

Prmt5 is required for germ cell survival during spermatogenesis in mice

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Figure S1

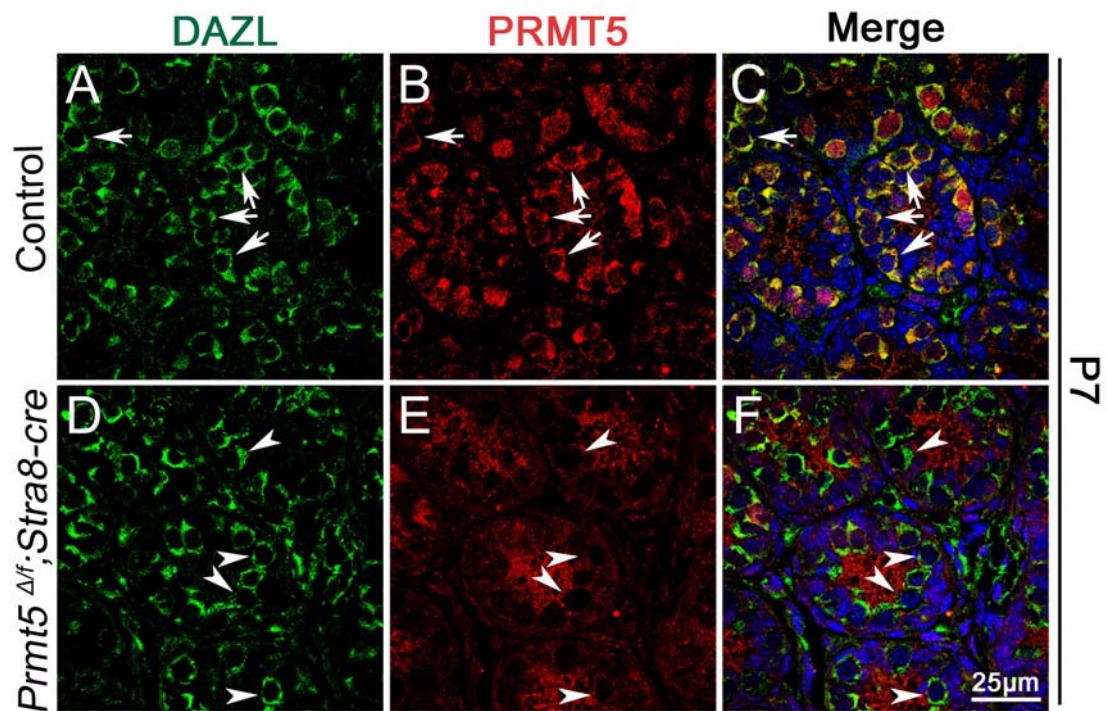


Figure S1. Prmt5 was inactivated as early as P7 in germ cells of *Prmt5*^{Δ/f}; *Stra8-Cre* testes. The expression of Prmt5 (red) in control and *Prmt5*^{Δ/f}; *Stra8-Cre* testes was examined by Immunofluorescence at P7. Germ cells were labeled with antibody against Dazl (green, white arrows). In control testes, Prmt5 was detected in cytoplasm of germ cells (B, C, white arrows), whereas no Prmt5 was detected in germ cells of *Prmt5*^{Δ/f}; *Stra8-Cre* testes (E,F, white arrowheads).

Figure S2

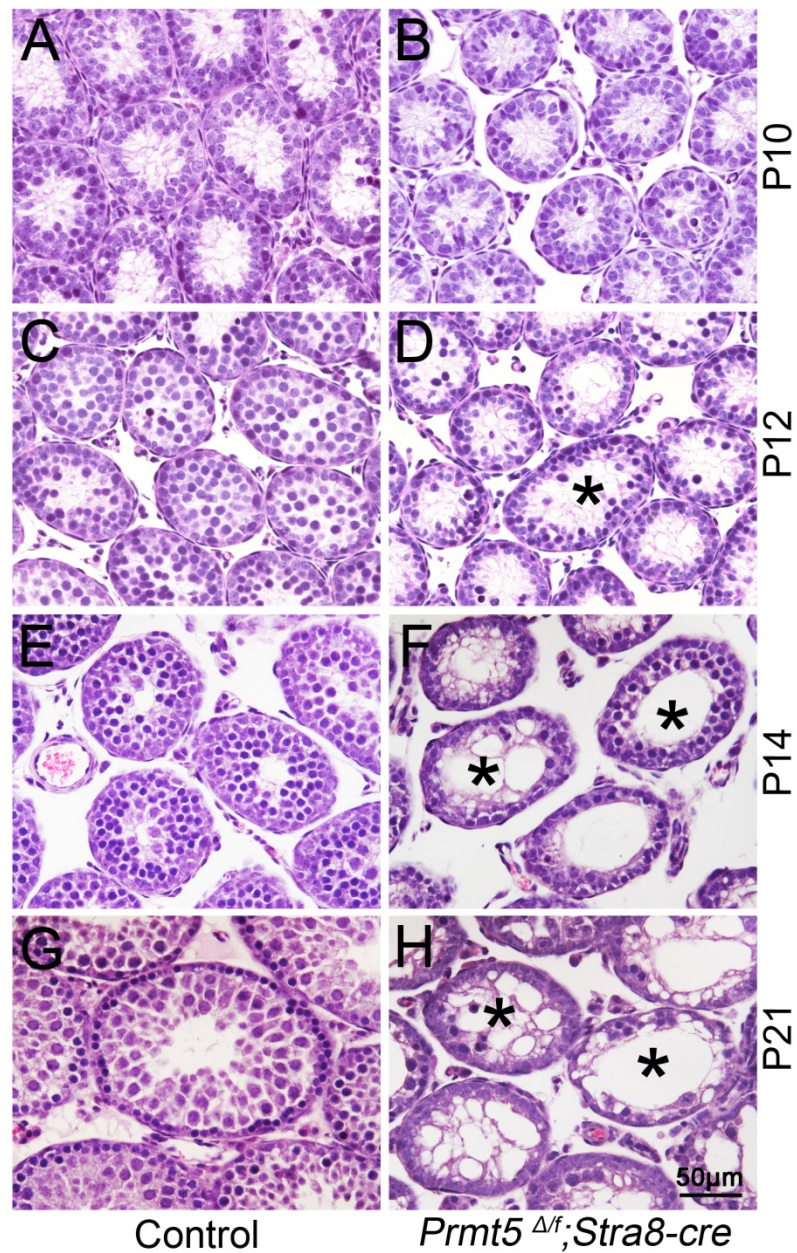


Figure S2. The defect of germ cell development was first observed in *Prmt5*^{Δf}; *Stra8-Cre* mice at P12. The results of H&E staining showed that the testes from *Prmt5*^{Δf}; *Stra8-Cre* mice (B) was grossly normal compared to control testes (A) at P10. Aberrant seminiferous tubules (asterisks) were first noted in *Prmt5*^{Δf}; *Stra8-Cre* testes (D, asterisks) at P12, and atrophic tubules (asterisks) were observed in *Prmt5*-deficient testes at P14 (F, asterisks) and P21 (H, asterisks).

Figure S3

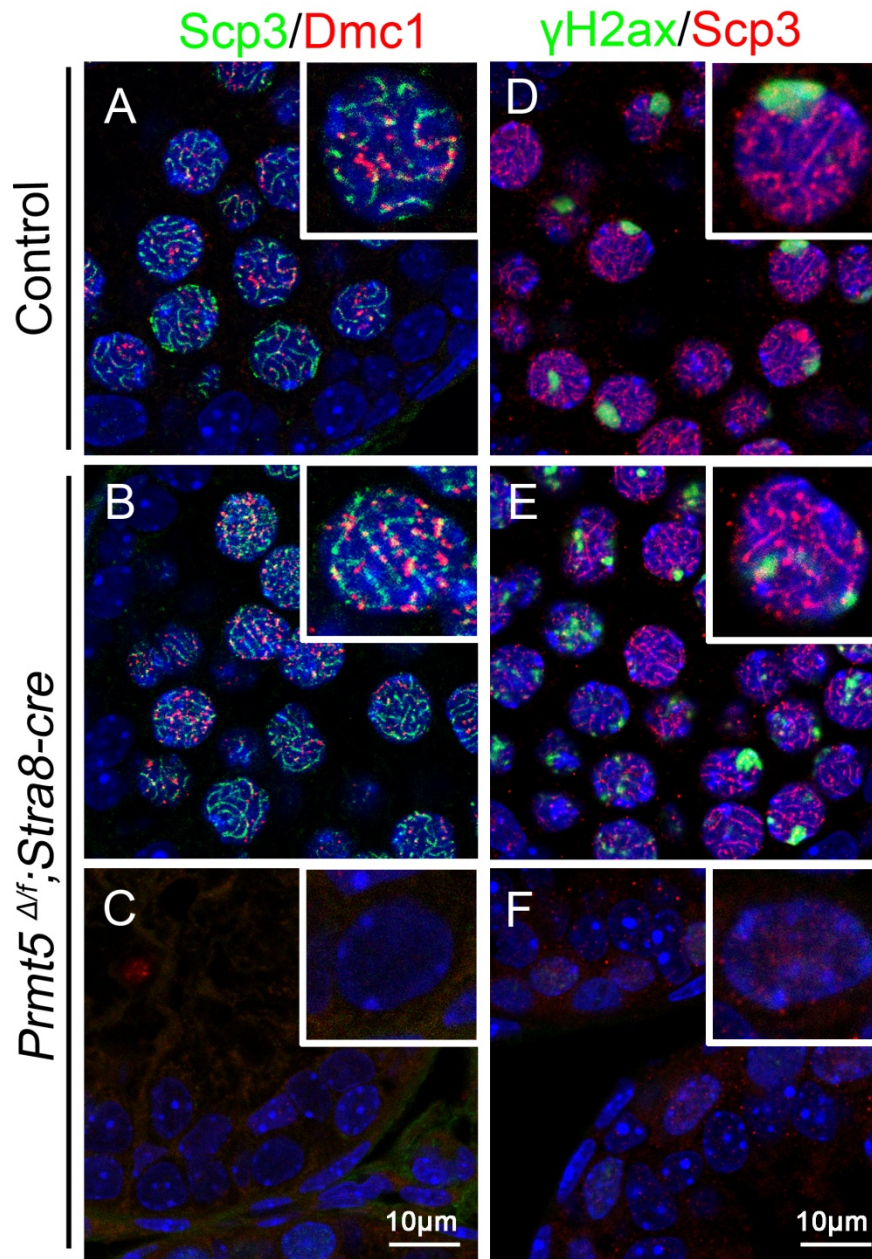


Figure S3. Immunofluorescence of Dmc1 and γ H2AX. The expression of Dmc1 and γ H2AX was examined by immunofluorescence. In control testes, Dmc1 protein (A, red) was detected in most of spermatocytes at P12, and co-localized with Scp3 (A, green). γ H2AX (green) was mainly observed in sex body (D) in control testes. In *Prmt5^{Δf}; Stra8-Cre* testes, Dmc1 protein (B, red) was only detected in a small number of germ cells, which was co-localized with Scp3 (B, green). γ H2AX (green) was also detected in a small number of germ cells of *Prmt5^{Δf}; Stra8-Cre* testes, whereas multiple foci were noted (E). Dmc1 (C) and γ H2AX (F) protein was absent in most of germ cells from *Prmt5^{Δf}; Stra8-Cre* testes.

Figure S4

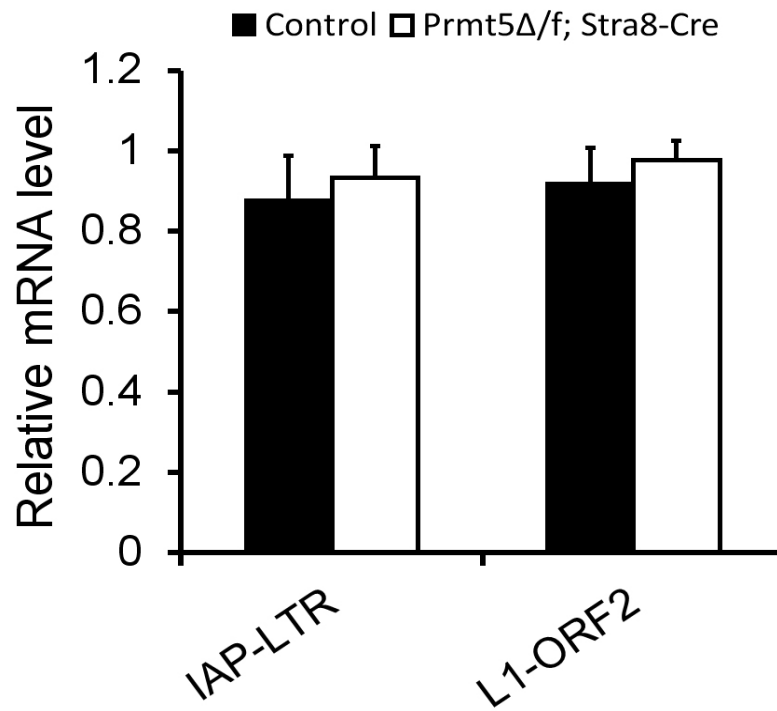


Figure S4. Loss of *Prmt5* did not result in up-regulation of transposable elements in germ cells. The expression of TEs in testes at P10 was examined by real-time PCR. Compared to control testes, the mRNA levels of *IAP-LTR* and *L1-ORF2* in *Prmt5* ^{Δ/f} ; *Stra8-Cre* testes were not increased.

Table S1

Gene	Forward primer sequence	Reverse primer sequence
<i>Prmt5</i>	TGGTGGCATAACTTTCGGACT	TCCAAGCCAGCGGTCAAT
<i>Dazl</i>	ATGTCTGCCACAACCTCTGAG	CTGATTTTCGGTTTCATCCATCCT
<i>Vasa</i>	CTAGGAAGACCAAATAGTGAATCTGAC	TCCAGAACCTGTTACTACTTCTTCATT
<i>Plzf</i>	CCCAGTTCTCAAAGGAGGATG	TTCCCACACAGCAGACAGAAG
<i>Stra8</i>	CTGTTGCCGGACCTCATGG	TCACTTCATGTGCAGAGATGATG
<i>Dmc1</i>	CCCTCTGTGTGACAGCTCAAC	GGTCAGCAATGTCCCGAAG
<i>Spo11</i>	CGTGGCCTCTAGTTCTGAGGT	GCTCGATCTGTTGTCTATTGTGA
<i>Rad51</i>	GTCCACAGCCTATTTACGGT	ACAGCCTCCACTGTATGGTAAC
<i>Rec8</i>	CTACCTAGCTTGCTTCTTCCCA	GCCTCTAAAAGGTGTCGAATCTG
<i>Syp3</i>	AGAAATGTATACCAAAGCTTCTTTCAA	TTAGATAGTTTTTCTCCTTGTTCTCA
<i>Syp1</i>	AAGTTTGATTCTAAAACAACCTCCTCA	ACTCTTTTTAGTTGGTGTCTTCACTGT
<i>Caspas3</i>	GCTGACTTCCTGTATGCTTAC	ATCCGTTGCCACCTTCC
<i>L1-ORF2</i>	CGGGAGACAGCACATACTAGCA	TCTCAAAGAACCAACTCCTCGT
<i>IAP-LTR</i>	ACATTCGCCGCCACAAGA	TAGTCGTAAATACCCTTGGCTCAT

Table S1. The primers used for real-time PCR analysis.