



2 5. Supplementary material



Data shown as $2^{(-\Delta\Delta Cq)}$. ($\Delta\Delta Cq = \Delta Cq_{endometriosis} - \Delta Cq_{control}$)



Rel. mRNA expression control HESC

Figure S2. Correlation between the mRNA expression of TRP channels in control hESC (n = 3) and endometriosis-derived hESC (n = 4). The correlation was assessed using the non-parametric Spearman correlation test.



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Figure S3. Representative traces of individual endometriosis-derived hESC illustrating TRP functionality (a-d) Application of respectively 50 μ M THC, 10 nM GSK, 250 nM EA and 100 μ M OAG elicit a robust and reversible influx of calcium in hESC derived from endometriosis-patients.



Rel. mRNA expression eutopic tissue

Figure S4. Correlation between the mRNA expression of TRP channels in paired eutopic and ectopic tissue samples (n=3). The correlation was assessed using the non-parametric Spearman correlation test.





ENDOMETRIAL BIOPSIES										
Sample	Endometriosis stadium	AFS score	Cycle phase	Age (years)	BMI (kg/m²)	Smoking	Time interval between menses			
1	II	10	Menstrual	38	23.7	No	Normal			
2	II	14	Menstrual	37	23.3	No	Normal			
3	II	8	Menstrual	31	16.2	No	Normal			
4	II	12	Menstrual	28	18.9	Yes	Normal			
5	II	15	Menstrual	35	26.6	No	Normal			
1	II	12	Follicular	32	21.8	No	Normal			
2	II	10	Follicular	34	26.6	No	Normal			
3	II	6	Follicular	21	19	Yes	Normal			
4	II	10	Follicular	33	21.6	No	Normal			
5	II	6	Follicular	29	21.9	No	Normal			
6	II	8	Follicular	34	25.3	No	Normal			
	I		Γ	1						
1	II	12	Early luteal	37	nd	nd	nd			
2	II	18	Early luteal	33	20.8	No	Normal			
3	II	10	Early luteal	30	29.8	No	Normal			
4	II	6	Early luteal	35 19.6		No	Normal			
1	II	8	Late luteal	31	20.4	No	Normal			
2	II	8	Late luteal	26	20.5	Yes	Normal			
3	II	10	Late luteal	31	17.6	No	Normal			

 Table S1. Demographics and information of cycle phases/days for each endometrial biopsy.

 Normal time between menses equals 24–38 days; nd: not determined due to lack of information.

PRIMARY HUMAN ENDOMETRIAL STROMAL CELLS											
	Endometriosis	AFS	Cycle	Age	BMI		Time interval	Experimental destination			
Sample	stadium	score	phase	(years)	(kg/m²)	Smoking	between menses	qPCR	Ca ²⁺	Prol.	Migr.
1	0	n/a	Luteal	29	19.2	Occasionally	Normal		Х		
2	0	n/a	Luteal	31	21.2	No	Normal		Х		
3	0	n/a	Luteal	26	31.6	Yes	Normal	х	х		
4	0	n/a	Luteal	40	18.6	No	Normal	х	Х	х	Х
5	0	n/a	Luteal	27	25.3	Yes	Infrequent	х	х	х	Х
6	0	n/a	Luteal	29	24.6	No	Normal		х	х	Х
7	0	n/a	Luteal	39	28	No	Normal		х	х	Х
1	II	8	Luteal	31	20.2	No	Normal		х	х	х
2	II	12	Luteal	28	23	No	Normal	х	х		
3	II	8	Luteal	36	19.1	No	Normal		Х	х	Х
4	II	8	Luteal	30	20.8	No	Normal	х	Х	х	х
5	II	8	Luteal	37	31.6	No	Normal	х	х		
6	II	8	Luteal	28	21.5	No	Normal	х	Х	х	Х
7	Ш	16	Luteal	34	24.5	No	Normal		х		
8	Ш	40	Luteal	27	19.8	No	Normal		х		
105	105 Table S2. Demographics and reproductive information for each endometrial biopsy . Normal time										
106	6 between menses equals 24–38 days; n/a: not applicable.										

	PAIRED EUTOPIC AND ECTOPIC TISSUE							
Sample	Endometriosis	AFS	Cycle	Age	BMI	Smoking	Time interval between	Ectopic lesion
	stadium	score	phase	(years)	(kg/m ²)		menses	-
1	II	8	Follicular	30	23.6	No	Normal	Superficial lesion; pararectal
2	II	8	Follicular	25	29.4	No	Normal	Superficial lesion; pararectal
3	II	10	Follicular	36	21.5	No	Normal	Superficial lesion; uterovesicular
126	Table S	53. Demo	graphics an	d reprod	uctive info	ormation of	the paired eutopic and ect	opic tissues.
127	Norma	l time bet	ween mense	s equals 2	24–38 days		1 1	1
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TAQMAN GENE EXPRESSION ASSAYS									
Gene name	Assay ID	RefSeq ID	Exon boundary	Assay location	Amplicon length				
TRPA1	Hs00175798_m1	NM_007332.2	2–3	441	124				
TRPV1	Hs00218912_m1	NM_018727.5	8–9	1540	94				
TRPV2	Hs00901640_m1	NM_016113.4	14–15	2591	72				
TRPV3	Hs01000530_m1	NM_001258205.1	14–15	2128	121				
TRPV4	Hs01099348_m1	NM_001177428.1	2–3	586	65				
TRPV5	Hs00219765_m1	NM_019841.4	7–8	1176	96				
TRPV6	Hs01114089_g1	NM_018646.4	7–8	1138	59				
TRPM1	Hs00931865_m1	NM_001252020.1	2–3	448	91				
TRPM2	Hs01066071_m1	NM_003307.3	9–10	1419	74				
TRPM3	Hs00257553_m1	NM_001007470.1	3–4	356	83				
TRPM4	Hs00214167_m1	NM_001195227.1	12–13	1852	60				
TRPM5	Hs00175822_m1	NM_014555.3	18–19	1795	79				
TRPM6	Hs01019353_m1	NM_001177310.1	13–14	1555	68				
TRPM7	Hs00918956_m1	NM_017672.4	38–39	5745	148				
TRPM8	Hs00375481_m1	NM_024080.4	2–3	157	106				
TRPC1	Hs00608195_m1	NM_001251845.1	6–7	1097	137				
TRPC3	Hs00162985_m1	NM_001130698.1	4–5	1416	58				
TRPC4	Hs00211805_m1	NM_001135955.1	8–9	2312	76				
TRPC5	Hs00202960_m1	NM_012471.2	6–7	2623	75				
TRPC6	Hs00989190_m1	NM_004621.5	3–4	1558	87				
TRPC7	Hs00220638_m1	NM_001167576.1	4–5	1458	55				
ACTB	Hs01060665_g1	NM_001101.3	2–3	208	63				
GAPDH	Hs02758991_g1	NM_001256799.2	6–7	752	93				
HPRT1	Hs02800695_m1	NM_000194.2	2–3	297	82				
PGK-1	Hs00943178_g1	NM_000291.2	5–6	697	73				
ТРВ	Hs00427620_m1	NM_001172085.1	2–3	578	91				

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Table S4. List of used Taqman genes (Applied Biosystems).