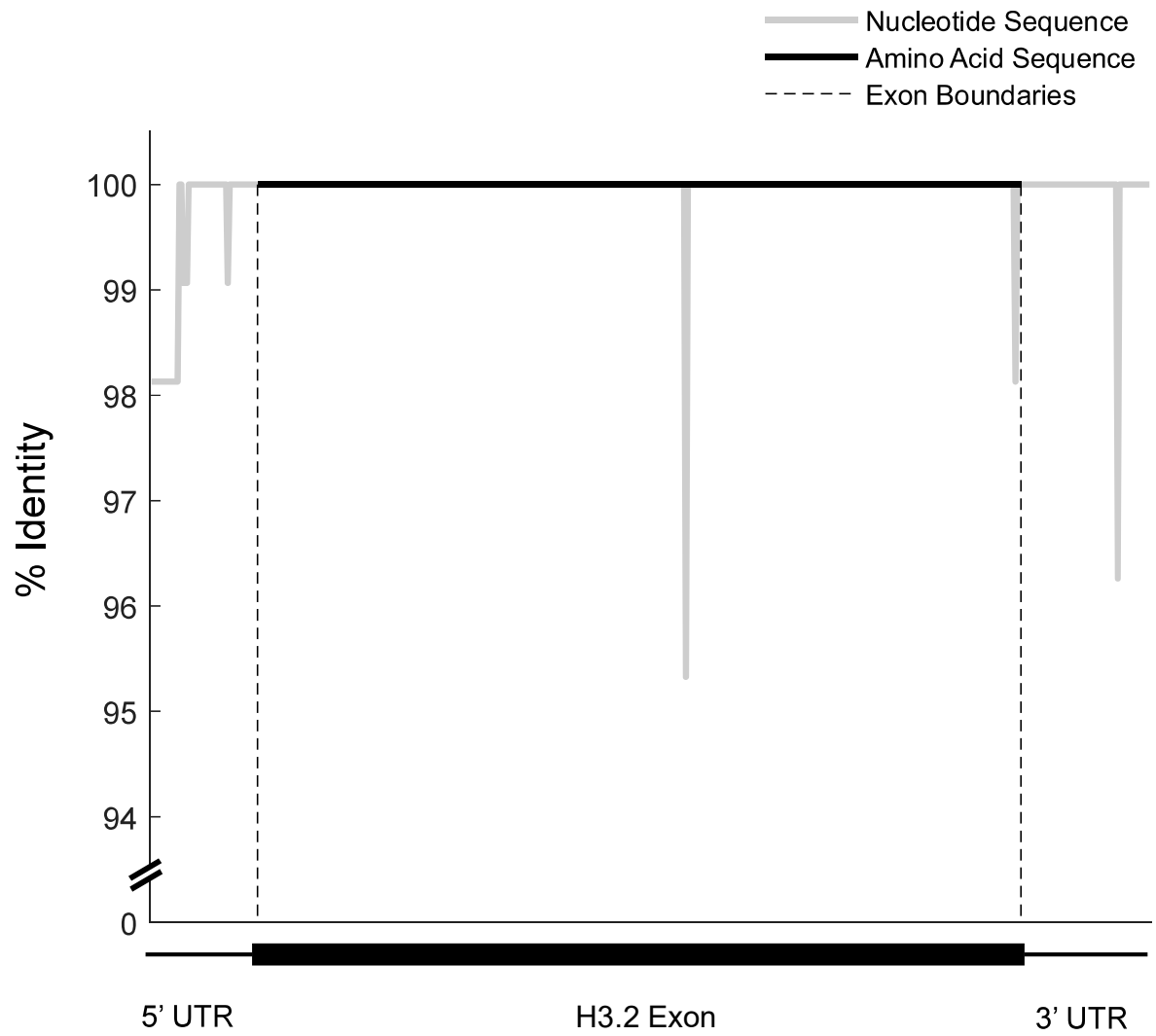
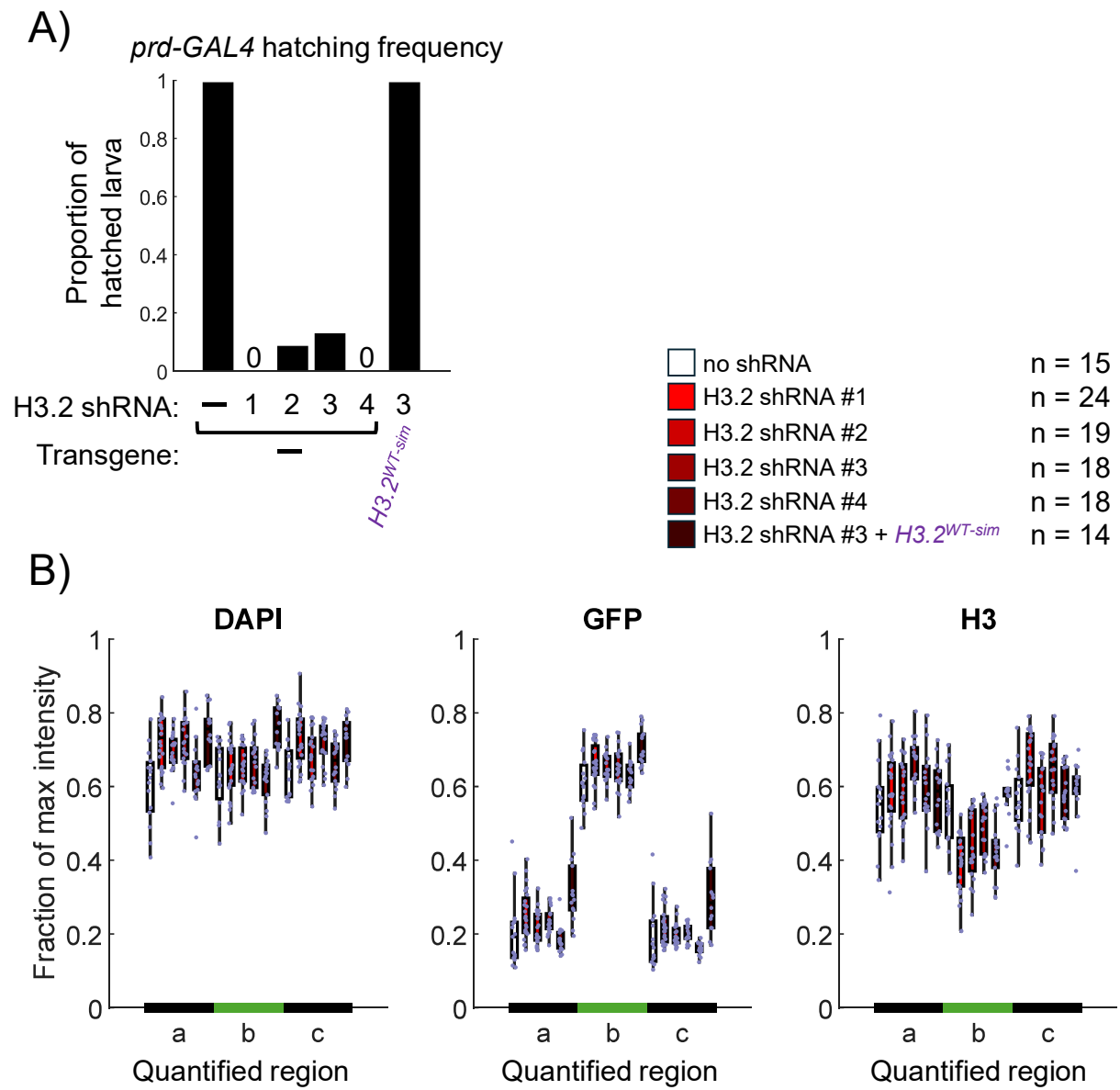
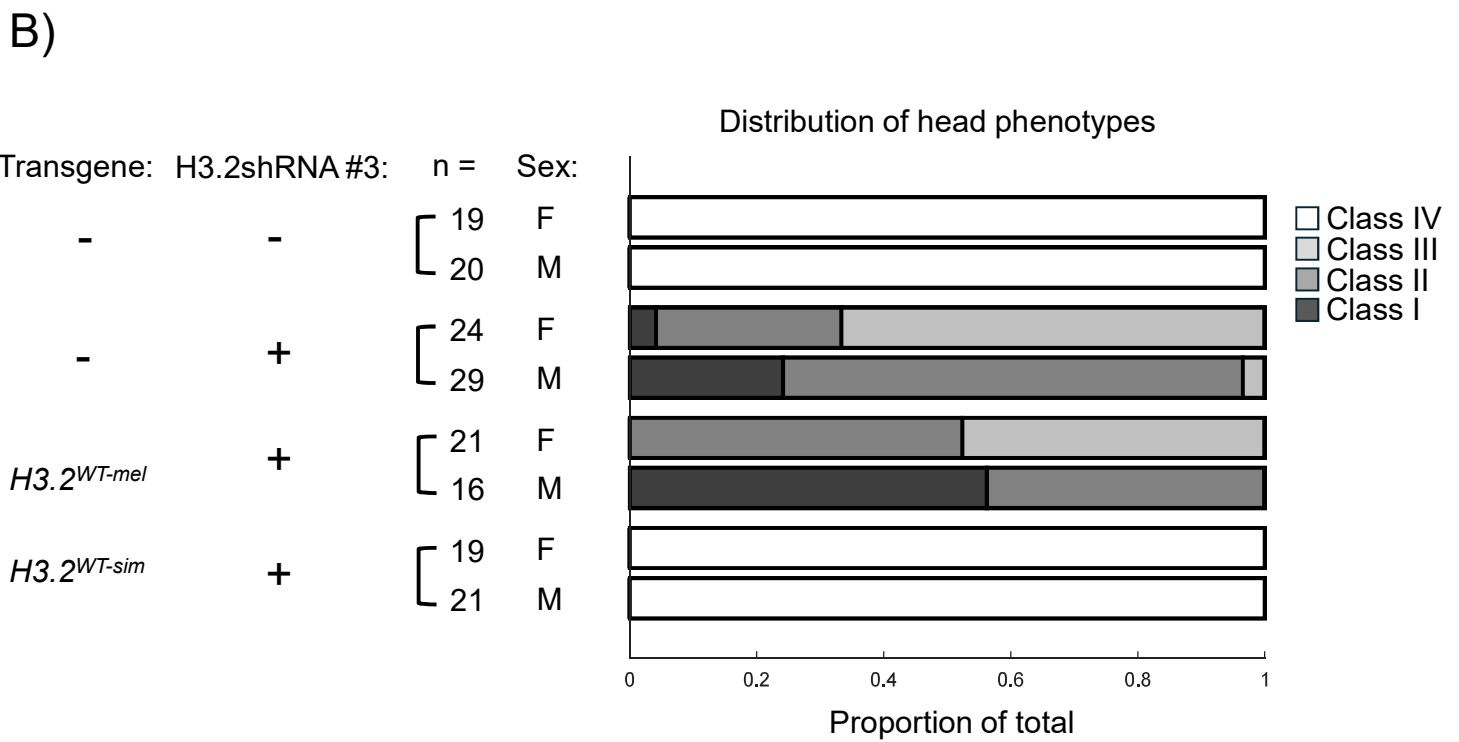
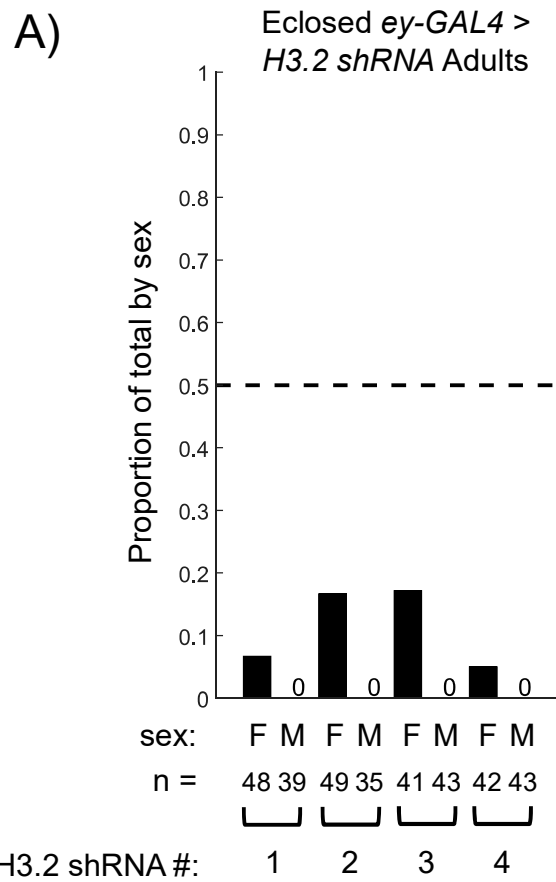


Sequence identity across 107 endogenous H3.2 copies

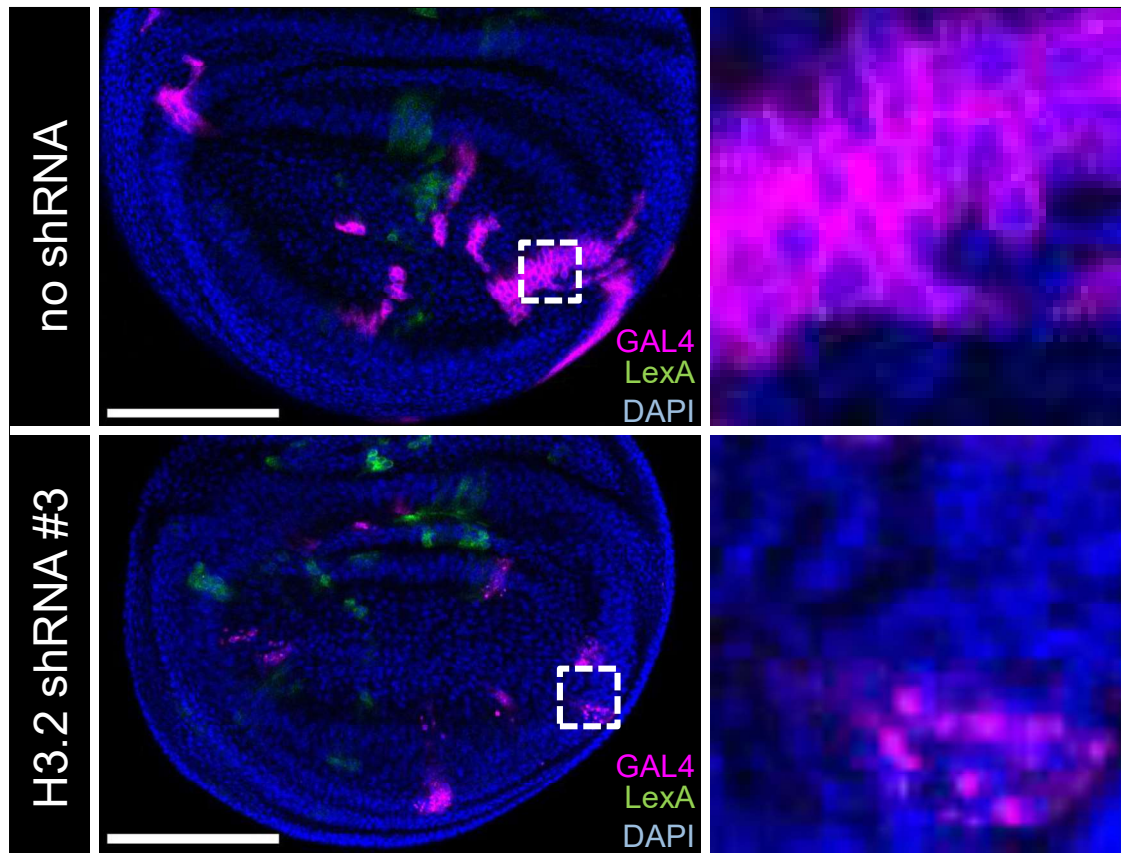




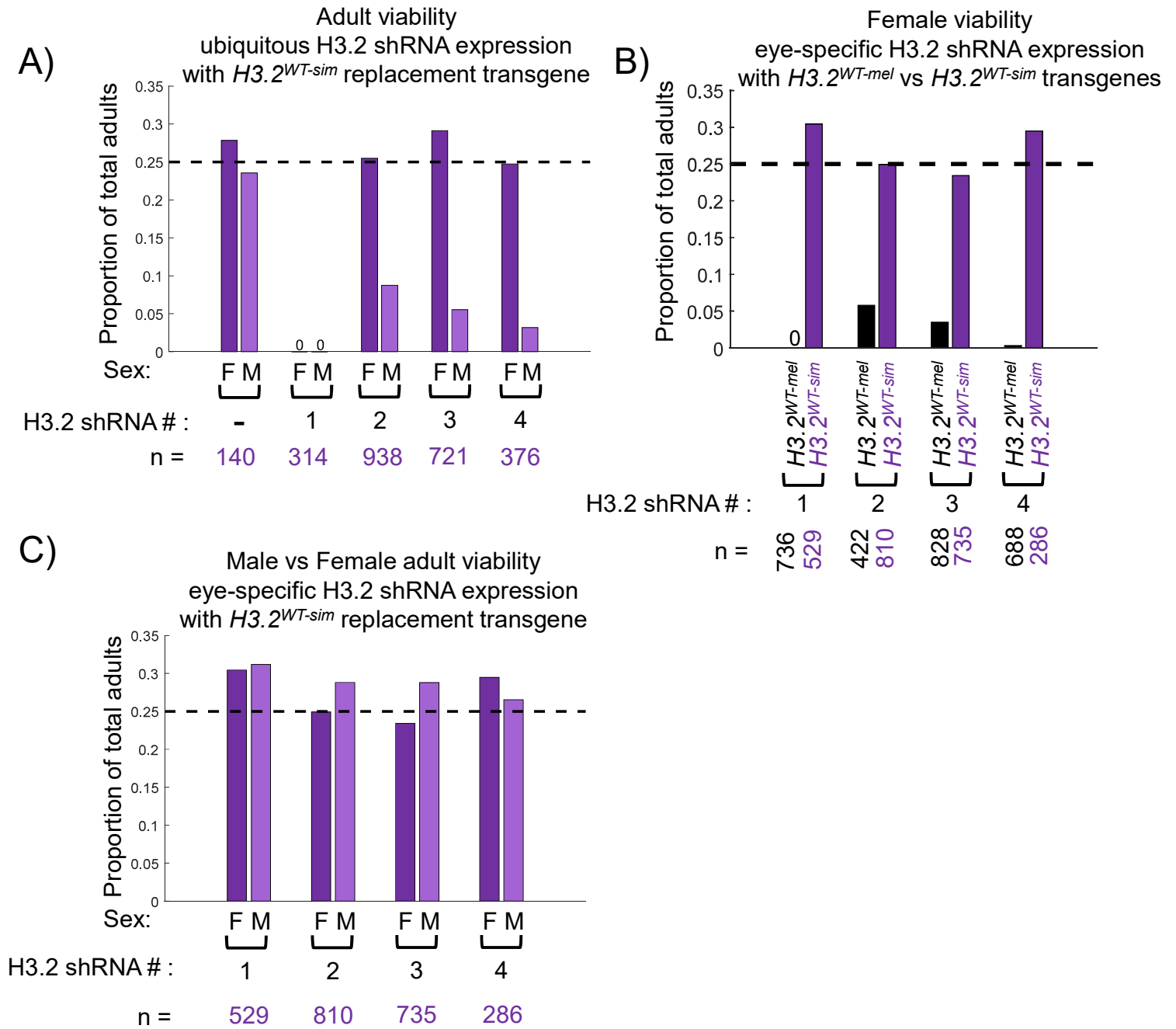
Supplemental Fig. 2



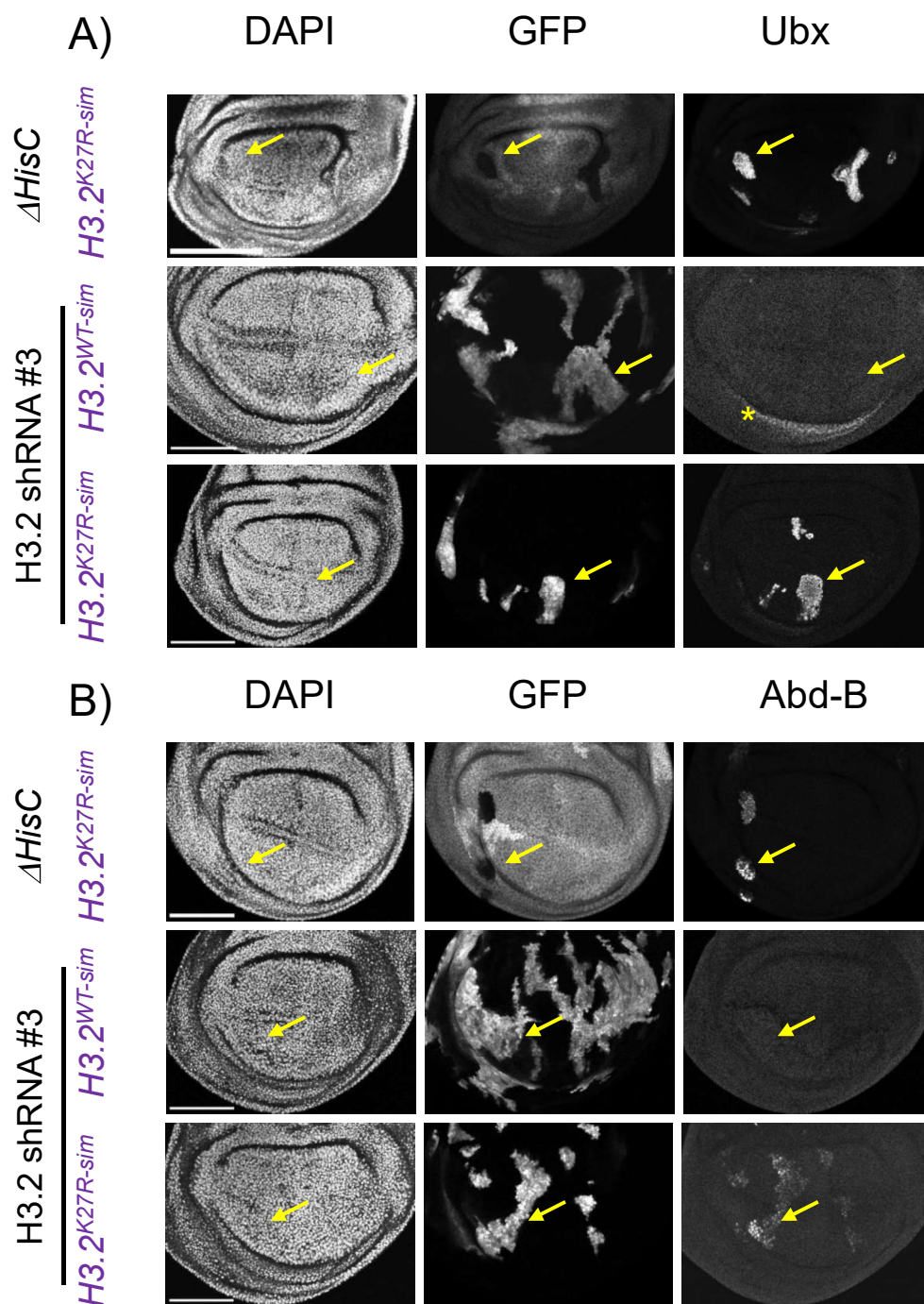
Supplemental Fig. 3



Supplemental Fig. 4



Supplemental Fig. 5



Supplemental Fig. 6