). Data
56 were expressed as mean \pm SD, analyzed using the unpaired t test including P-values (*<0.05, **<0.01,
57 ***<0.001, ****<0.0001, NS; not significant). **(H)** Expansion and morphological observation of mouse
58 endometrial organoids from each passage at day 1 and 7. Scale bar; 25um. **(I)** Growth curve of mouse
59 endometrial organoids up to passage 15. **(J)** Flow cytometry analyses of CD326 (red) and CD44 (red)
60 versus isotype controls (grey) in mouse endometrial organoids. **(K)** Immunostaining images of H&E, Ki67,
61 CDH1, and CK14 in mouse endometrial organoids and its parental tissues (Four independent sets of human
62 endometrial organoids were used for characterization). Scale bar; Tissue-100um (H&E), 20um (Ki67, CDH1
63 and CK14), Organoid-100um (H&E), 30um (Ki67, CDH1 and CK14).

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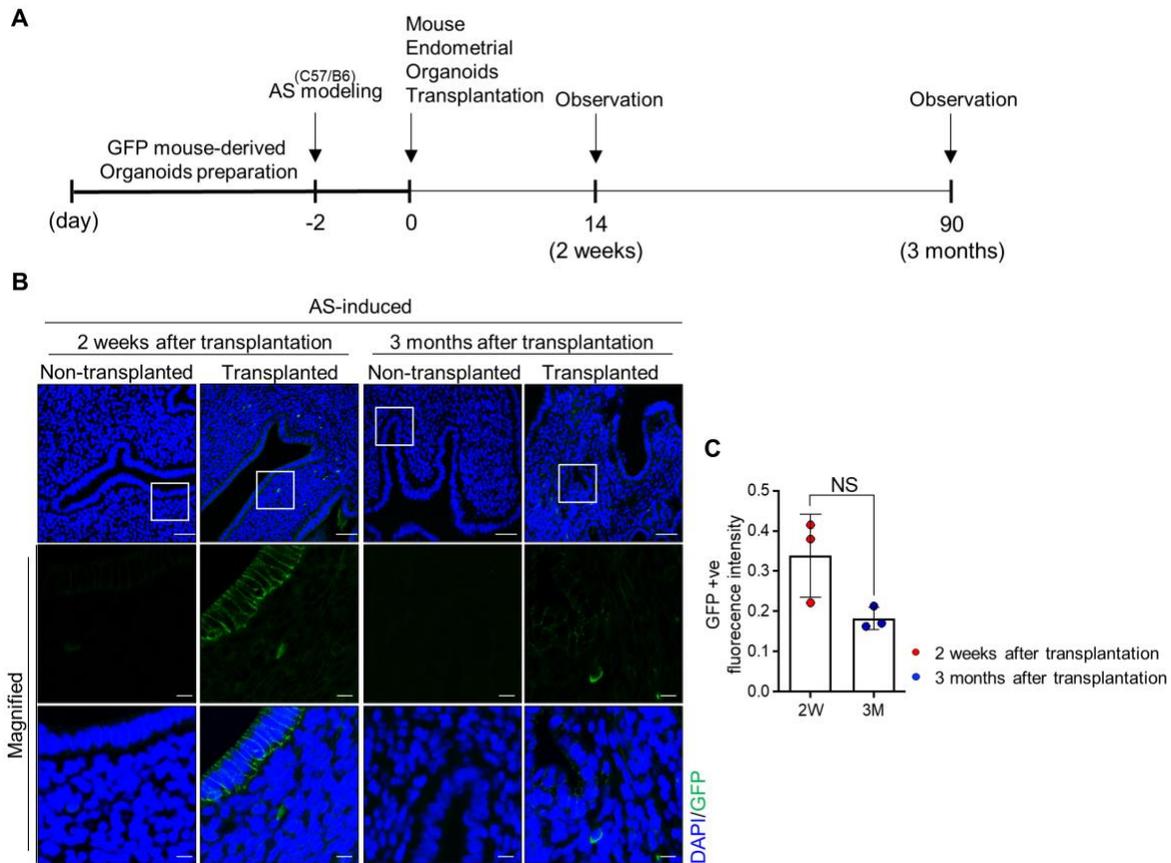
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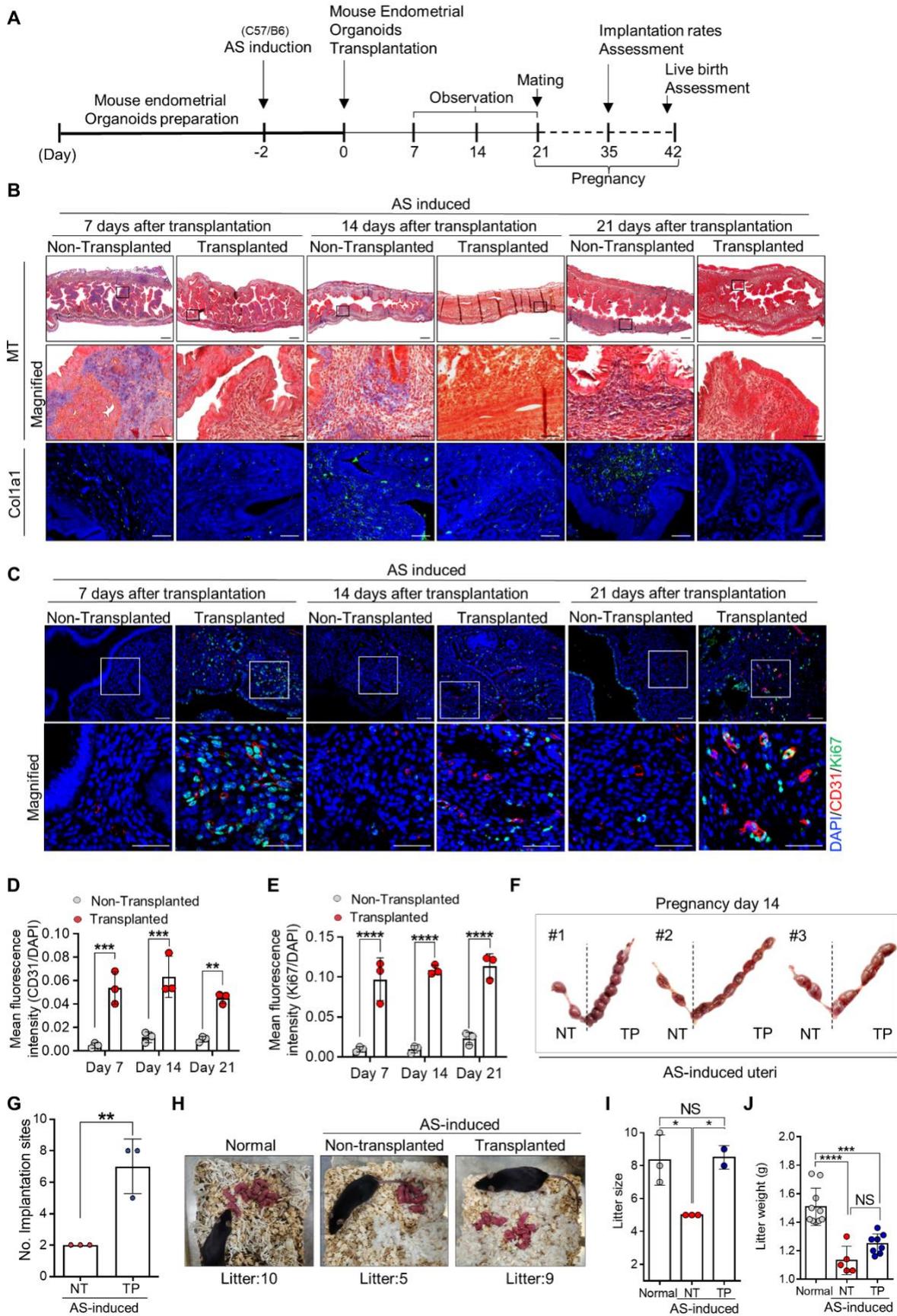
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74 **Supplementary Figure 3. Lasting duration time of mouse endometrial organoid transplantation in**
 75 **AS-induced endometrium**

76 **(A)** An experimental design of GFP-mouse endometrial organoid transplantation to AS-induced uteri of
 77 C57/B6 mice. **(B)** Immunofluorescent analyses of GFP (green) and DAPI (blue) in GFP-mouse endometrial
 78 organoid-transplanted AS-induced endometrium 3 months after transplantation compared with 2-week time
 79 point or non-transplanted AS-induced endometrium (Total number of mice=6; 3 mice per group). Scale bar;
 80 50um, 10um (Magnified). Quantified intensity of GFP signals detected in AS-induced endometrium 3
 81 months or 2 weeks after transplantation of GFP mouse-derived endometrial organoids was shown in a
 82 graph **(C)**. Data were expressed as mean \pm SD, analyzed using the unpaired t test including P-values (NS;
 83 not significant).



85 **Supplementary Figure 4. Therapeutic effects of mouse endometrial organoid transplantation in AS-**
86 **induced endometrium**

87 **(A)** An experimental design of mouse endometrial organoid transplantation to AS-induced uteri of C57/B6
88 mice and further analyses including fertility assessments. **(B)** Immunohistochemical analyses of MT and
89 Col1a1 in mouse endometrial organoid-transplanted AS-induced endometrium compared with non-
90 transplanted AS-induced endometrium at indicated days (Day 7, 14 and 21) (Total number of mice=48; 3
91 mice per group; triplicates and three independent sets of mouse organoids were used for each
92 transplantation). Scale bar; 25um. **(C)** Immunofluorescent analyses of CD31 (red) and Ki67 (green) with
93 DAPI (blue) in in mouse endometrial organoid-transplanted AS-induced endometrium compared with non-
94 transplanted AS-induced endometrium at indicated days (Day 7, 14 and 21). Scale bar; 25um.
95 Immunofluorescent CD31 and Ki67 intensity were quantified in graphs of **(D-E)**. **(F)** Representative images
96 of uteri with implantation sites on pregnancy day 14 (n=3). **(G)** The total number of implantation sites was
97 counted in each horn and quantified. **(H)** Representative images of mouse litters with their mother with
98 endometrial organoids-transplanted AS-induced endometrium compared to groups with non-transplanted
99 AS-induced endometrium and normal endometrium. Comparisons of sizes **(I)** and weights **(J)** of litters from
100 mice with endometrial organoids-transplanted AS-induced endometrium compared to groups with non-
101 transplanted AS-induced endometrium and normal endometrium. Data were expressed as mean \pm SD,
102 analyzed using the unpaired t test including P-values (*<0.05, **<0.01, ***<0.001, ****<0.0001, NS; not
103 significant).

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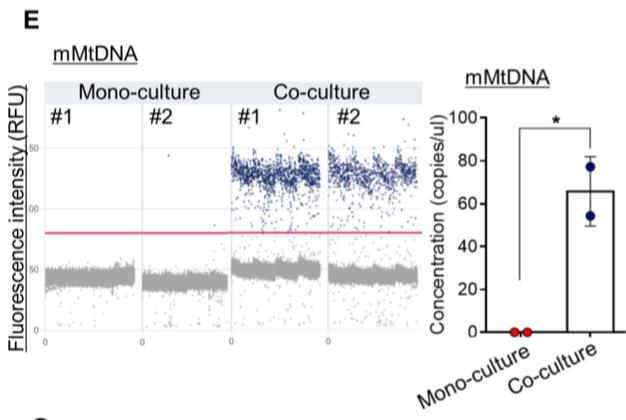
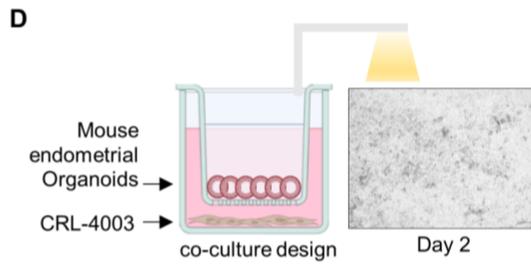
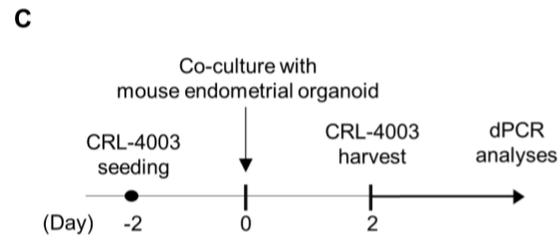
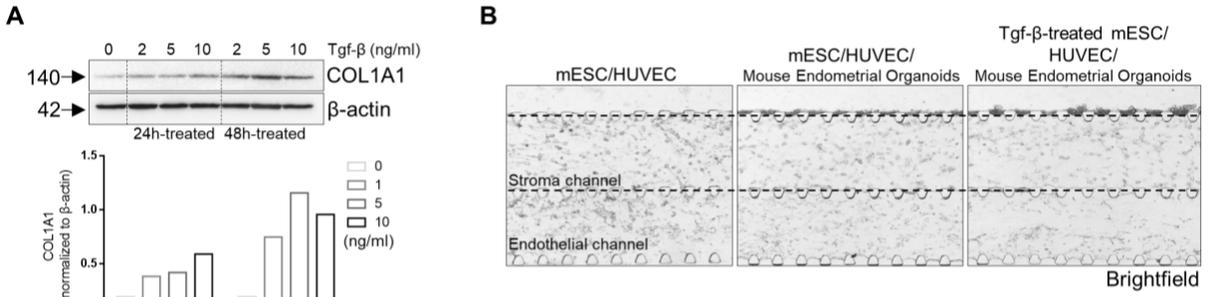
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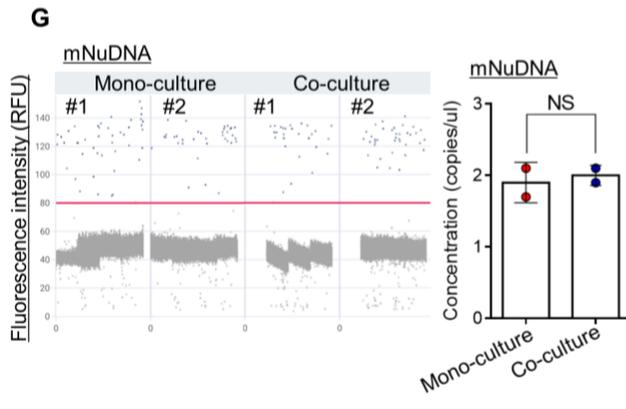
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Target	Sample	Partitions			
		Valid	+	-	
mMtDNA	Mono-culture	#1	25286	0	25286
		#2	25442	1	25441
	Co-culture	#1	25403	1403	24000
		#2	25255	983	24272



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Target	Sample	Partitions			
		Valid	+	-	
mNuDNA	Mono-culture	#1	25441	45	25396
		#2	25491	35	25456
	Co-culture	#1	25476	39	25437
		#2	25476	42	25434

112 **Supplementary Figure 5. Endometrial organoid-originated mitochondrial migration into Tgf- β -**
113 **induced damaged cells**

114 **(A)** Immunoblotting analysis of COL1A1 in mESCs with Tgf- β treatment at indicated concentration (0, 2, 5,
115 and 10 ng/ml) and timepoints (0, 24, and 48 h). **(B)** Brightfield images about serial steps of cell loading for
116 establishment of the endometrium-on-a-chip. Scale bar; 25um. **(C)** An experimental design of co-culture
117 of CRL-4003 with mouse endometrial organoids. **(D)** An experimental design of transwell assay and
118 brightfield image of co-cultured CRL-4003 with mouse endometrial organoids at day 2. **(E-F)** digital PCR
119 analyses of mMtDNA (mouse mitochondrial DNA) and partitioning information in co-cultured CRL-4003 with
120 mouse endometrial organoids compared with monocultured CRL-4003. **(G-H)** dPCR digital PCR analyses
121 of mNuDNA (mouse nucleus DNA) and partitioning information in co-cultured CRL-4003 with mouse
122 endometrial organoids compared with monocultured CRL-4003. Data were expressed as mean \pm SD,
123 analyzed using unpaired t test including P-values (*<0.05, **<0.01, ***<0.001, ****<0.0001, NS; not
124 significant).

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Supplementary Table 1. Primer sequence pairs used for QRT-PCR and RT-DPCR.

Species	Gene	Direction	Sequence
Mouse	<i>Rpl7</i>	Forward	TCAATGGAGTAAGCCCAAAG
		Reverse	CAAGAGACCGAGCAATCAAG
	<i>Actb</i>	Forward	GTGACGTTGACATCCGTAAAGA
		Reverse	GCCGGACTCATCGTACTCC
	<i>Tgfb1</i>	Forward	GTGAAACGGAAGCGCATCGAAG
		Reverse	CATAGTAGTCCGCTTCGGGCTCC
	<i>Timp1</i>	Forward	CTTGGTTCCTGGCGTACTC
		Reverse	ACCTGATCCGTCCACAAACAG
	<i>Col1a1</i>	Forward	CTGGCGGTTCCAGTCCAAT
		Reverse	TTCCAGGCAATCCACGAGC
	<i>Pgc1a</i>	Forward	ACGGTTTACATGAACACAGCTGC
		Reverse	CTTGTTTCGTTCTGTTCAGGTGC
	<i>Nrf1</i>	Forward	GAACGCCACCGATTTCACTGTC
		Reverse	CCCTACCACCCACGAATCTGG
	<i>Mfn1</i>	Forward	TGCATGTTTCACCACAGTTTC
		Reverse	GTAGCTCACAACCACCTGTAA
	<i>Fis1</i>	Forward	AGGCTCTAAAGTATGTGCGAGG
		Reverse	GGCCTTATCAATCAGGCGTTC
	<i>mMtDNA</i>	Forward	AGGCATGAAAGGACAGCACA
		Reverse	TTGGGGTTTGGCATTAAAGAGGA
	<i>mNuDNA</i>	Forward	GAATTCAGATTTGTGCATACACAGTGACT
		Reverse	AACATTTTTTCGGGAATAAAAGTTGAGT
	<i>Tnfa</i>	Forward	CCCTCACACTCAGATCATCTTCT
		Reverse	GCTACGACGTGGGCTACAG
	<i>Hk2</i>	Forward	GGAGGAACCAATTTCCCTGCTGCT
		Reverse	CCTTTGATCCCCATGTATCCAAGA
	<i>Scd1</i>	Forward	TTCTTGCGATACACTCTGGTGC
		Reverse	CGGGATTGAATGTTCTTGTCGT
	<i>Gpr84</i>	Forward	CTCCTGCTACCATGAGTCTGT
		Reverse	GTGCAGTAGAGTAGATCAGCCA
Human	<i>ACTB</i>	Forward	CATGTACGTTGCTATCCAGGC
		Reverse	GCCTTAATGTCACGCACGAT
	<i>CDH1</i>	Forward	AGTCACTGACACCAACGATAAT

	Reverse	ATCGTTGTTCACTGGATTTGTG
<i>EPCAM</i>	Forward	GTCTGTGAAAACACTACAAGCTGG
	Reverse	CAGTATTTTGTGCACCAACTGA
<i>VIM</i>	Forward	CGGCTGCGAGAGAAATTGC
	Reverse	CCACTTTCCGTTCAAGGTCAAG
<i>ITGB3</i>	Forward	AGTAACCTGCGGATTGGCTTC
	Reverse	GTCACCTGGTCAGTTAGCGT
<i>SPP1</i>	Forward	GAAGTTTCGCAGACCTGACAT
	Reverse	GTATGCACCATTCAACTCCTCG

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