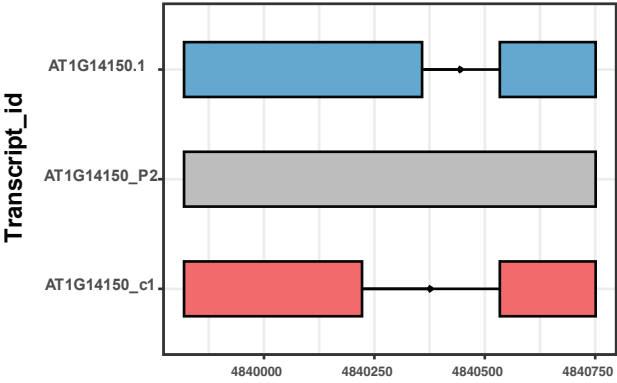
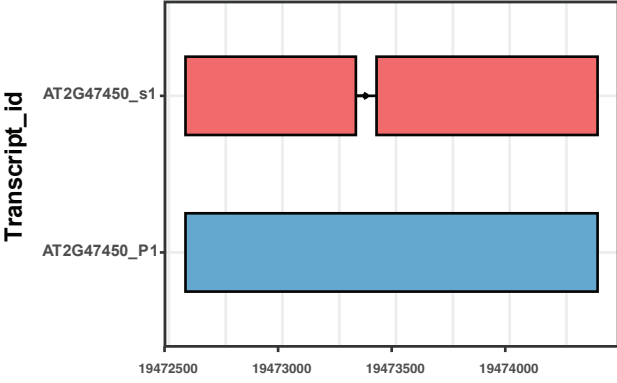


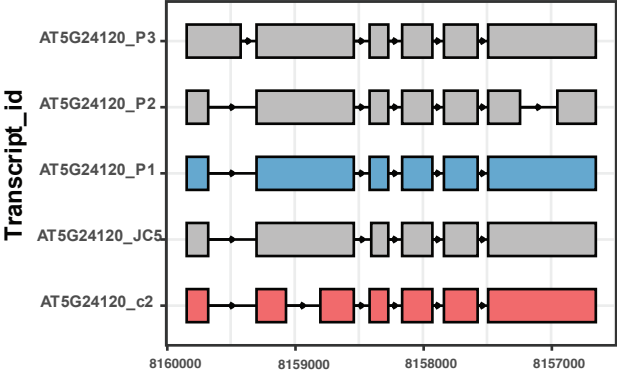
(a)



(b)



(c)



S3 Figure. Transcript isoforms. Splice variants models for PNSL2, CHAOS and SIG5 (a, b,c, respectively). Total transcript models according to the AtRTD2 reference database (Zhang et al., 2017). Transcript models undergoing isoform switch with light exposure are represented in red and green. Transcripts that do not undergo isoform switch are represented in gray. Arrows indicate directionality of transcripts, from N terminal to the C terminal in the corresponding proteins.