

SI Correction

PLANT BIOLOGY

Correction to Supporting Information for “Recruitment of the NineTeen Complex to the activated spliceosome requires AtPRMT5,” by Xian Deng, Tiancong Lu, Lulu Wang, Lianfeng Gu, Jing Sun, Xiangfeng Kong, Chunyan Liu, and Xiaofeng Cao, which appeared in

issue 19, May 10, 2016, of *Proc Natl Acad Sci USA* (113:5447–5452; first published April 25, 2016; 10.1073/pnas.1522458113).

The authors note that Fig. S6 appeared incorrectly. The SI has been corrected online.

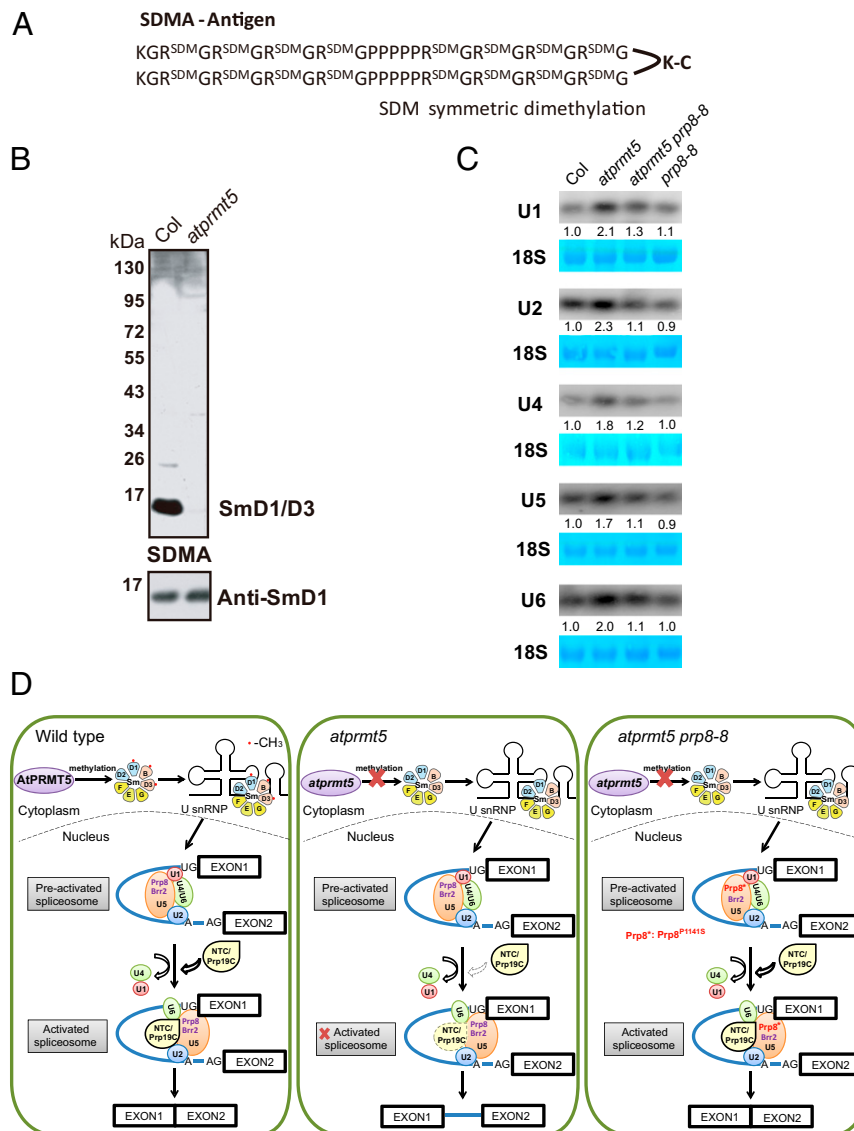


Fig. S6. Preparation of anti-SDMA antibody, U snRNAs analysis, and working model. (A) Low complex GAR motifs harboring symmetric dimethylarginines (SDMA). (B) Total proteins extracted from Col and *atprmt5* mutants were immunoblotted with anti-SDMA and anti-SmD1 antibodies. (C) Levels of U1, U2, U4, U5, and U6 snRNAs are increased in *atprmt5* mutants and restored in *atprmt5 prp8-8*. 18S rRNAs were used as a loading control. (D) Model for the role of AtPRMT5 in Prp19C/NTC complex recruitment to the spliceosome during pre-mRNA splicing.

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