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Supplementary Materials for

Disruption in ACTL7A causes acrosomal ultrastructural defects in human and mouse sperm as a novel male factor inducing early embryonic arrest

Aijie Xin, Ronggui Qu, Guowu Chen, Ling Zhang, Junling Chen, Chengqiu Tao, Jing Fu, Jianan Tang, Yanfei Ru, Ying Chen, Xiandong Peng, Huijuan Shi*, Feng Zhang*, Xiaoxi Sun*

*Corresponding author. Email: xiaoxi_sun@aliyun.com (X.S.); zhangfeng@fudan.edu.cn (F.Z.); shihuijuan2011@163.com (H.S.)

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Other Supplementary Material for this manuscript includes the following:

(available at advances.sciencemag.org/cgi/content/full/6/35/eaaz4796/DC1)

Movies S1 to S4

Supplementary Materials

Supplementary figures



Figure S1. Sperm morphology of the affected brothers under light microscopy. Sperm samples of brother 1 (A) and brother 2 (B) were spread over slides and dried at room temperature, fixed in 95% ethanol for Papanicolaou stain. Spermatozoa were then scored by $\times 100$ oil-immersion bright-field objective according to the standard of the fifth edition of the WHO guidelines. A1, A2, B1 and B2 correspond to enlarged spermatozoa with normal morphology. Photo credit: Aijie Xin, Fudan University.



Figure S2. Day 3 embryos of one brother couple by ICSI treatment.

Seven oocytes were fertilized with sperm of brother 1 by ICSI. The images A, B and C showed the Day 3 embryos. They all arrested at 2-cell, 4-cell or 5-cell stage. Photo credit: Jing Fu, Fudan University.



Figure S3. *Actl7a* expression was significantly decreased in testes of *Actl7a*^{KI/KI} mice.

The relative mRNA level of *Actl7a* in testes of the *Actl7a*-mutated mice (*Actl7a*^{KI/KI}) was significantly decreased by approximately 30% analyzed by one-way ANOVA. Nine testicular samples of each genotype group were used for biological replicates. Three duplicates of each sample were performed by RT-qPCR assays. *Gapdh* served as a reference gene. * P < 0.05; NS, not significant. Data are represented as the mean \pm SEM.





(A) Motility parameters of spermatozoa in the cauda region were compared among the mice with wile-type (WT), $Actl7a^{+/KI}$ or $Actl7a^{KI/KI}$ genotypes. Three sperm samples in each group were used for biological replicates. The semen parameters showed no significant difference among the three genotypes. Data are represented as the mean \pm SEM. (B) The morpha and size of testes from $Actl7a^{KI/KI}$ mice showed no major change compared to those of WT and $Actl7a^{+/KI}$ mice. (C) Testis weight presented no significant difference among WT, $Actl7a^{+/KI}$ and $Actl7a^{KI/KI}$ mice. Five mice in each group were used for biological replicates. Data are represented as the mean \pm SEM. (D) H&E stained cross sections of testis and cauda epididymis of WT and $Actl7a^{KI/KI}$ mice. The histomorphology of seminiferous tubules and caudal epididymis in $Actl7a^{KI/KI}$ mice appeared similar to those of the WT male mice. Scale bars: 100 µm. Photo credit: Aijie Xin, Fudan University.



Figure S5. Spermatozoa from the *Actl7a*^{KI/KI} mice failed to fertilize normal mouse oocytes by IVF and ICSI. Photo credit: Aijie Xin, Fudan University.



Figure S6. The thickness of perinuclear theca in brother and mouse sperm with homozygous *Actl7a* mutation.

(A) Measurement of PT thickness. The four red lines are the measuring area of the subacrosomal layer of the perinuclear theca (SAL-PT); the four blue lines are the measuring area of the postacrosomal sheath of the perinuclear theca (PAS-PT). Taken the average value as the thickness of SAL-PT and PAS-PT. (B) Significant increase of the thickness of SAL-PT and PAS-PT in brother sperm. (C) Significant increase of the thickness of SAL-PT and PAS-PT in sperm from the male mice carrying homozygous *Actl7a* mutation. Photo credit: Aijie Xin, Fudan University.



Figure S7. Genotyping and fertility of the offspring from *Actl7a*^{KI/KI} mice.

(A) The number represented the mice marked with dotted circle in figure 5D. Dotted rectangle indicate the position of *Actl7a* point mutation. (B) The fertility rate presented no significant difference between the offspring of the $Actl7a^{KI/KI}$ male mice and the wild-type (WT) controls. Male, n=5; Female, n=5. Two-tailed Student's *t* test; NS, not significant.

Supplementary tables

	Brother 1	Brother 2	Low reference limits of WHO criteria
Age	29	28	/
Semen volume (ml)	1.8	7.0	1.5 (1.4–1.7)
Sperm concentration (10 ⁶ /ml)	325	162	15 (12–16)
Total sperm (10 ⁶)	585	1134	39 (33–46)
Progressive motility (PR, %)	54.7	68.4	32 (31–34)
Total motility (PR+NP, %)	76.6	78.1	40 (38–42)

Table S1. Semen parameters of the two brothers affected by a homozygous *ACTL7A* mutation.

	Brother 1				Brother 2			
Cycle ID	1	2		3	4	1	2	3
No. of oocytes retrieved	21	19	1	7	19	5	8	7
No. of mature oocytes (M II)	17	17	1	4	19	4	6	5
Method of ART	Later-ICSI	ICSI	ICSI	DIVF	DIVF	IVF	ICSI	DIVF
No. of fertilization (2PN)	9	9	7	6	17	0	3	5
No. of embryo for transfer*	0	0	0	6	11	0	0	4
No. of good quality embryo†	0	0	0	6	9	0	0	3
Live birth	0	0	0	0	2	0	0	1

Table S2. The ART history of two affected couples before the genetic diagnosis ofthis study.

* Embryo for transfer: Grade 2, \geq 5 cells.

[†]Good quality embryos: Grade 1, \geq 7-8 cells.

Abbreviations: ICSI, intracytoplasmic sperm injection; IVF, in vitro fertilization; DIVF, donor sperm in vitro fertilization.

	WT	KI/KI	SrCl ₂	SrCl ₂ + KI/KI
2 PN (%)*	93.0 ± 3.9	0	93.3±4.0	64.4 ± 3.2
2-cell (%) †	86.0 ± 14	/	91.4 ± 2.5	94.0 ± 4.3
4-cell (%) †	84.0 ± 8.0	/	71.8 ± 3.1	84.0 ± 6.4
Morula (%) †	84.0 ± 8.0	/	71.8 ± 3.1	89.8 ± 3.7
Blastula (%) †	79.8 ± 3.9	/	29.9 ± 3.3	49.0 ± 7.4

Table S3. The development rate of mouse embryos in different development stagesafter AOA.

* Percentages of 2PN were calculated from the total number of injected oocytes.

[†] Percentages of 2-cell embryos to blastula were calculated from 2PN.

Table S4. Sequence of single-guide RNA (sgRNA) and the single-stranded oligonucleotide (ssODN).

Material	Sequence (5' - 3')
sgRNA	ACGGGGAGGCTAGACTACGC
ssODN	TATGAGGGTTATCCTTTGCCCAGCATCACGGGGAGGCTAGACTACACAGGT
	TCTGACCTAACGACCTACCTGATGAACCTGATGAACAA

Note: The protospacer adjacent motif (PAM) sequence is shown in red, and the mutated target is shown in blue.

Supplementary movies

Movies S1 to 4. The embryonic development from day 0 to day 3 following ICSI with AOA in the case of brother 2 and his wife.