



Y357

Mmu CELLVWY GDEYQQELGI

Rat RELLVWY GDEYQQELGI

KO516U GKSSIGPAGSSGQVASFWSGMGMSMAKSWASSGEAR.

KO516L RELLVWY GDEYQQELGI

KO8 SGLVWG.

KO2 RASGLVWG.

Hsa CELLVWY GDEYQQELGI

Dre QEFRVWY ADEYAQGLGA

Bfl TELMVWY GNEYAKDLGI

Cin TELLVWY GDQYAQSLGI

Spu RELLVYY GEQYAKDLGI

Nve TELLVWY GDKYANDLGI

Hsa- *Homo sapiens*Dre- *Danio rerio*Bfl- *Branchiostoma floridae*Cin- *Ciona intestinalis*Spu- *Strongylocentrotus purpuratus*Nve- *Nematostella vectensis*

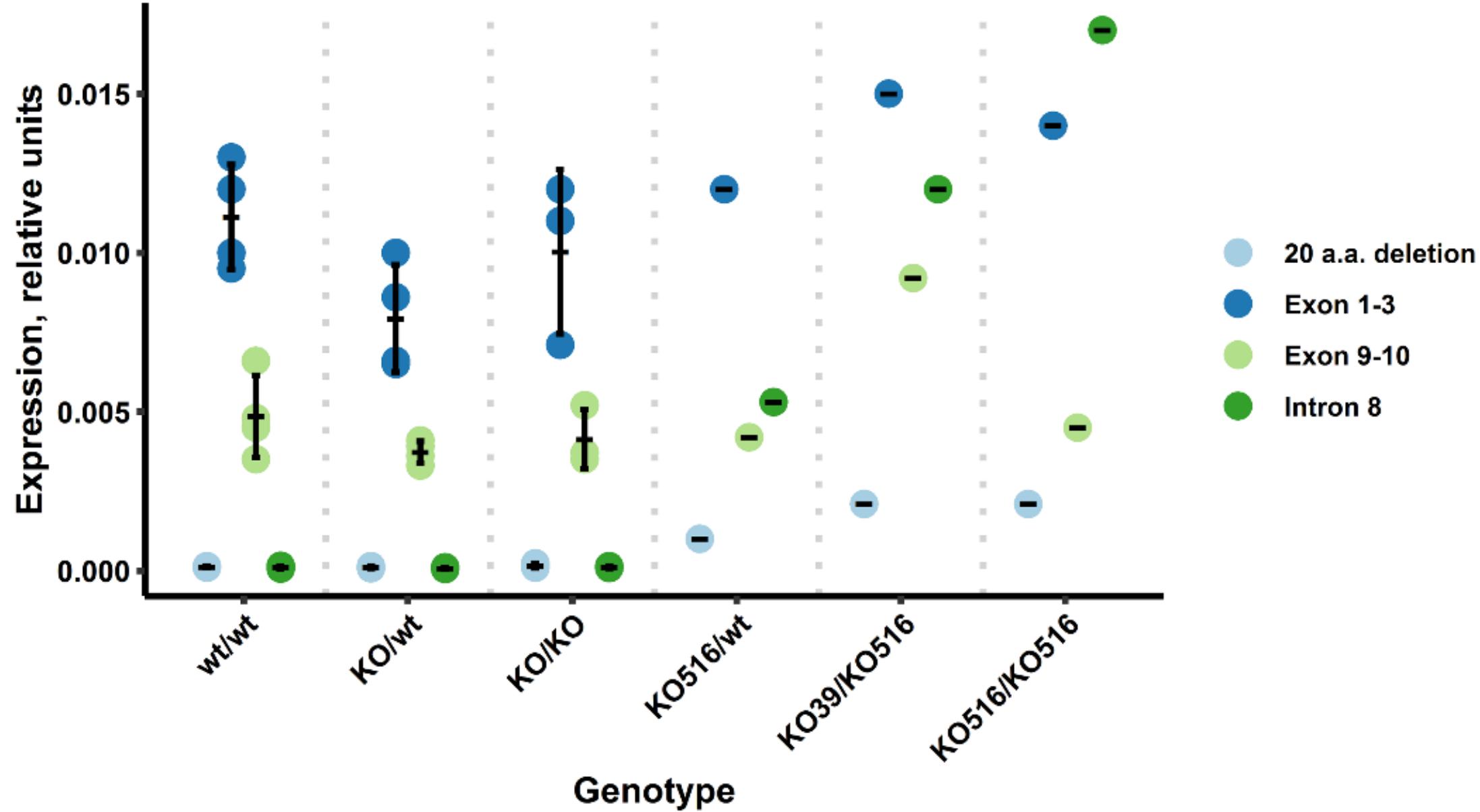
Mutant founders:

KO516 - ♀ deletion 516 bp

KO39 - ♂ deletion 39 bp

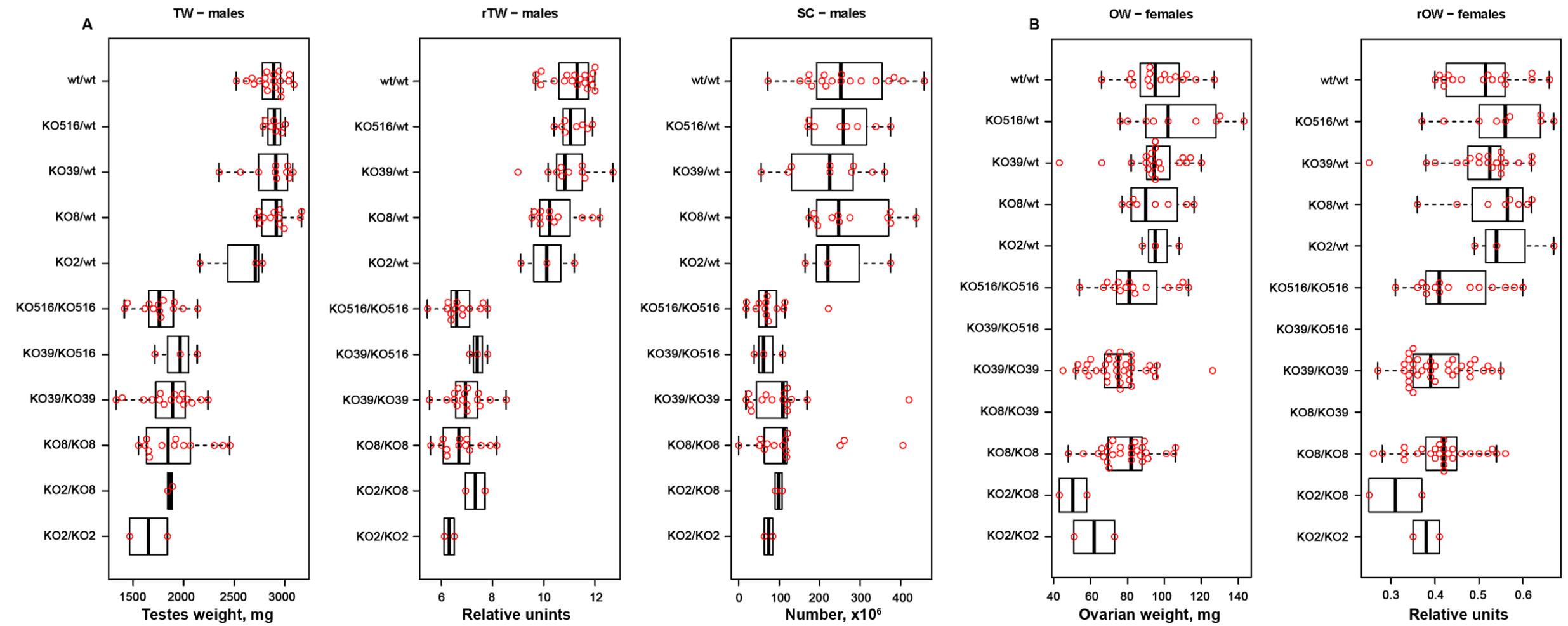
KO8 - ♂ deletion 8 bp

KO2 - ♀ deletion 2 bp

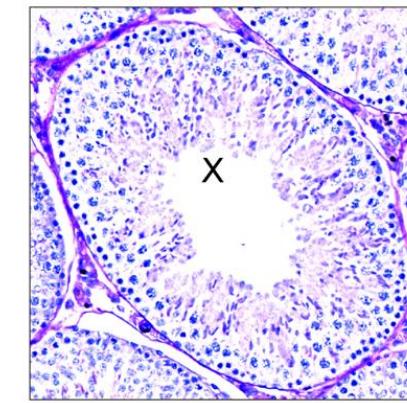
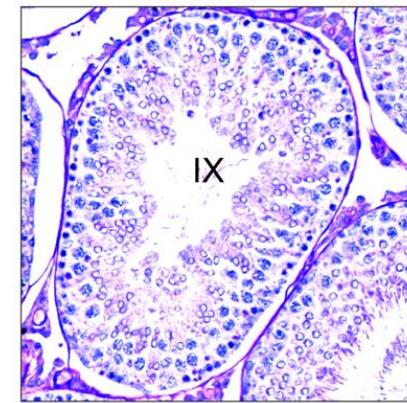
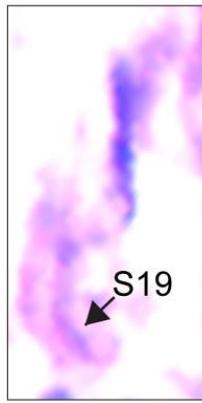
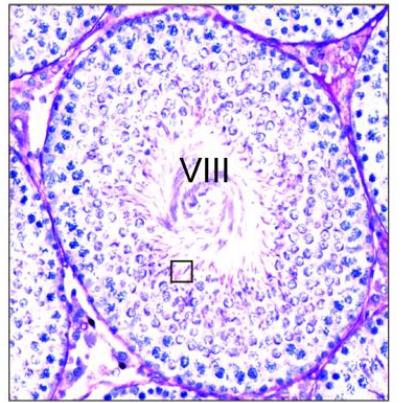


Supplemental Fig. S2

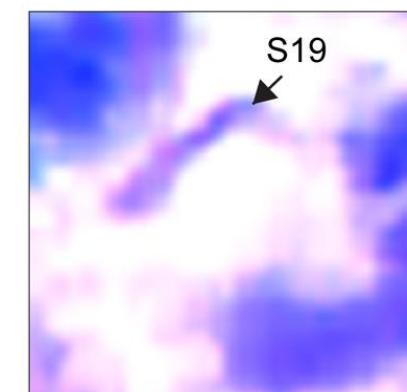
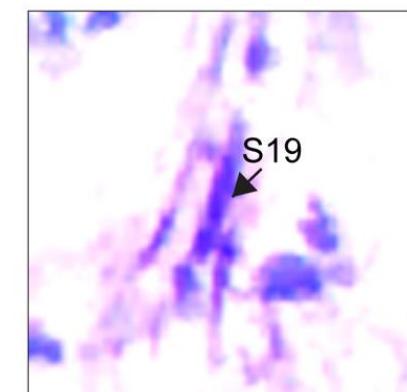
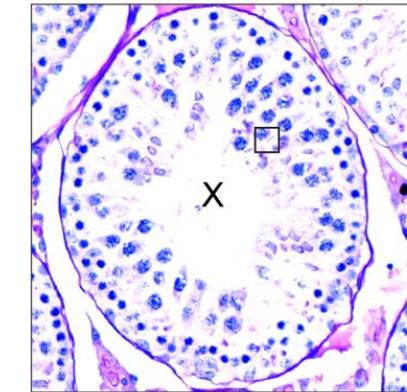
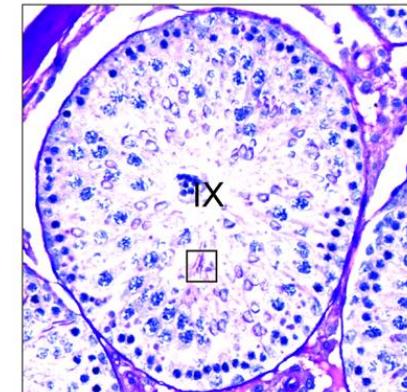
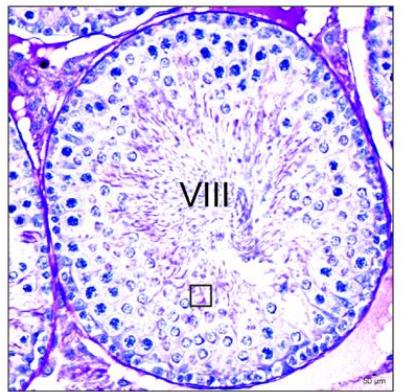
Supplemental Fig. S3



Prdm9^{wt/wt}

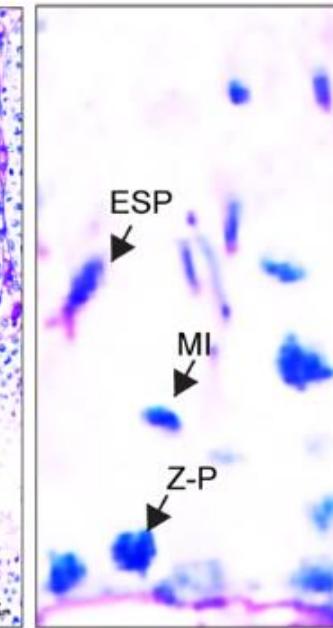
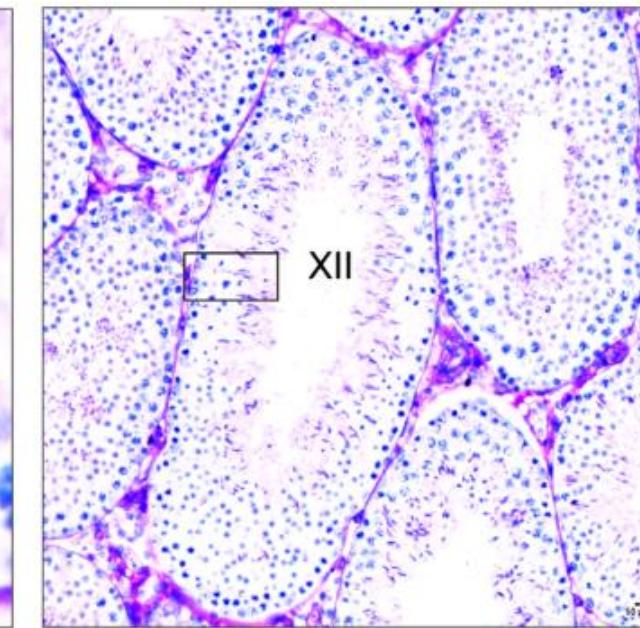
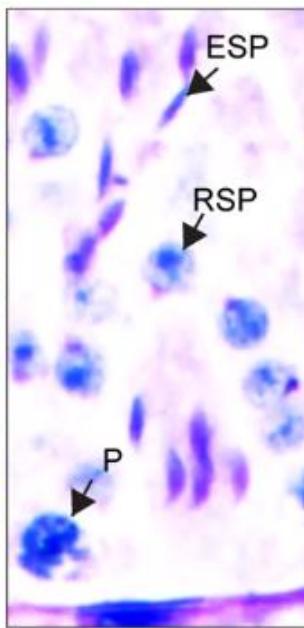
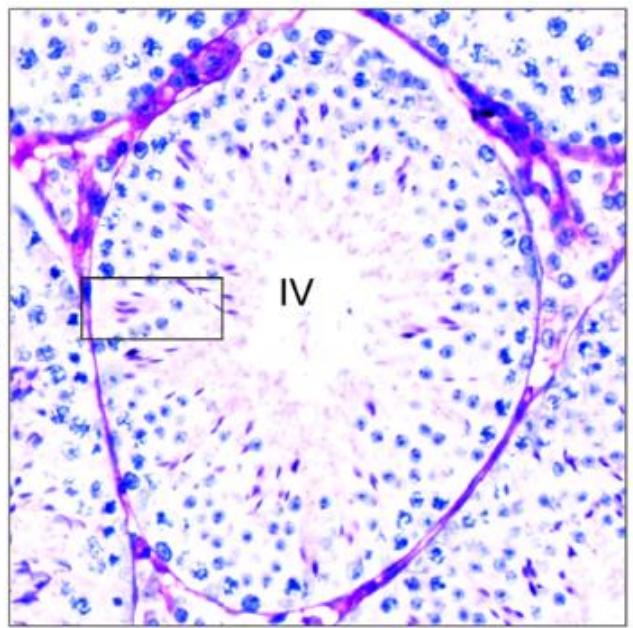


Prdm9^{KO/KO}

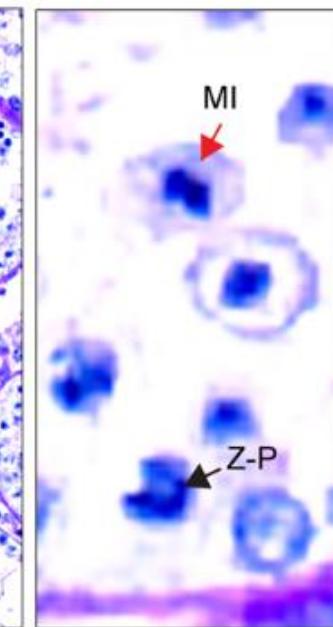
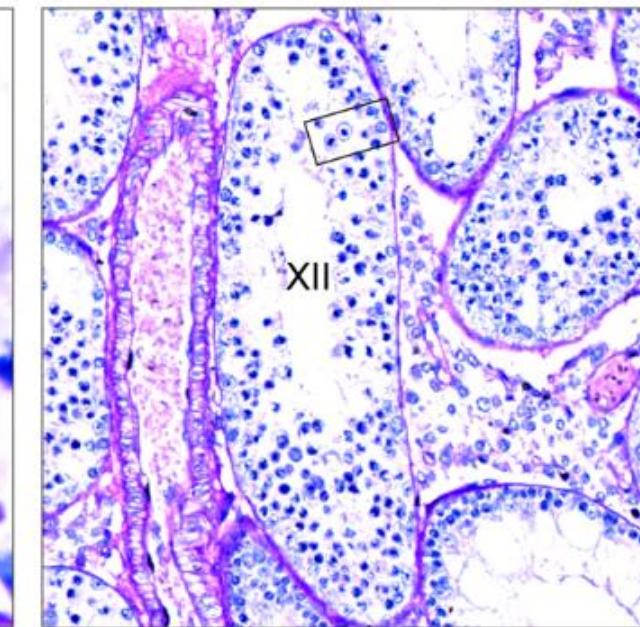
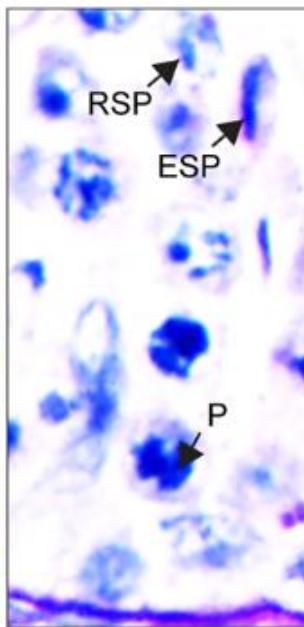
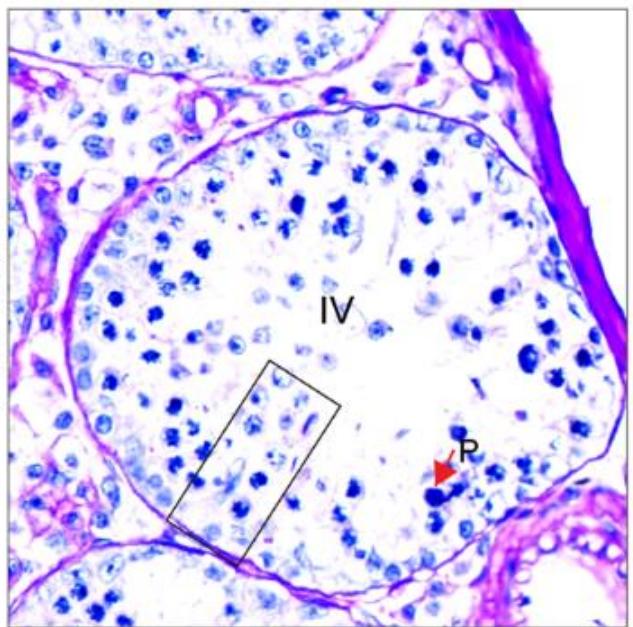


Supplemental Fig. S4

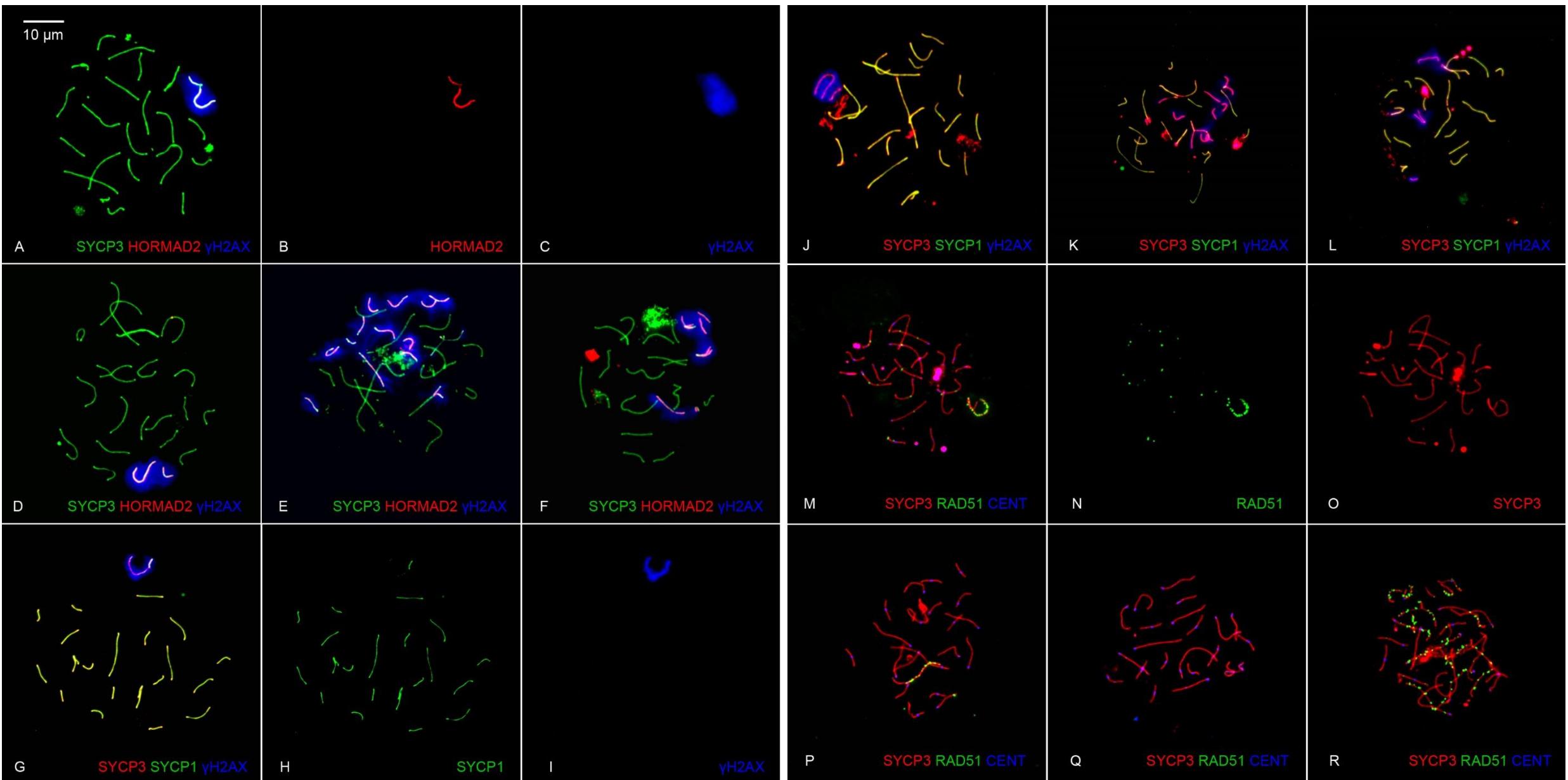
PWD-*Prdm9*^{wt/wt}



PWD-*Prdm9*^{ko/ko}

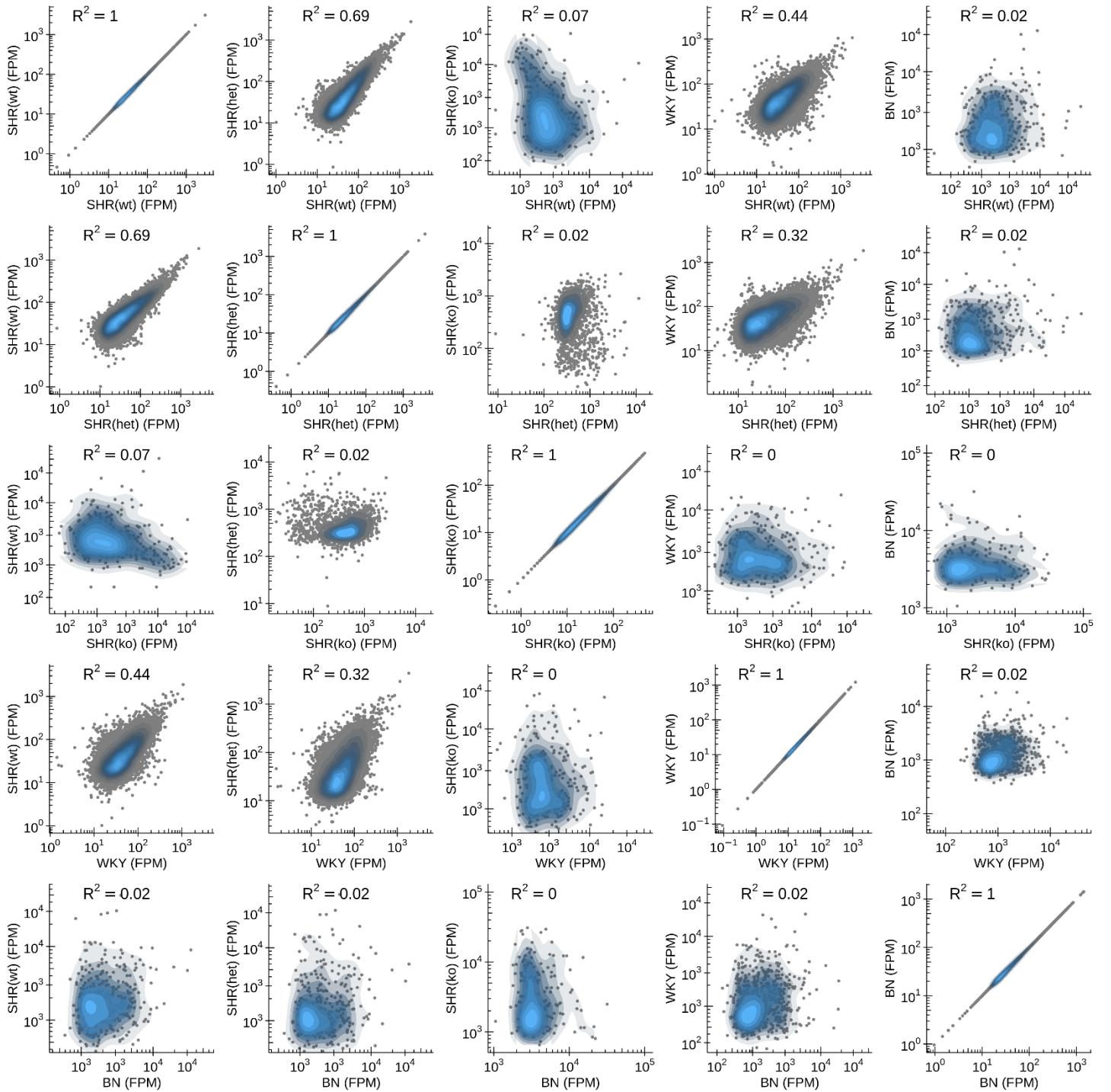


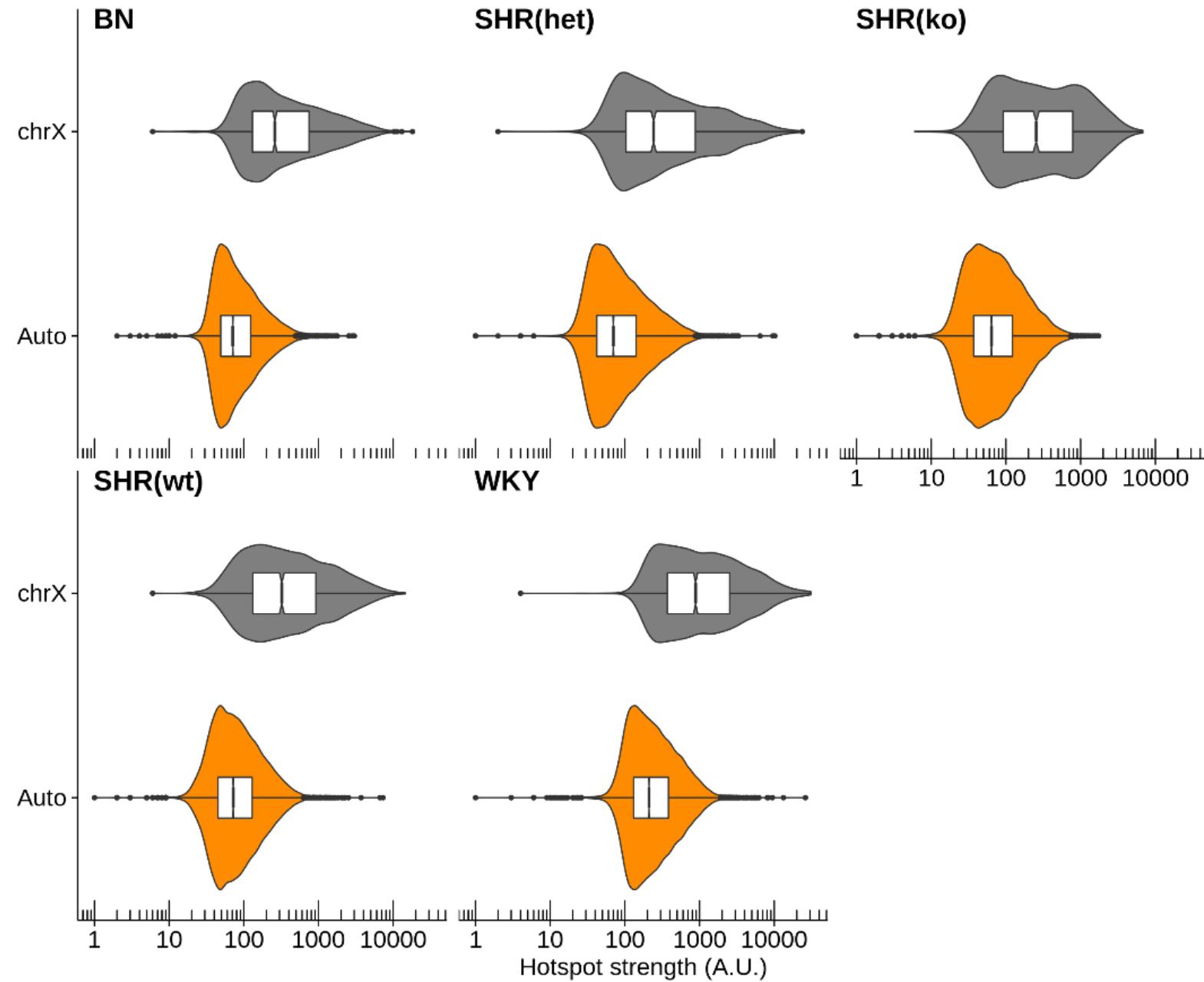
Supplemental Fig. S5

10 μ m

Supplemental Fig. S6

Supplemental Fig. S7





Supplemental Fig. S8