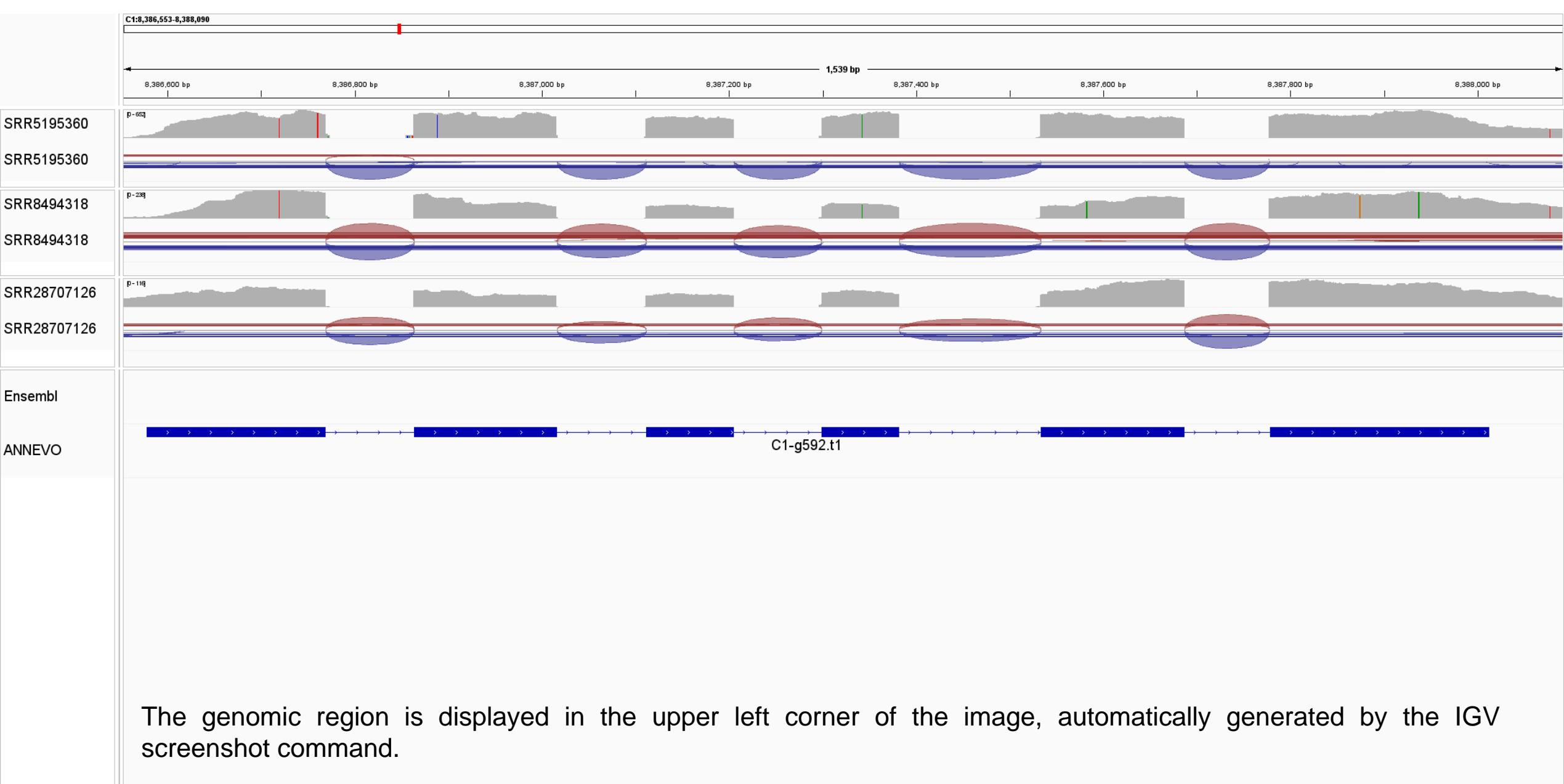
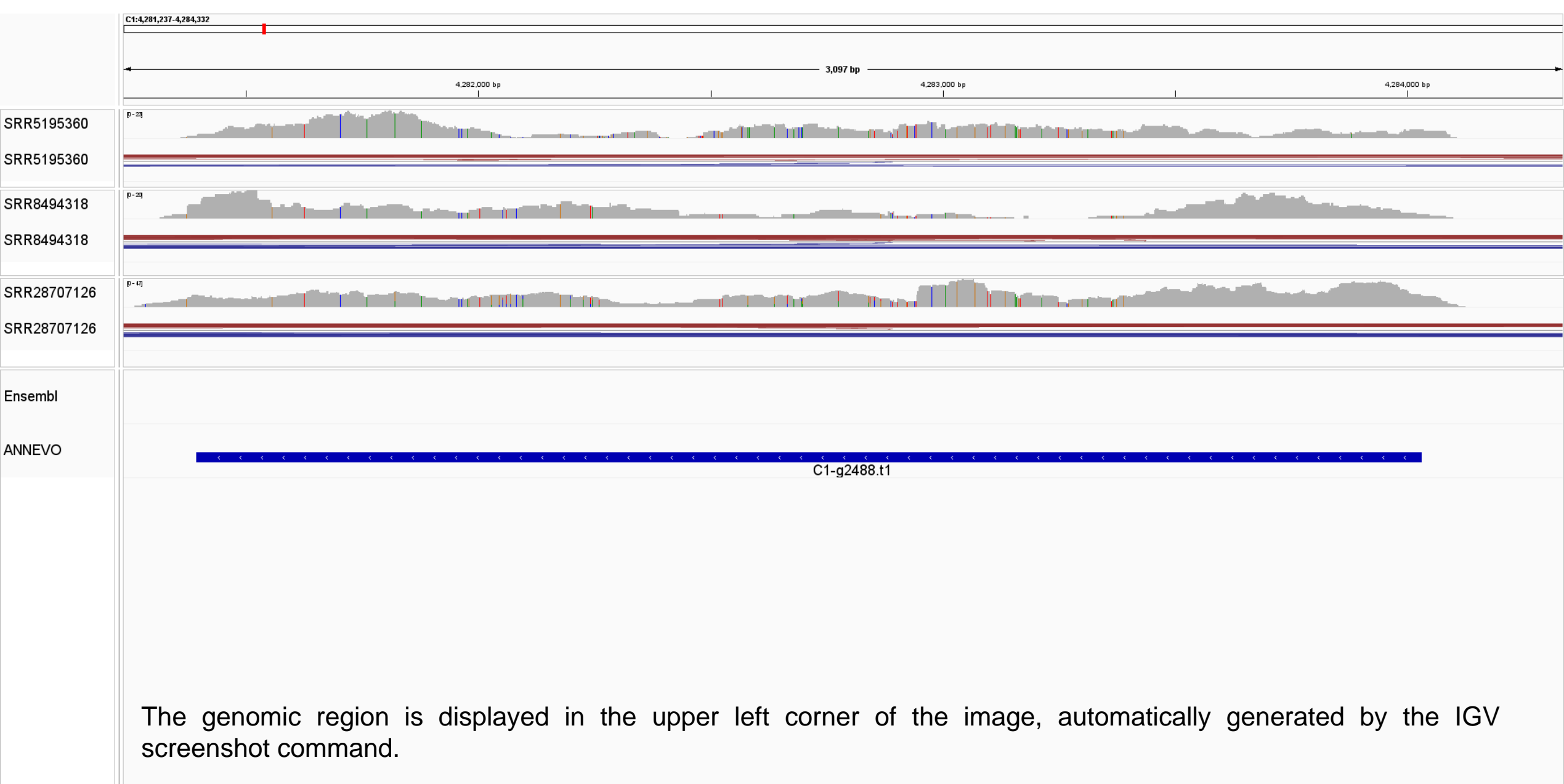
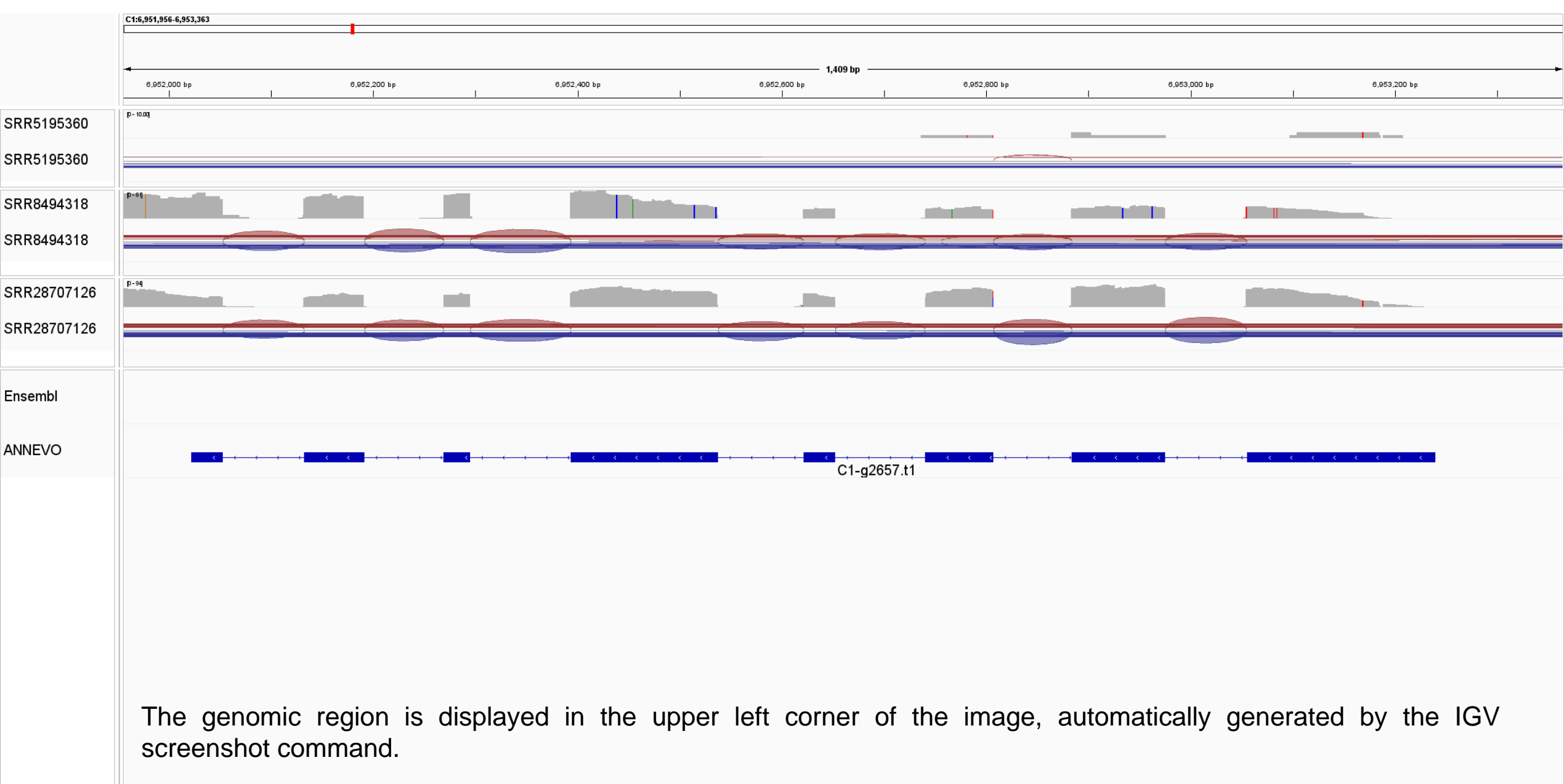


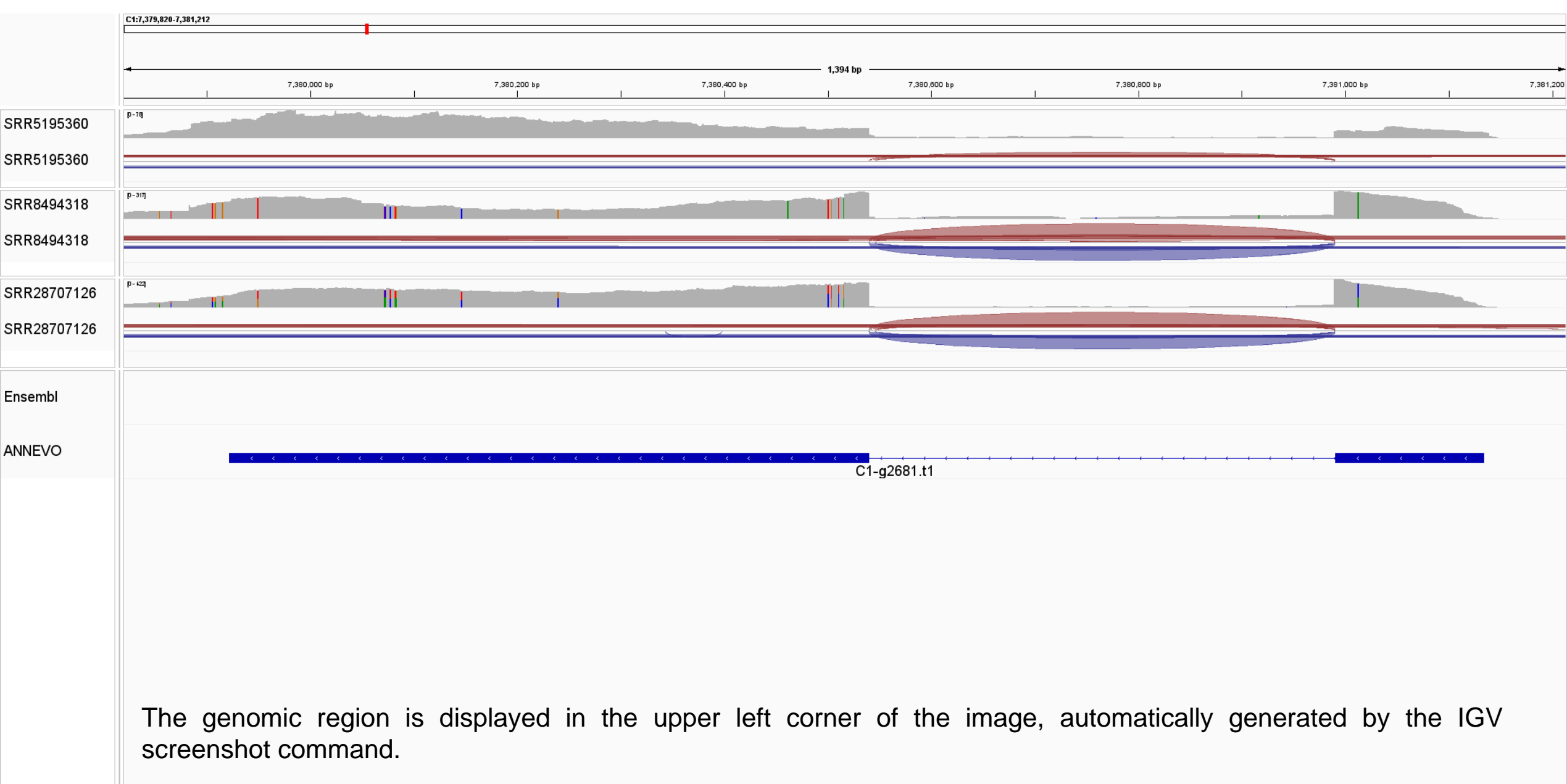
Additional BUSCO Genes Predicted by ANNEVO and Fully
Supported by RNA-Seq

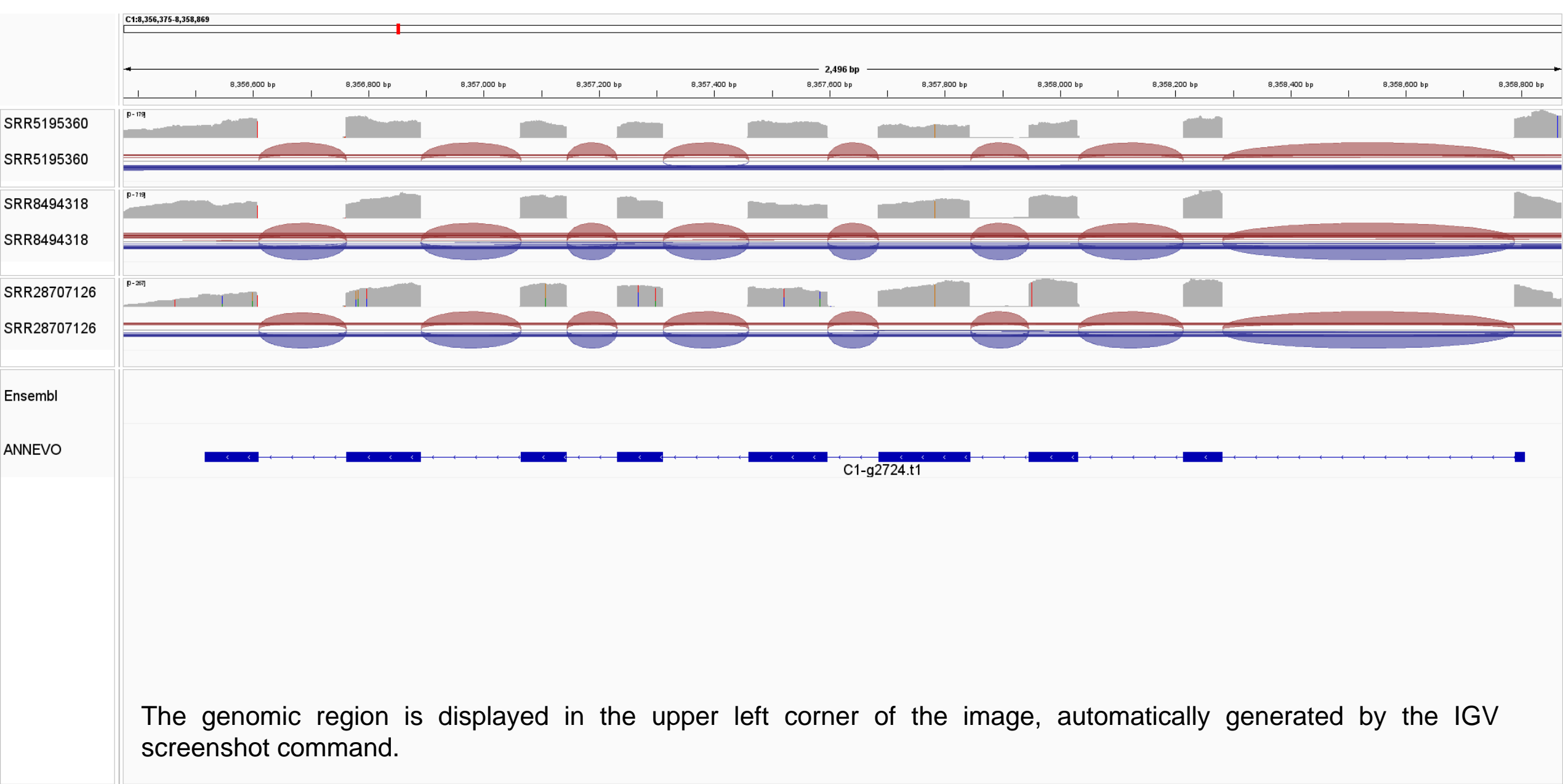
Missing BUSCO Gene in Ensembl







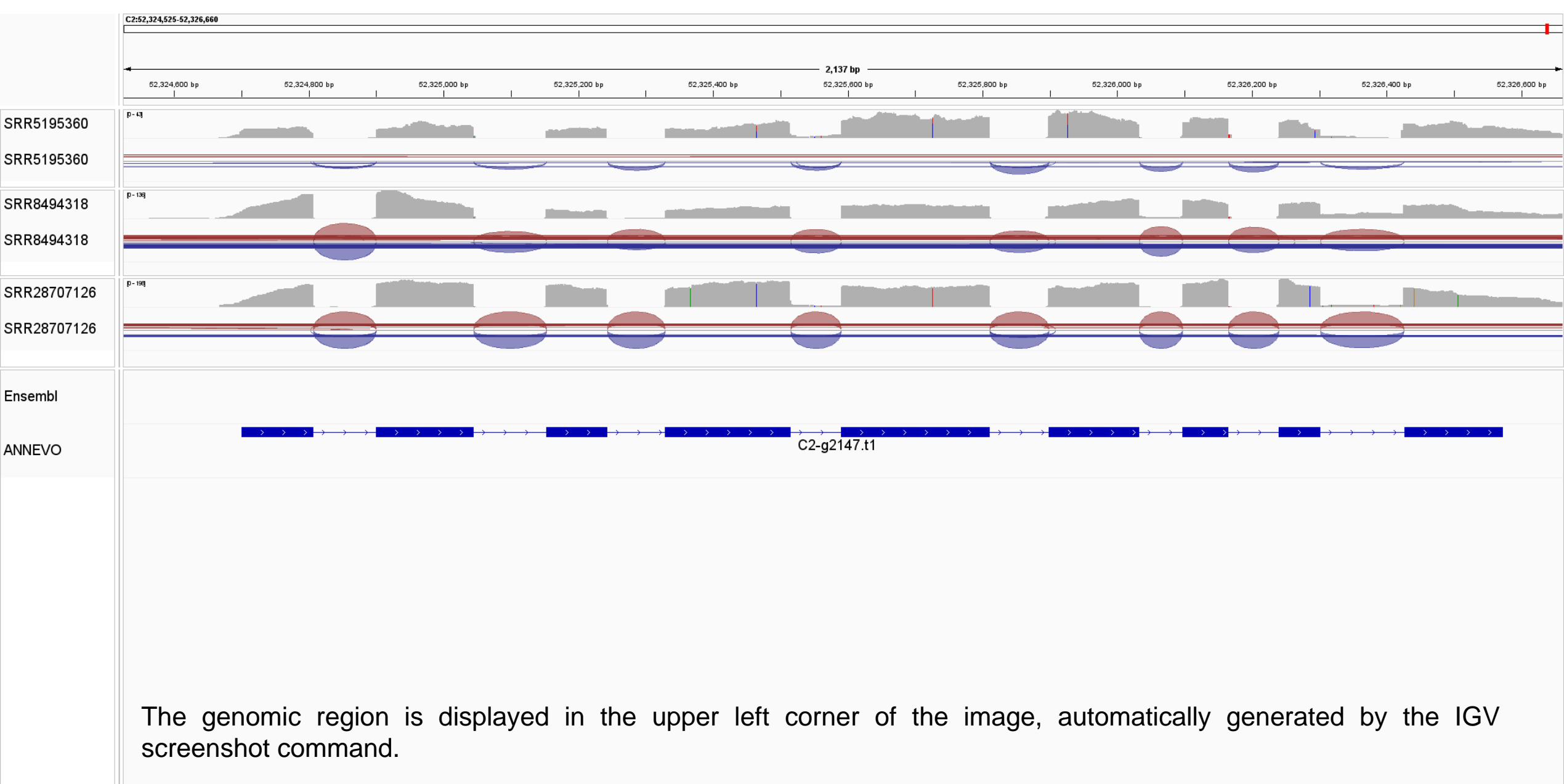




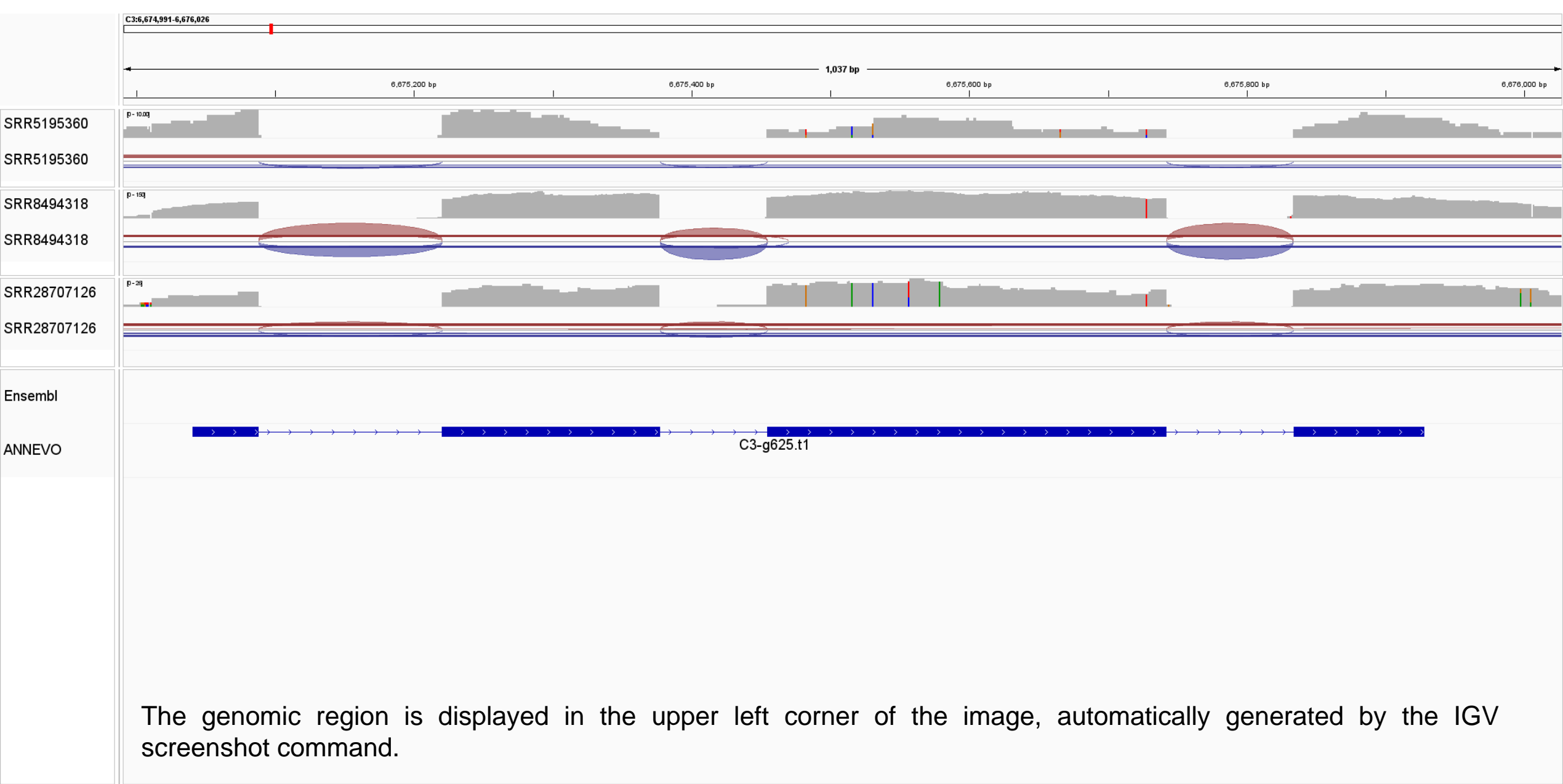


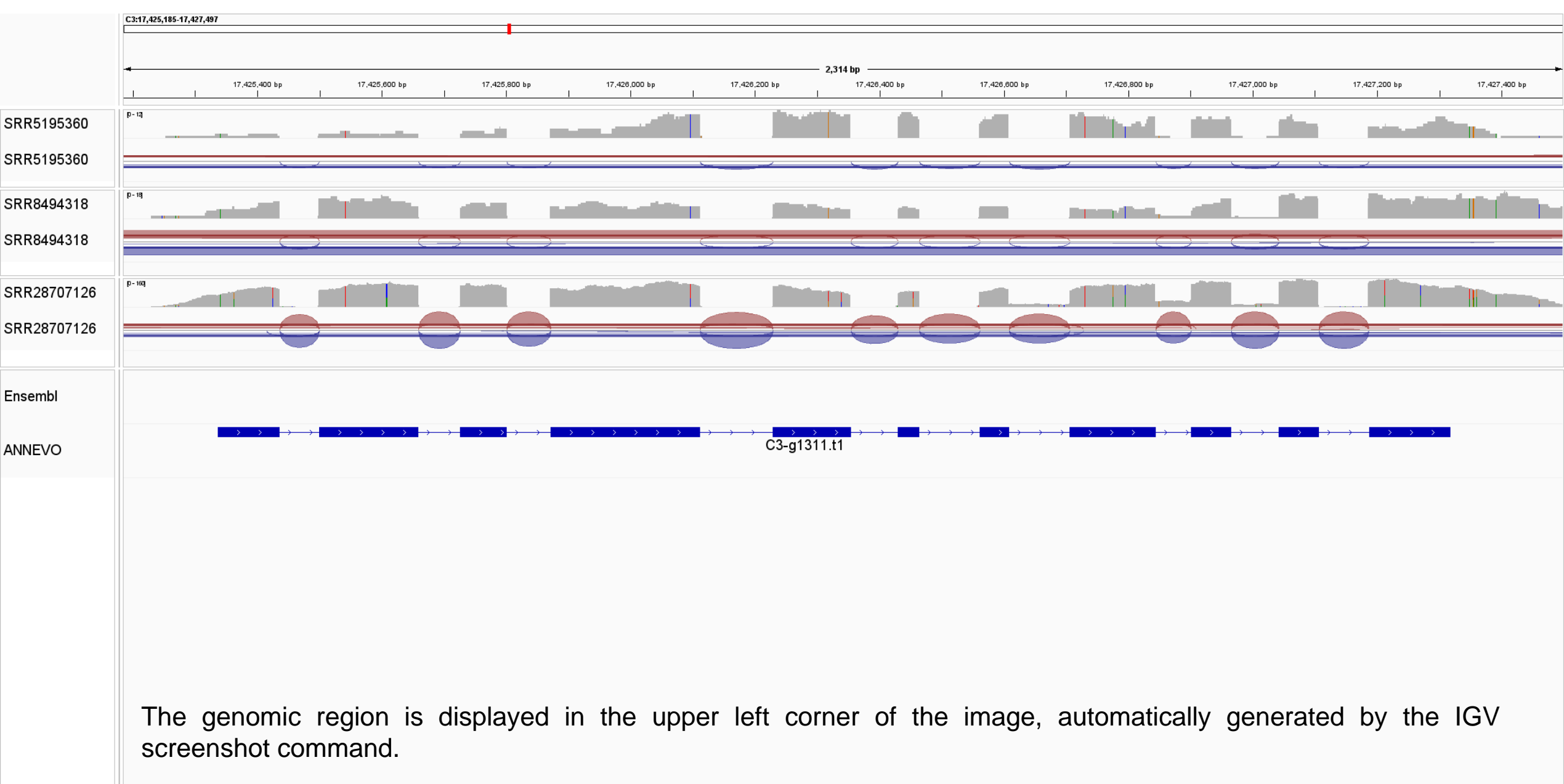


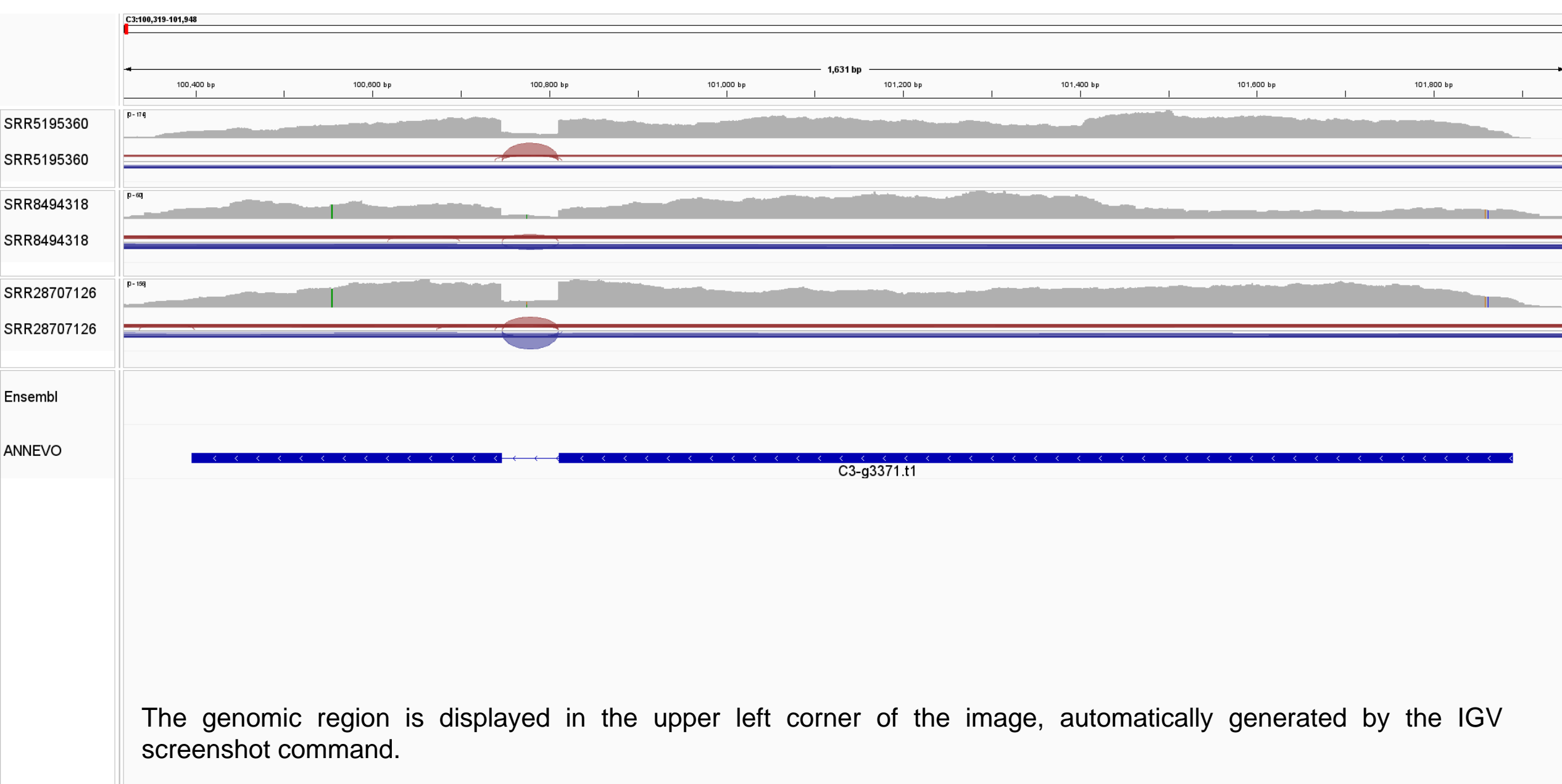


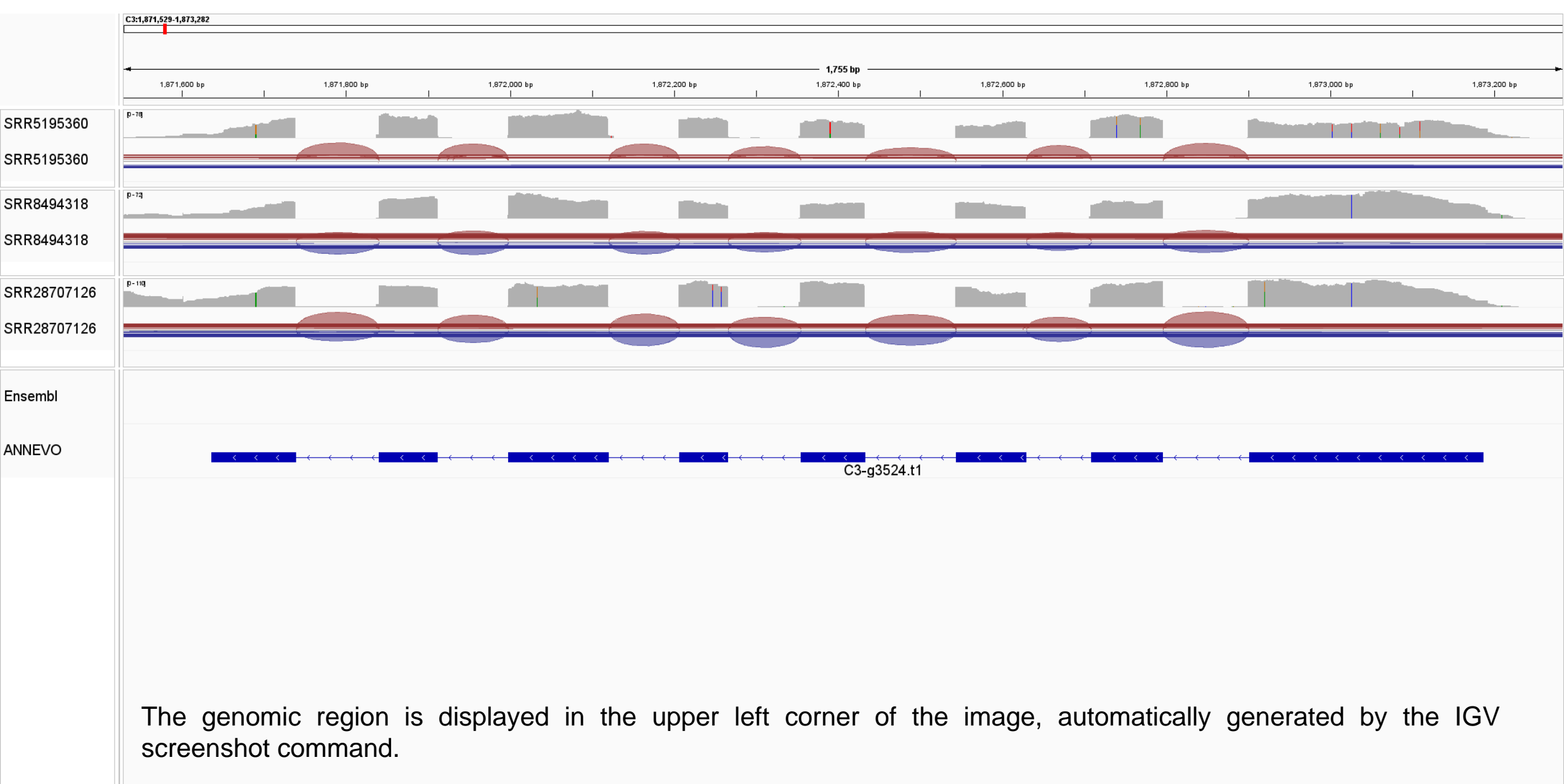


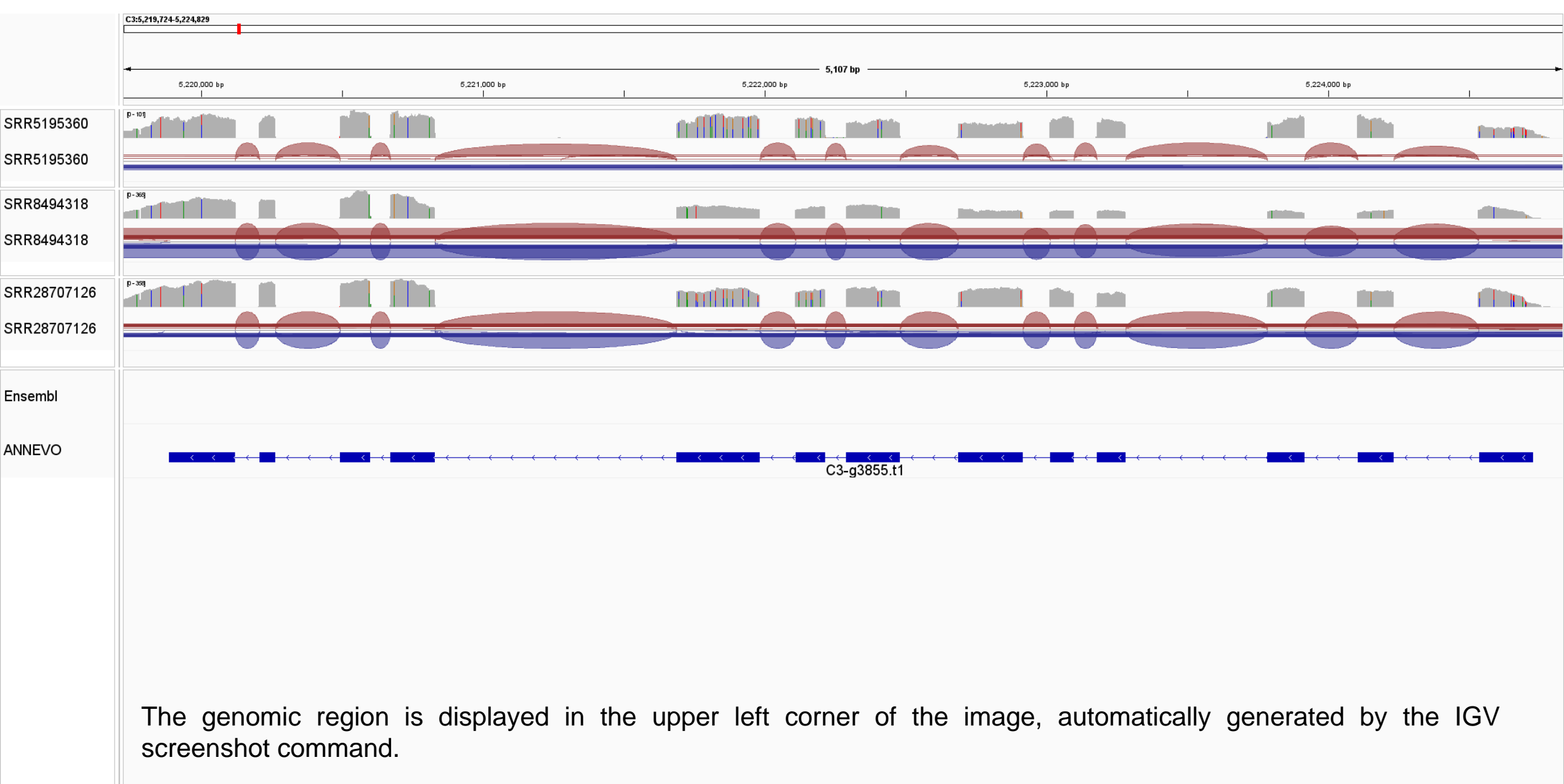


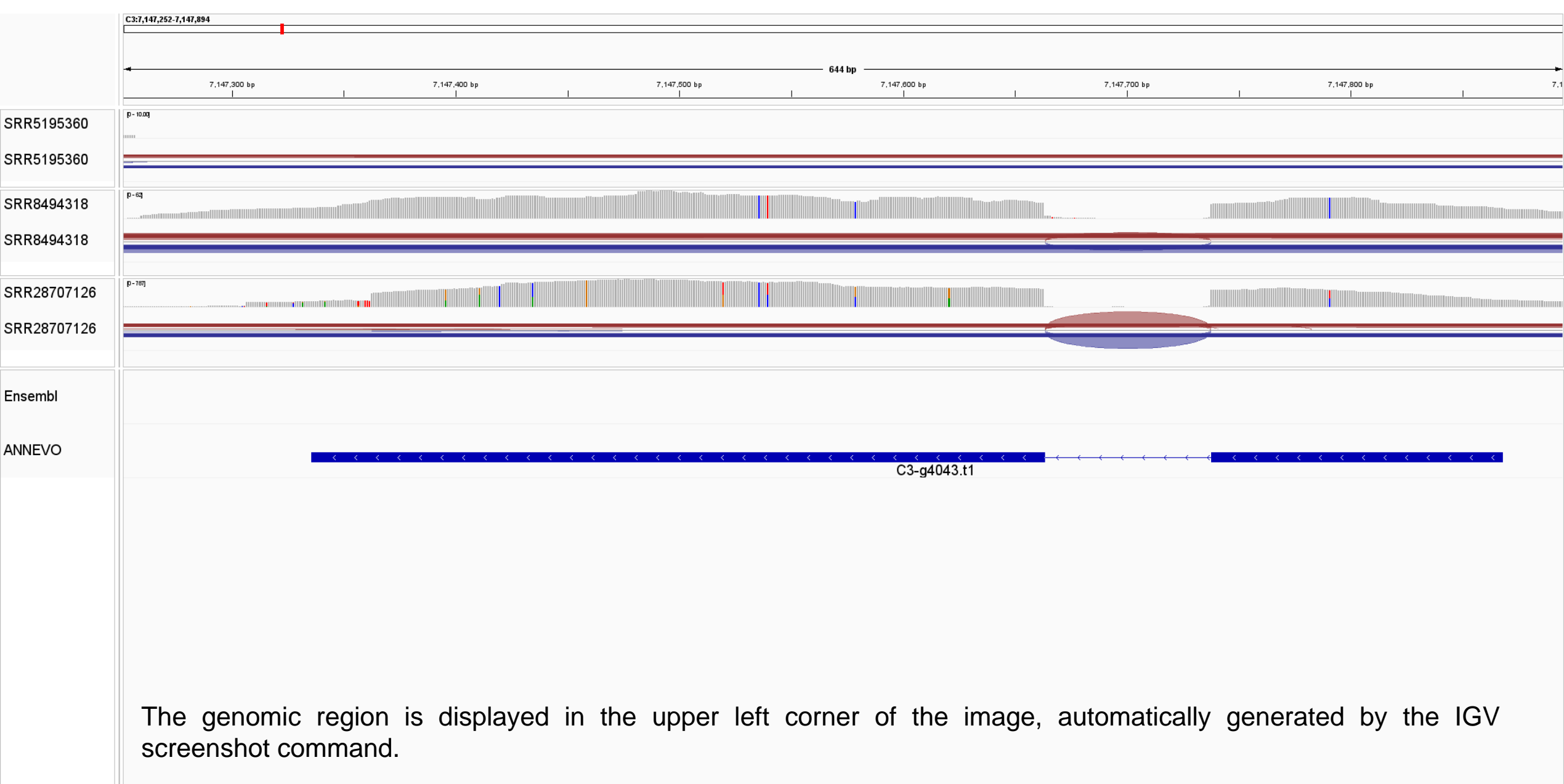


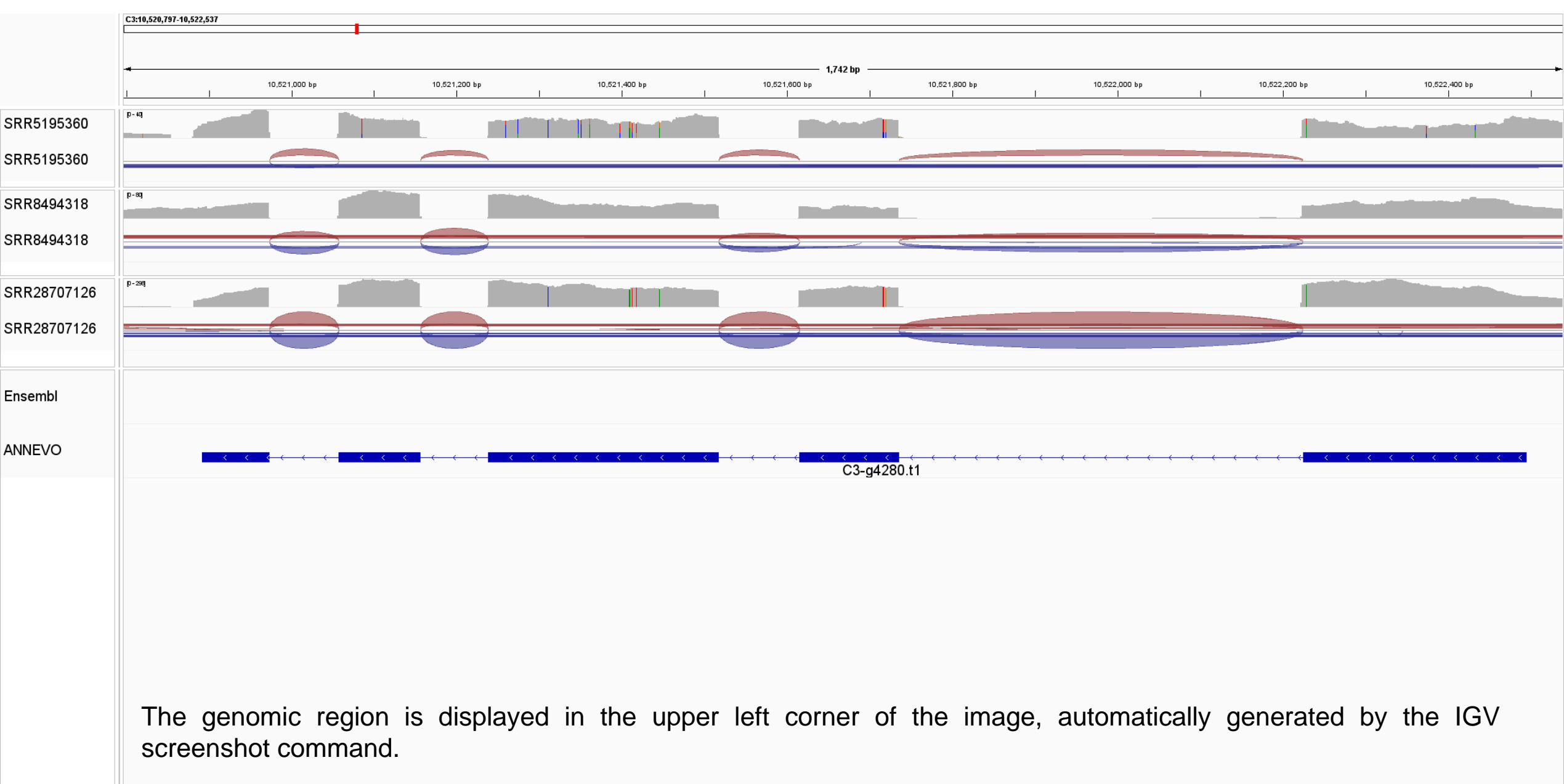


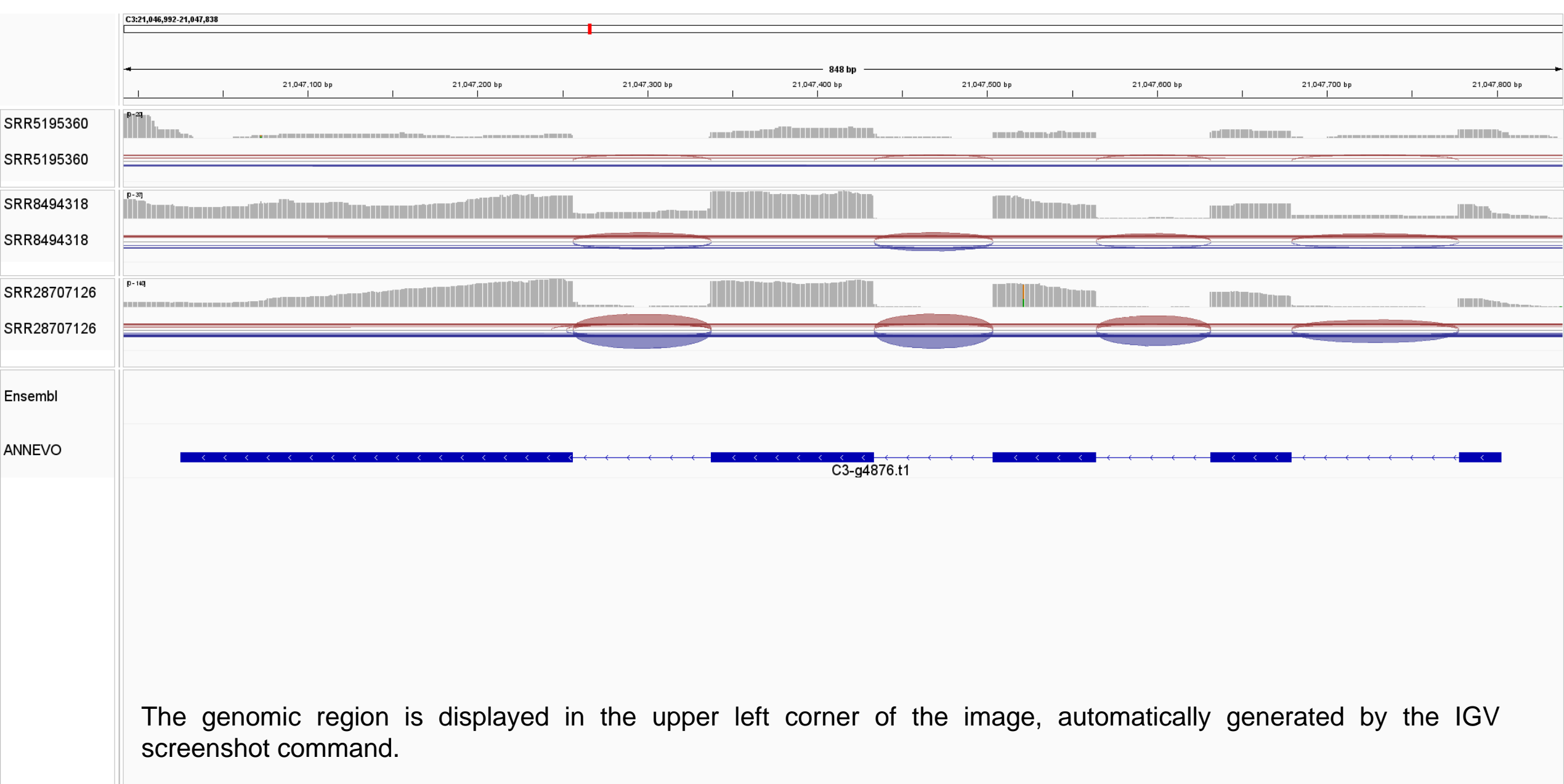


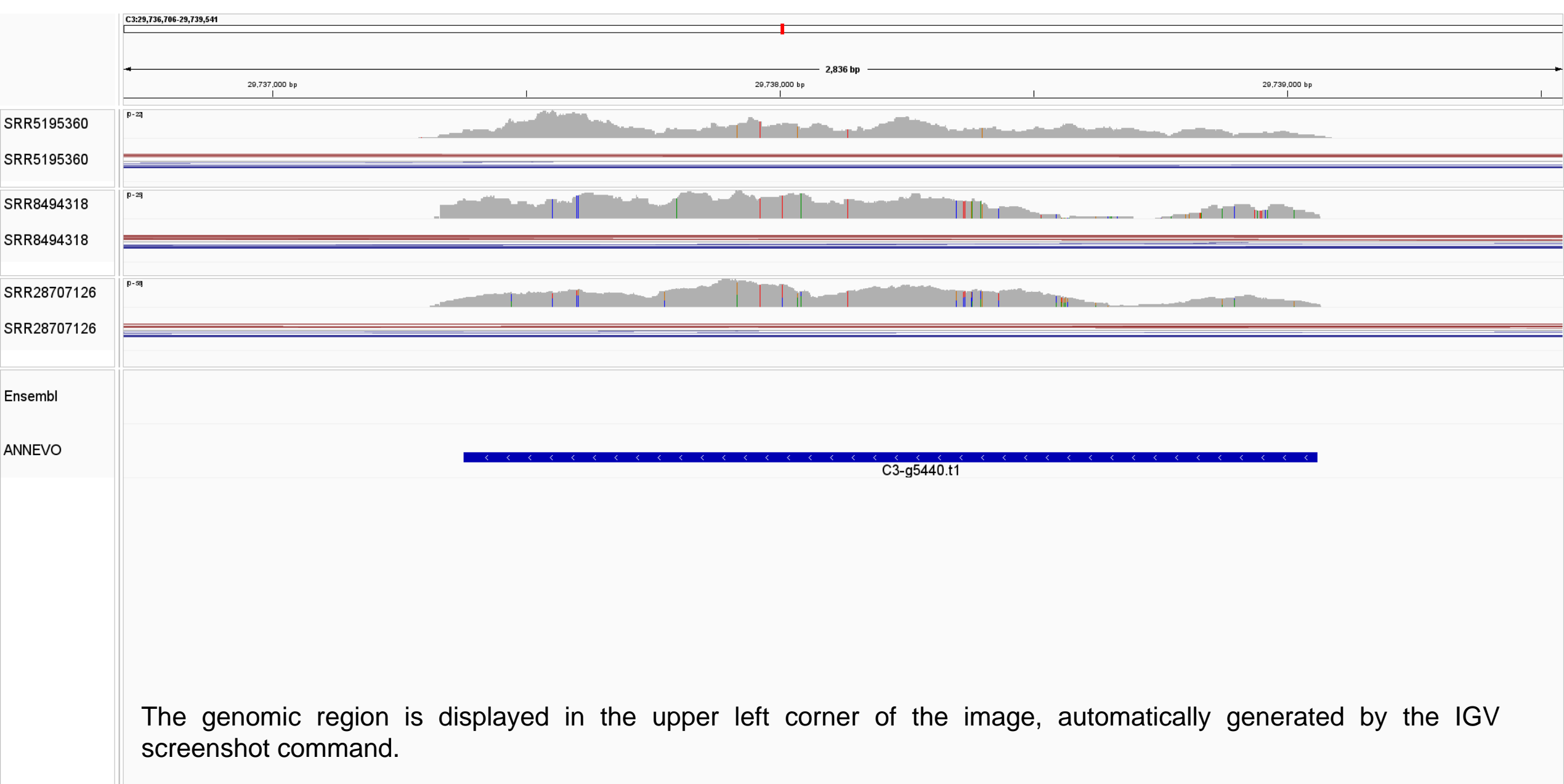


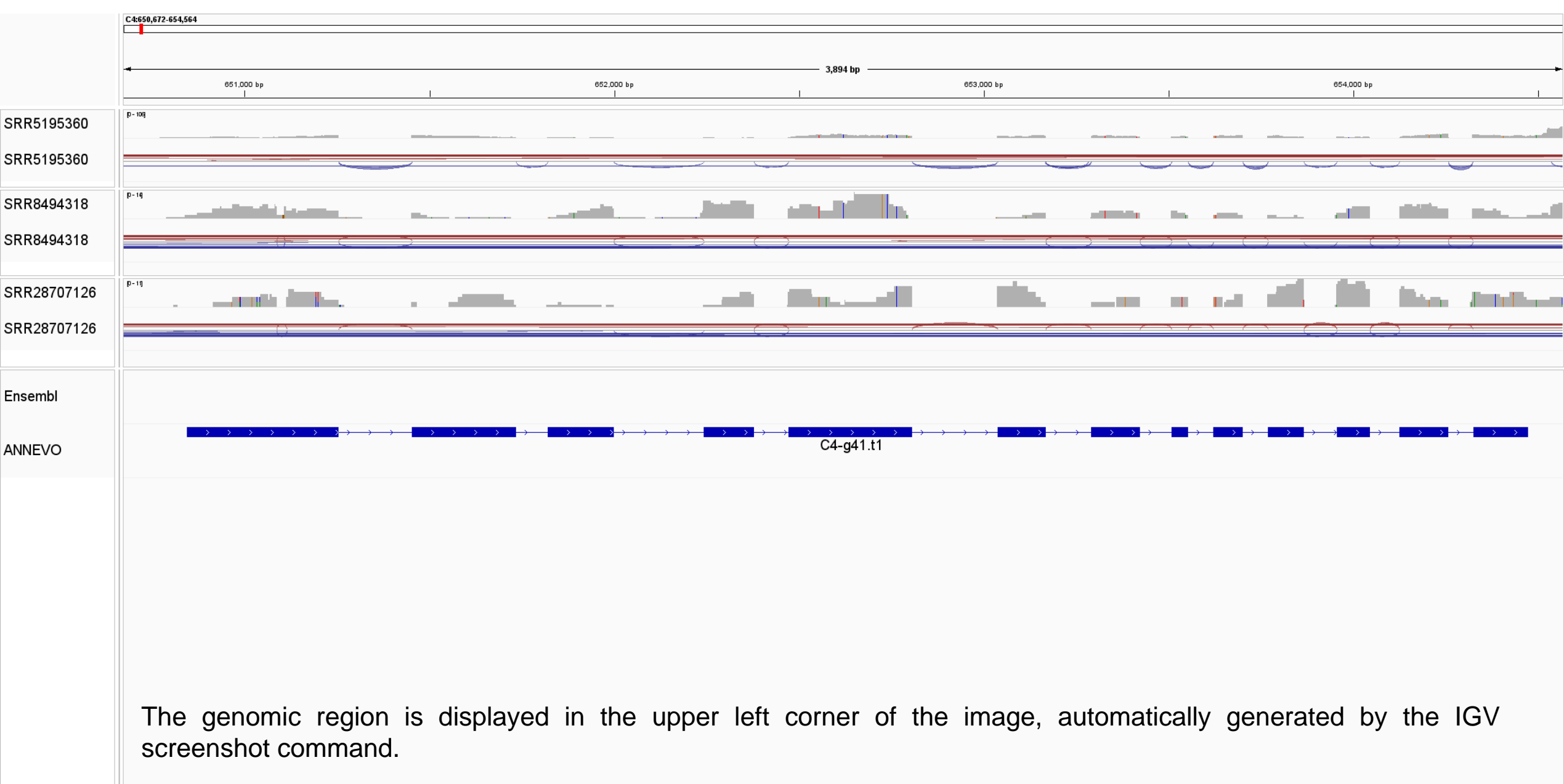




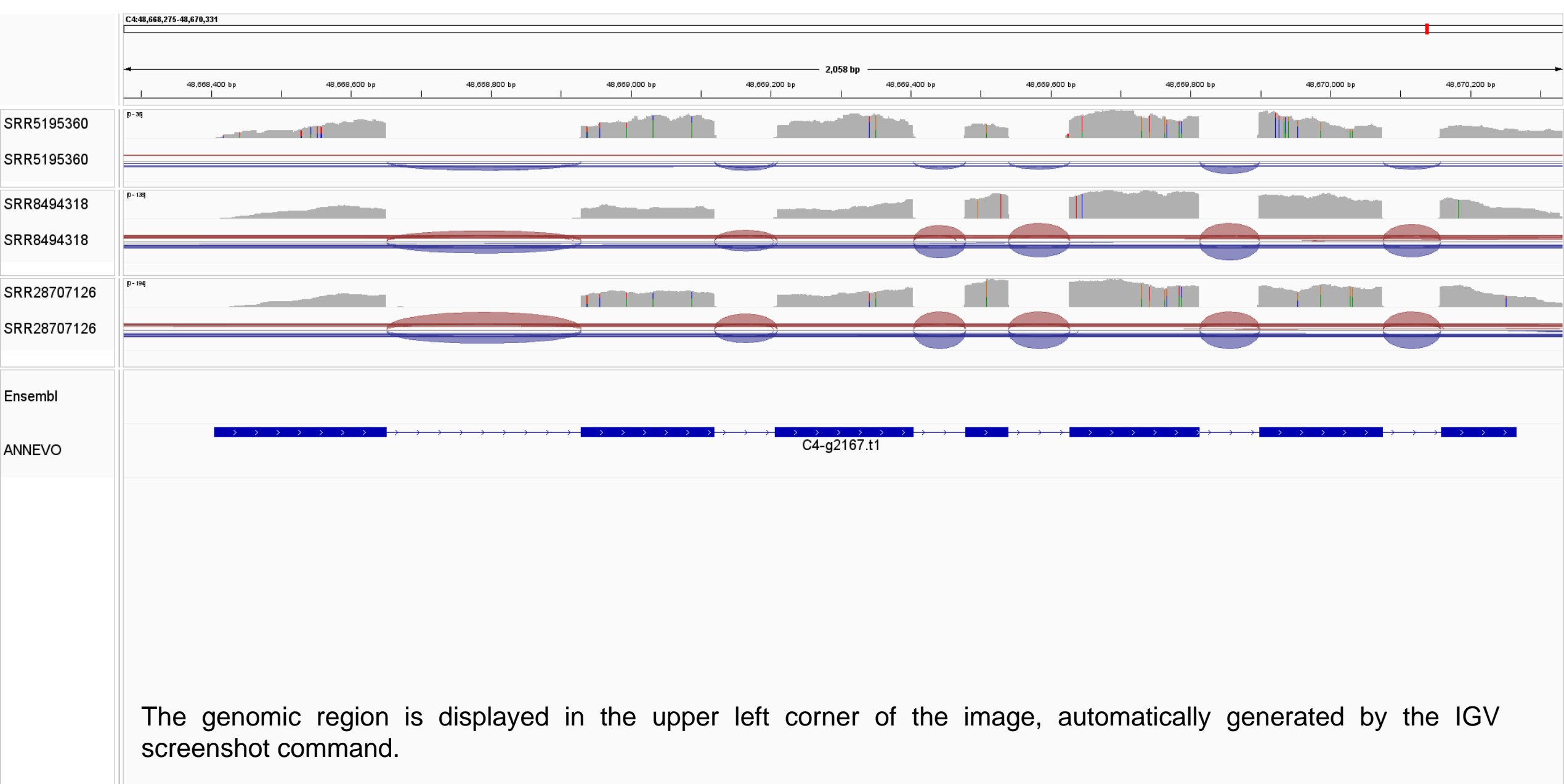


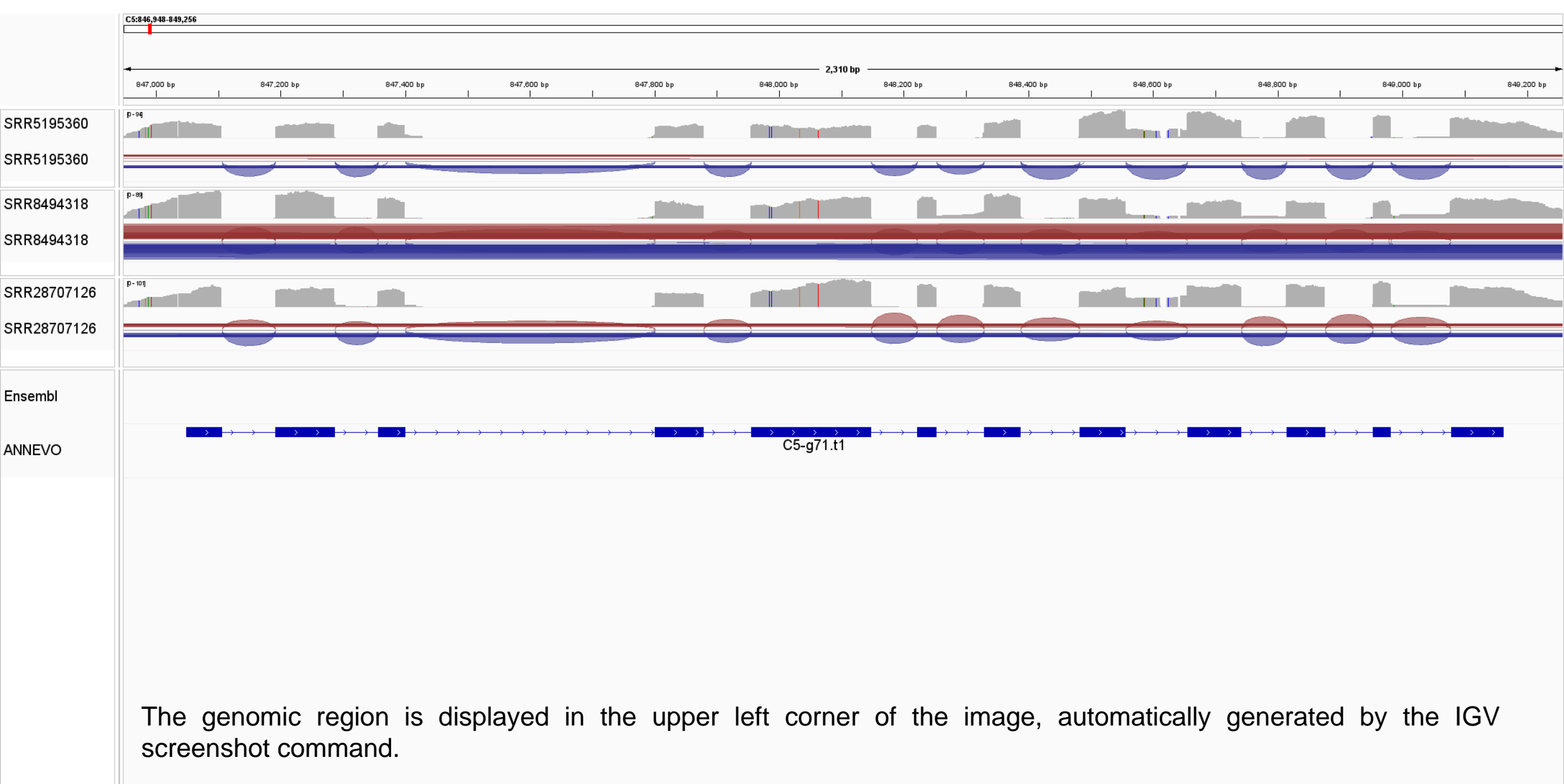






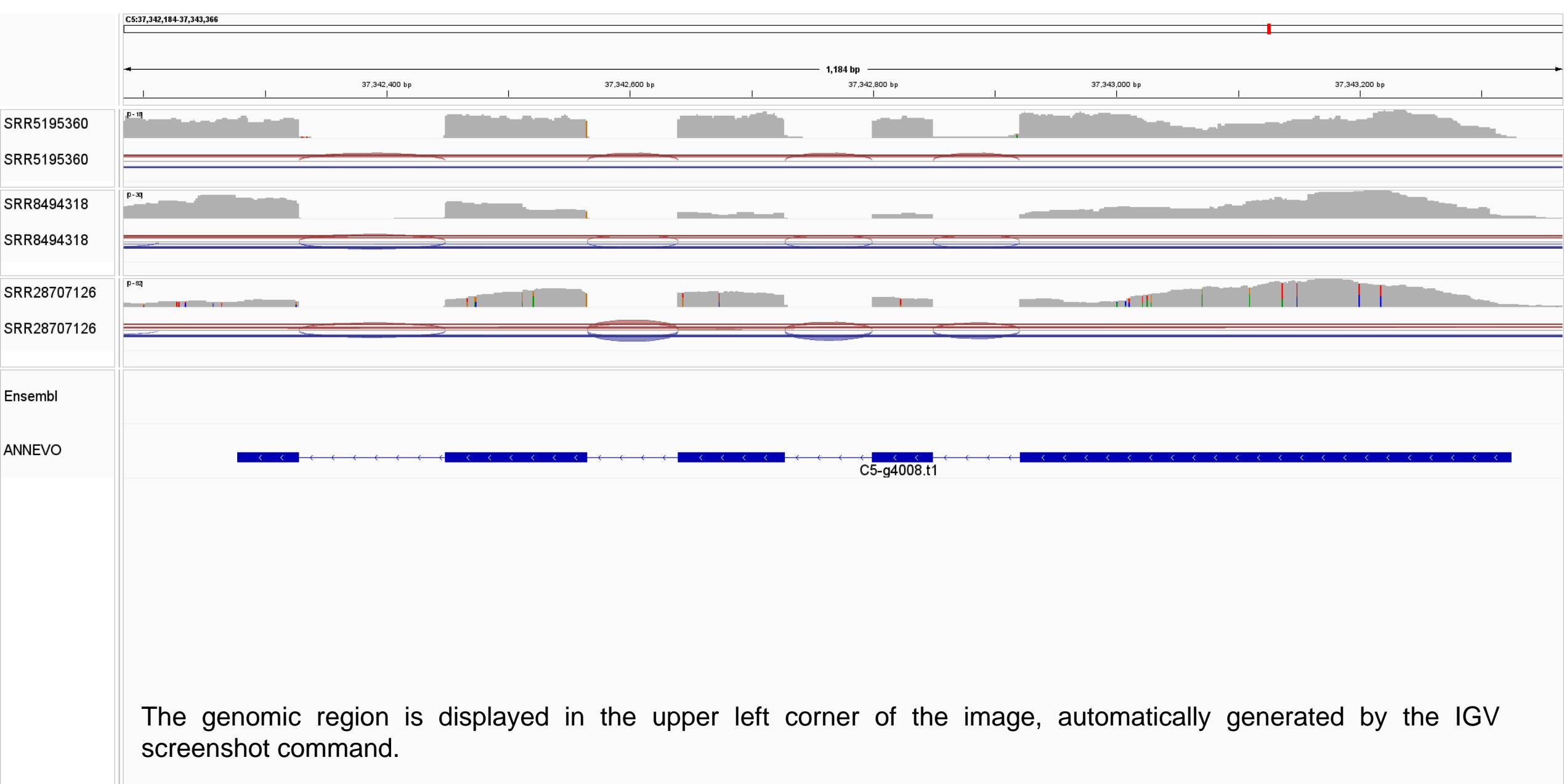


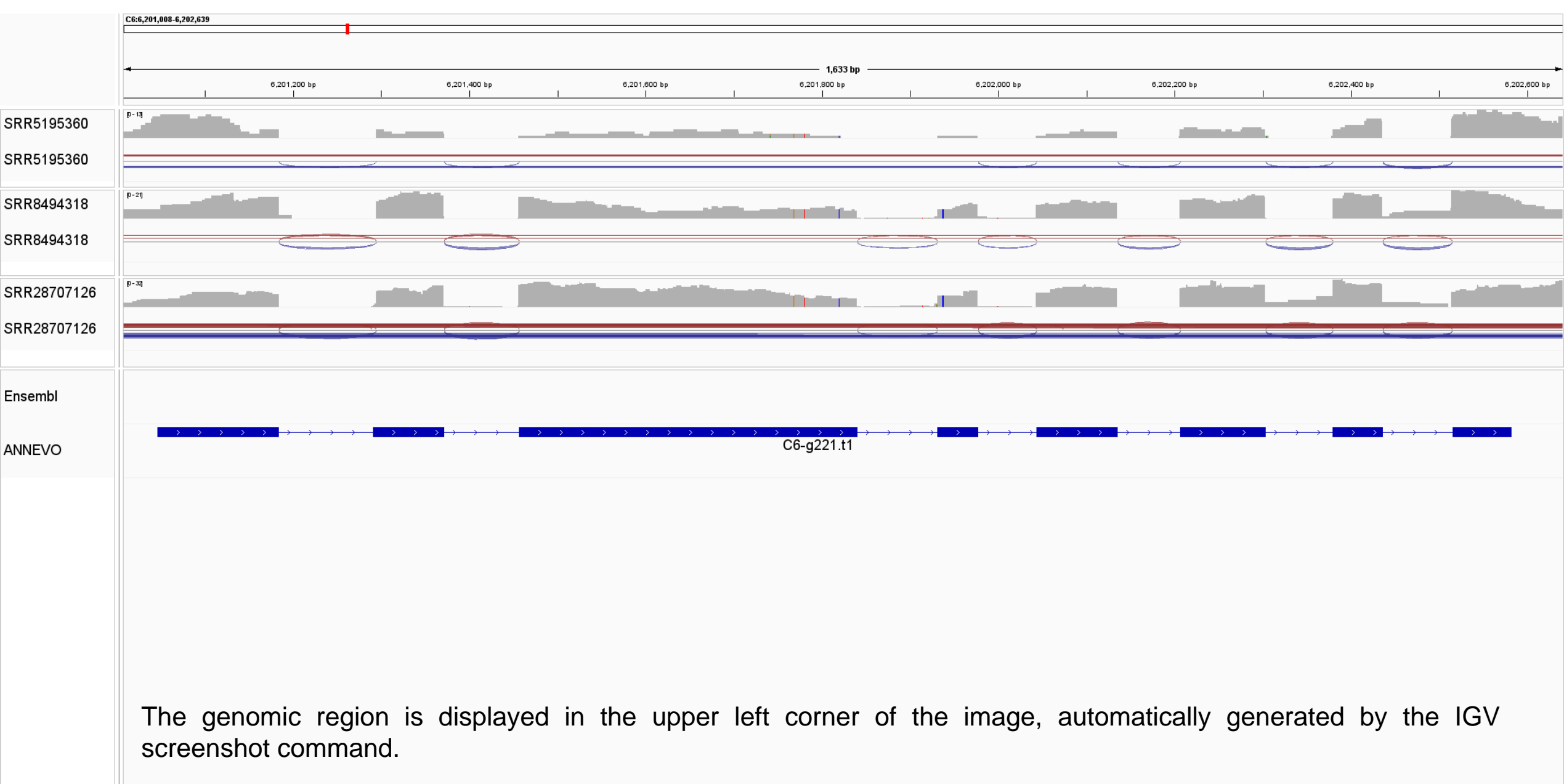


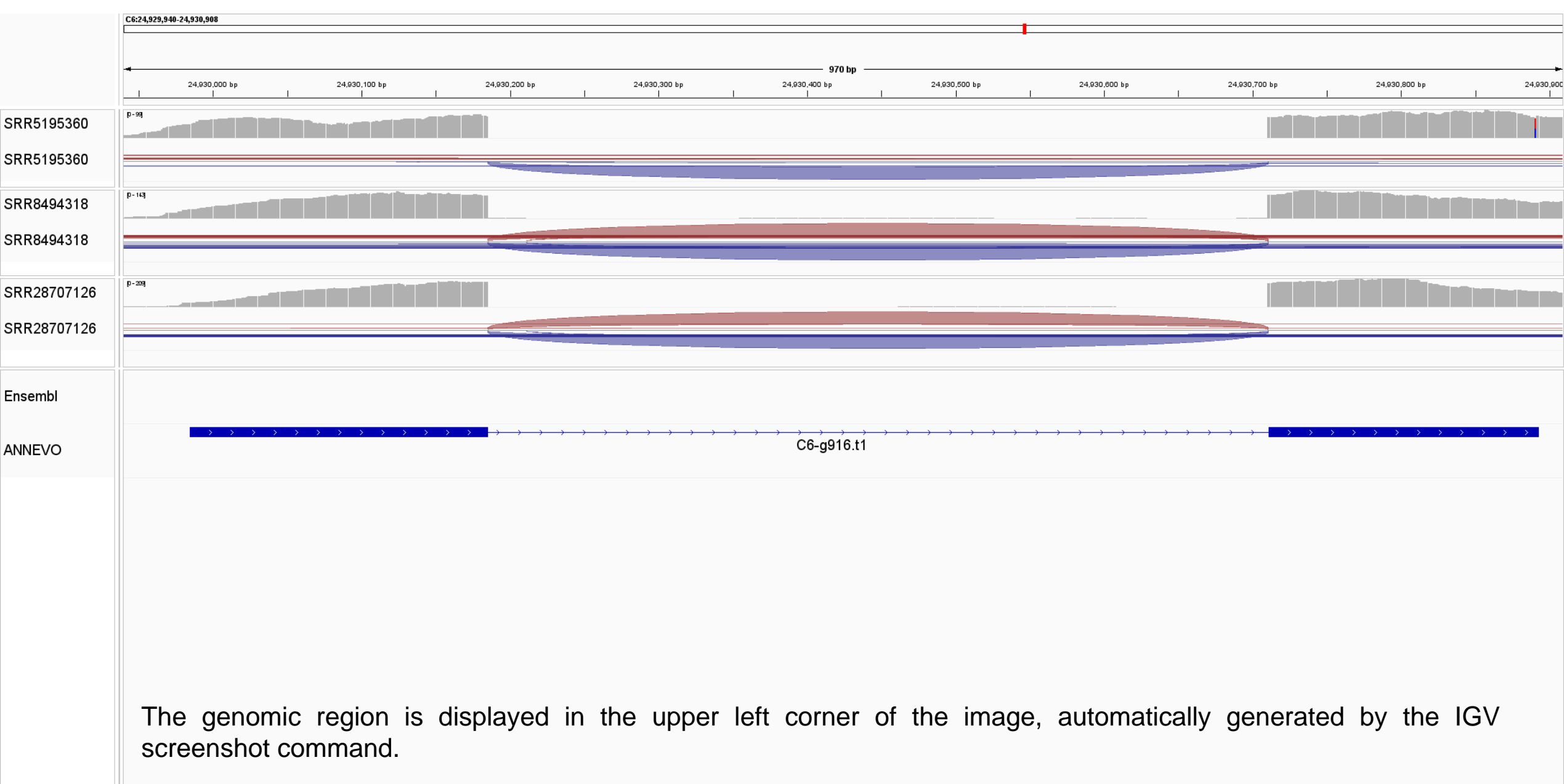






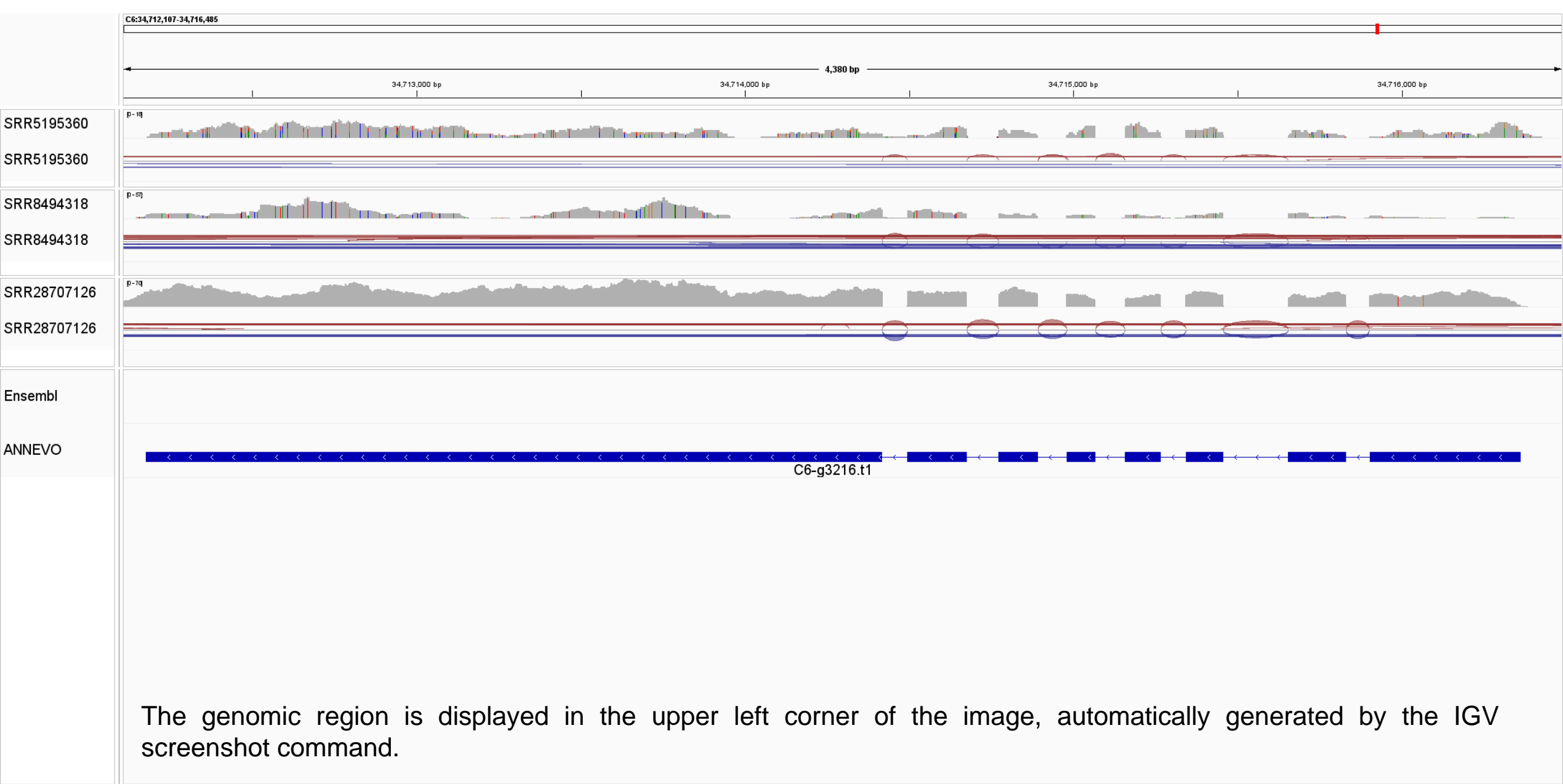


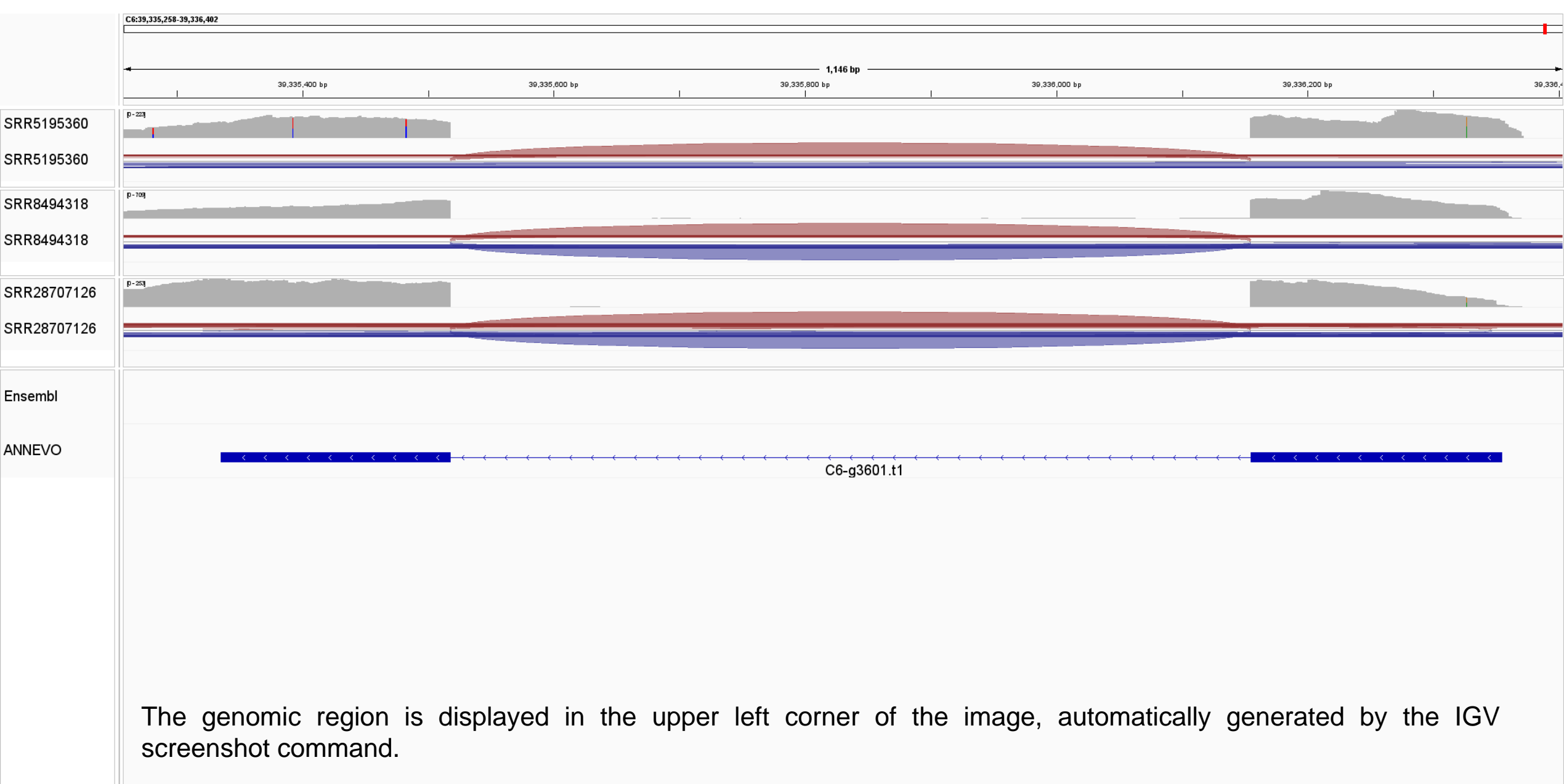


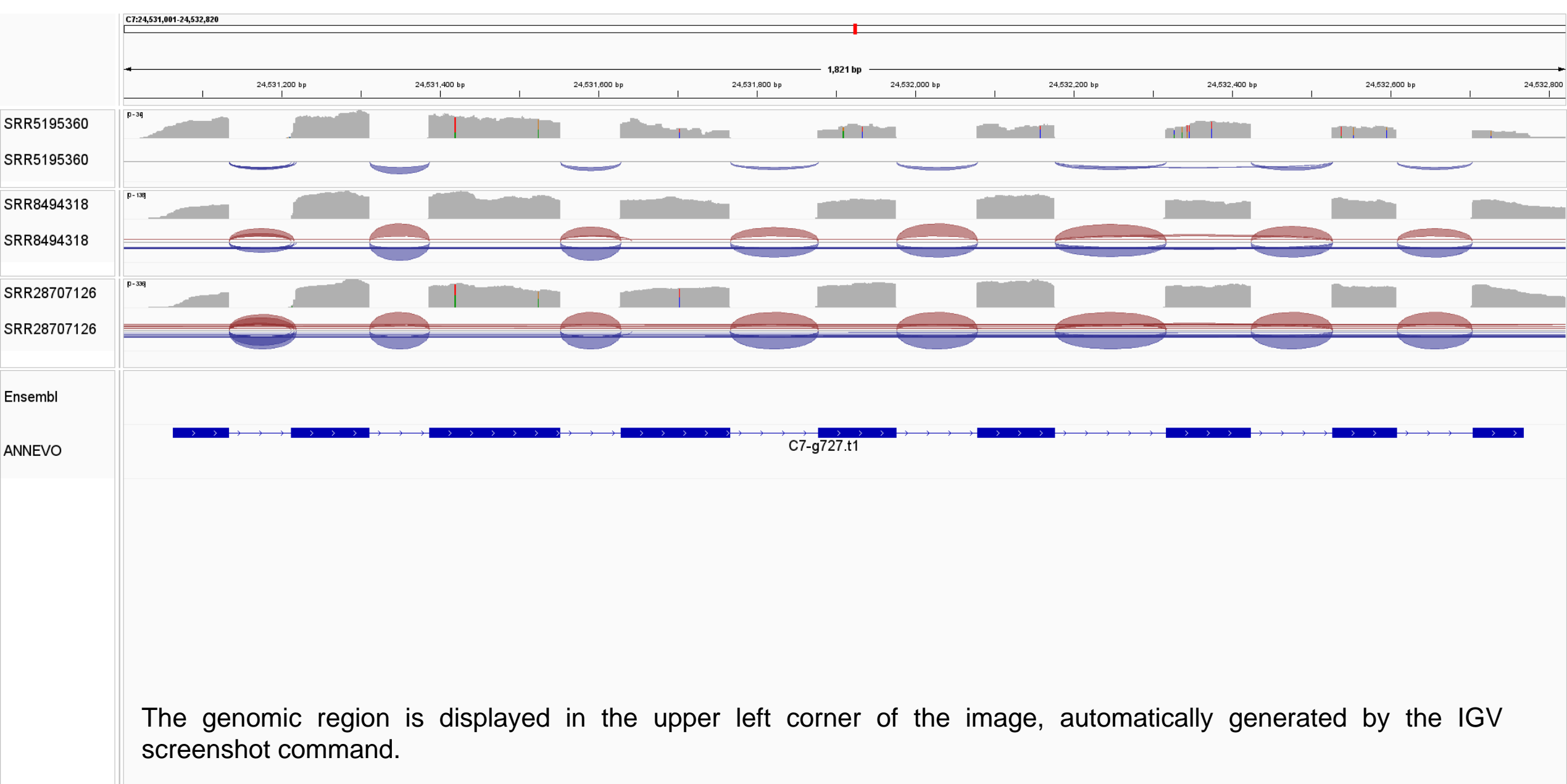


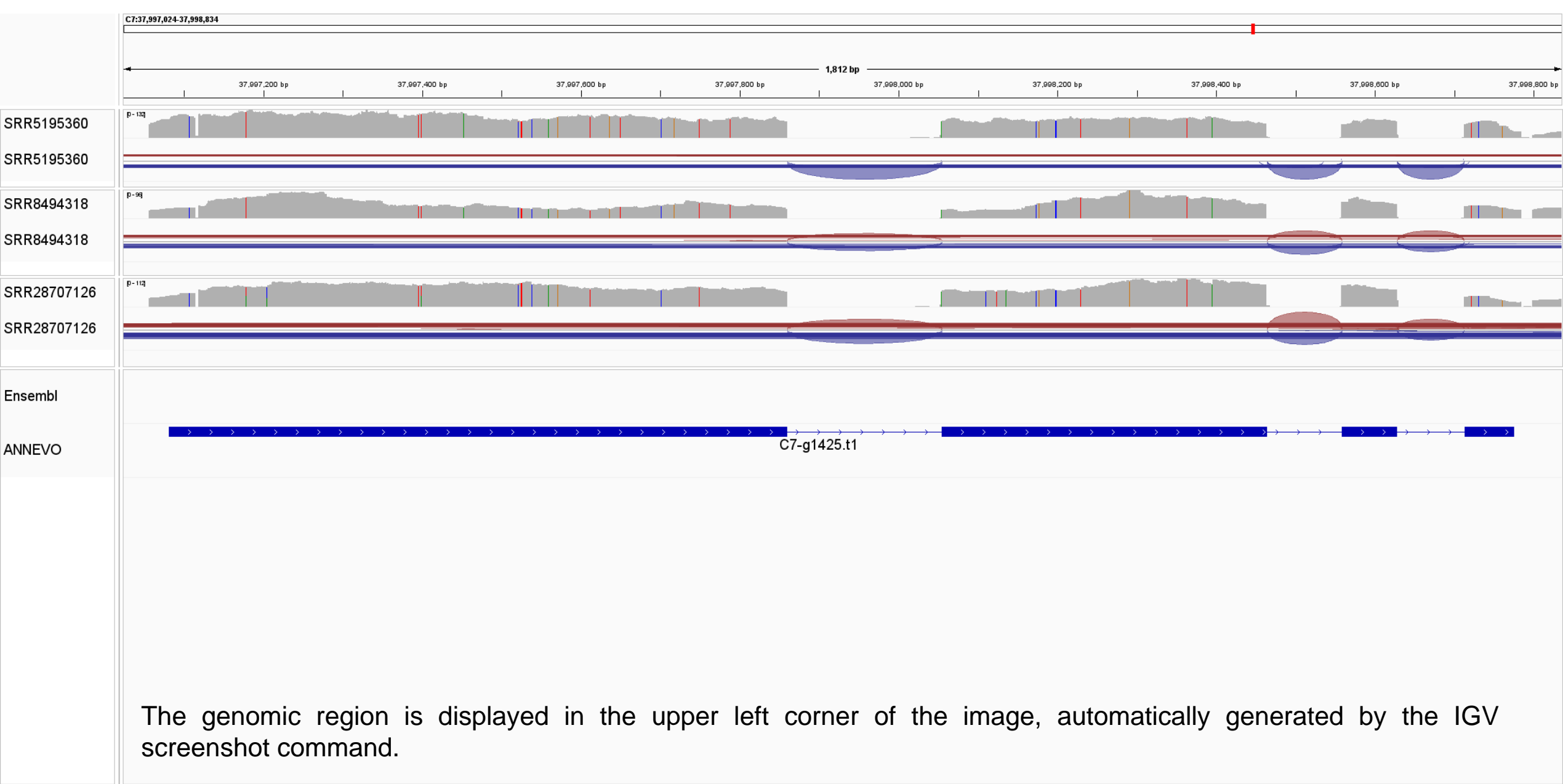


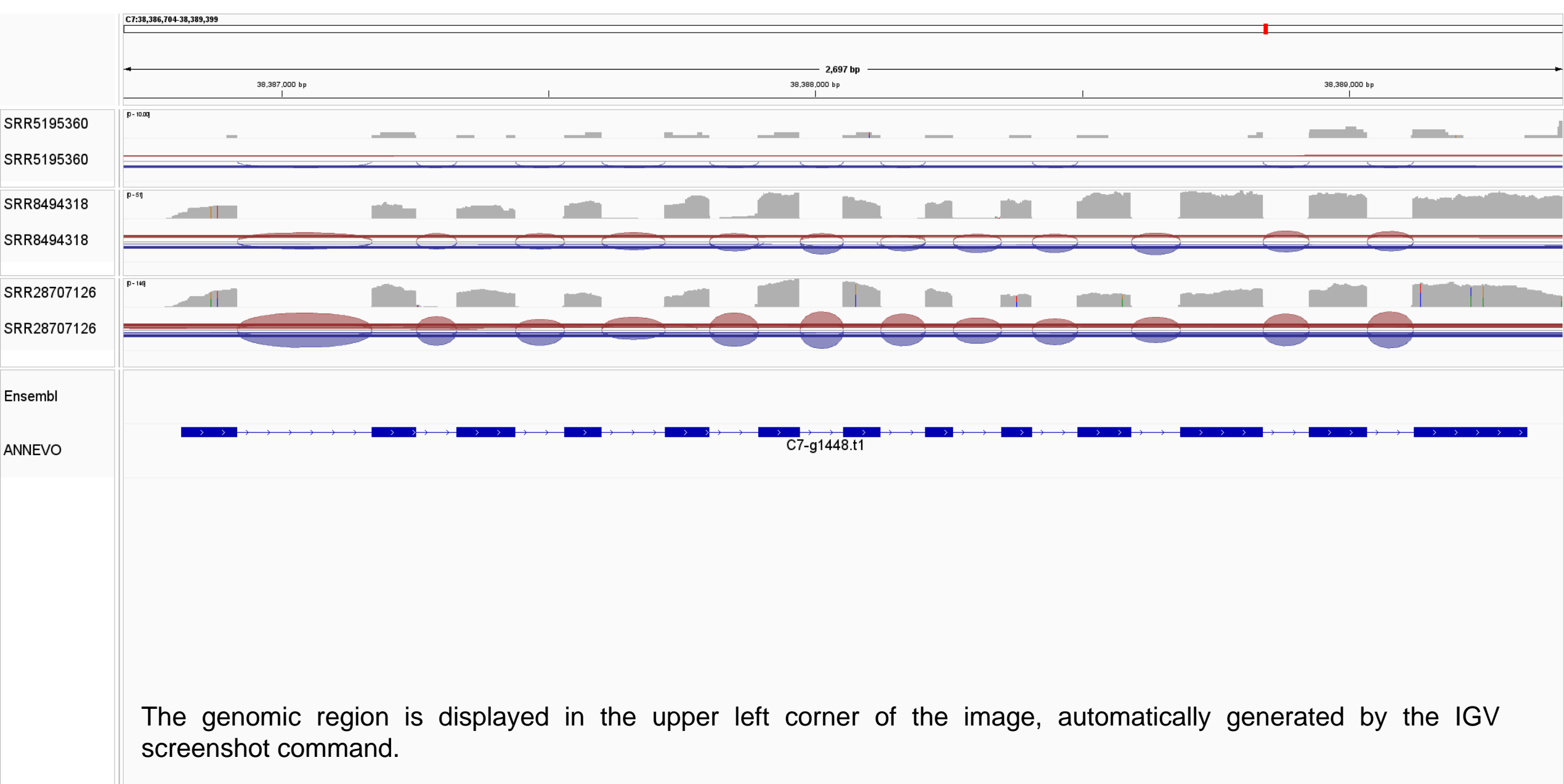


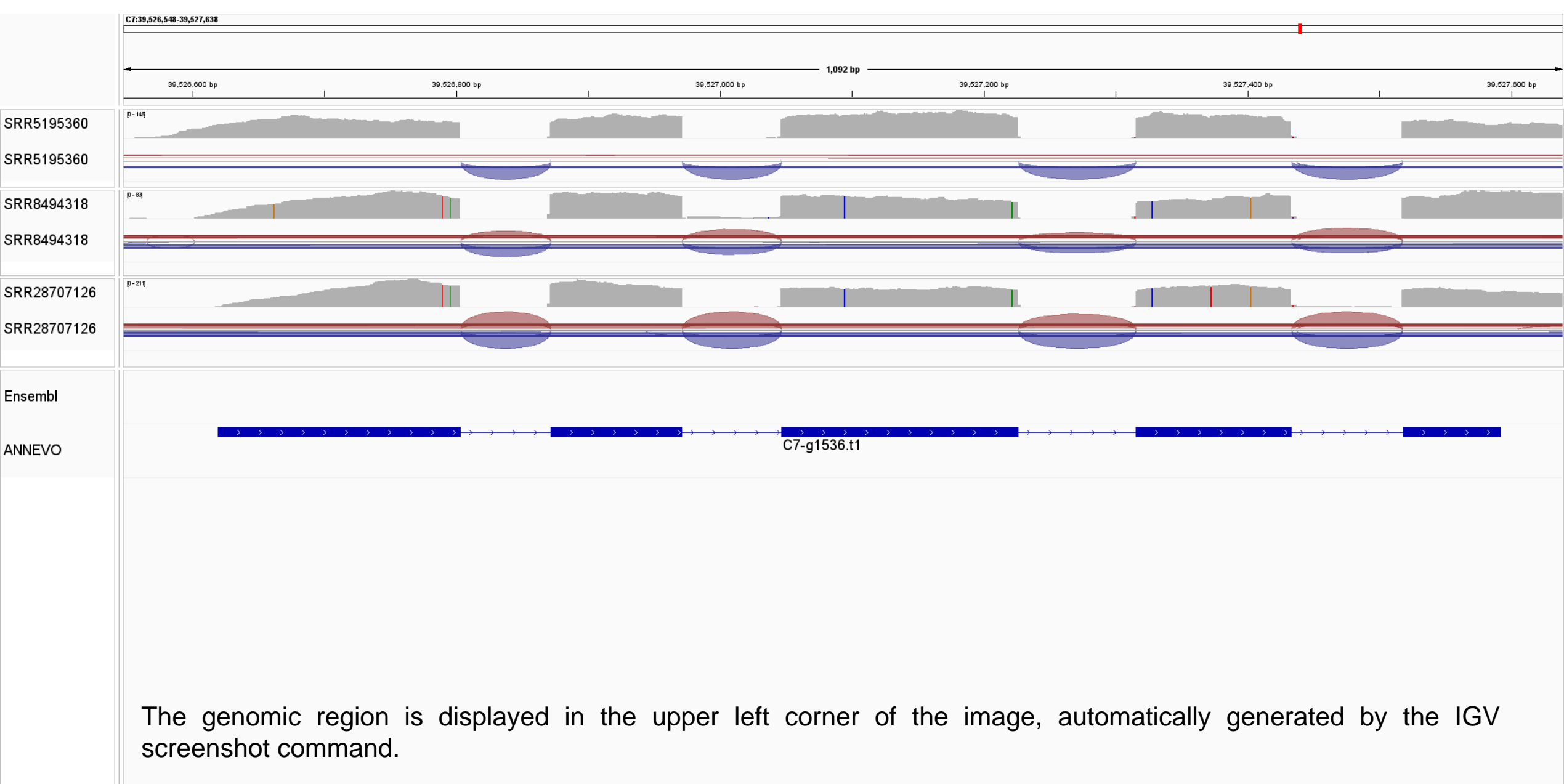


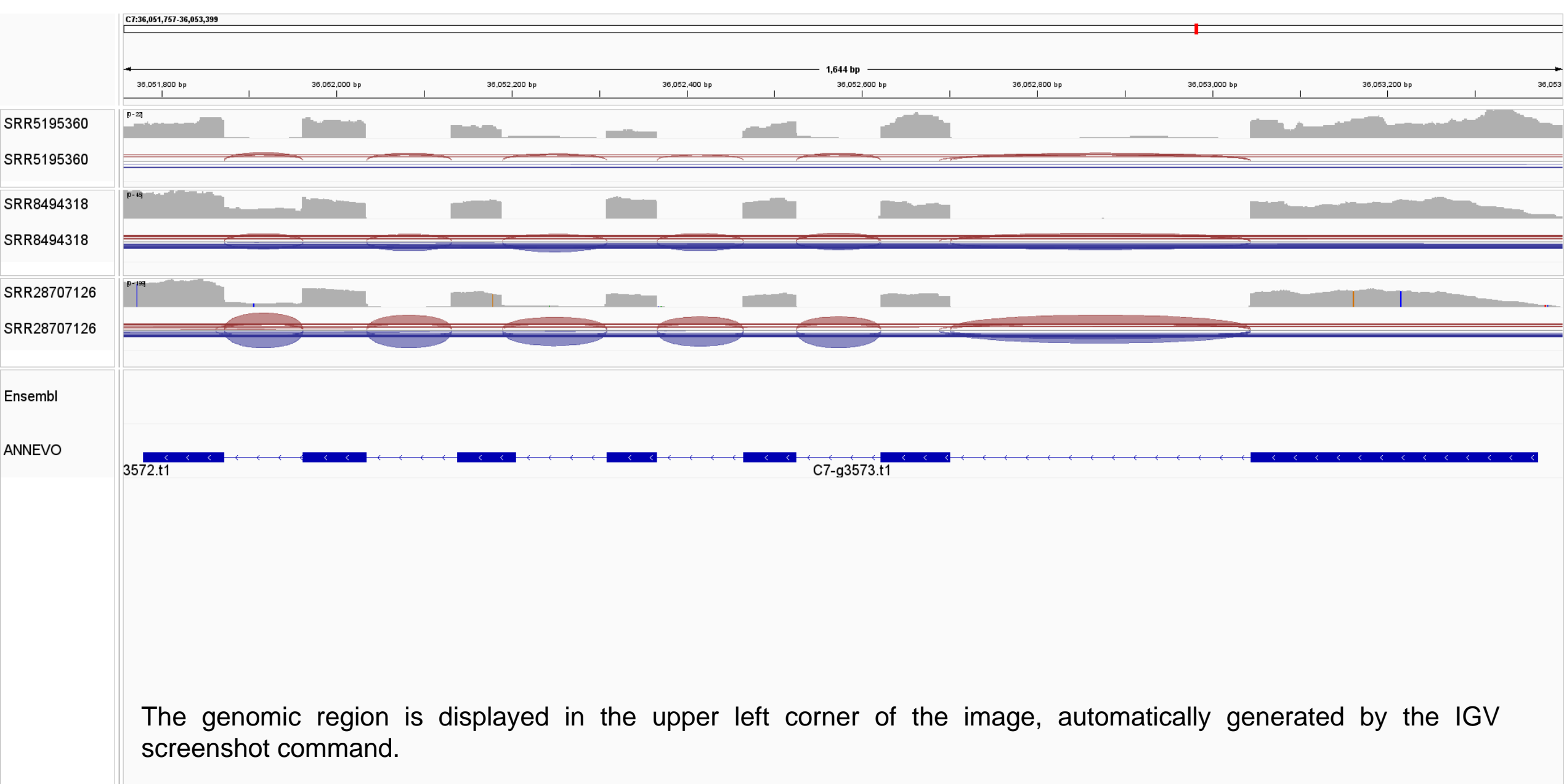




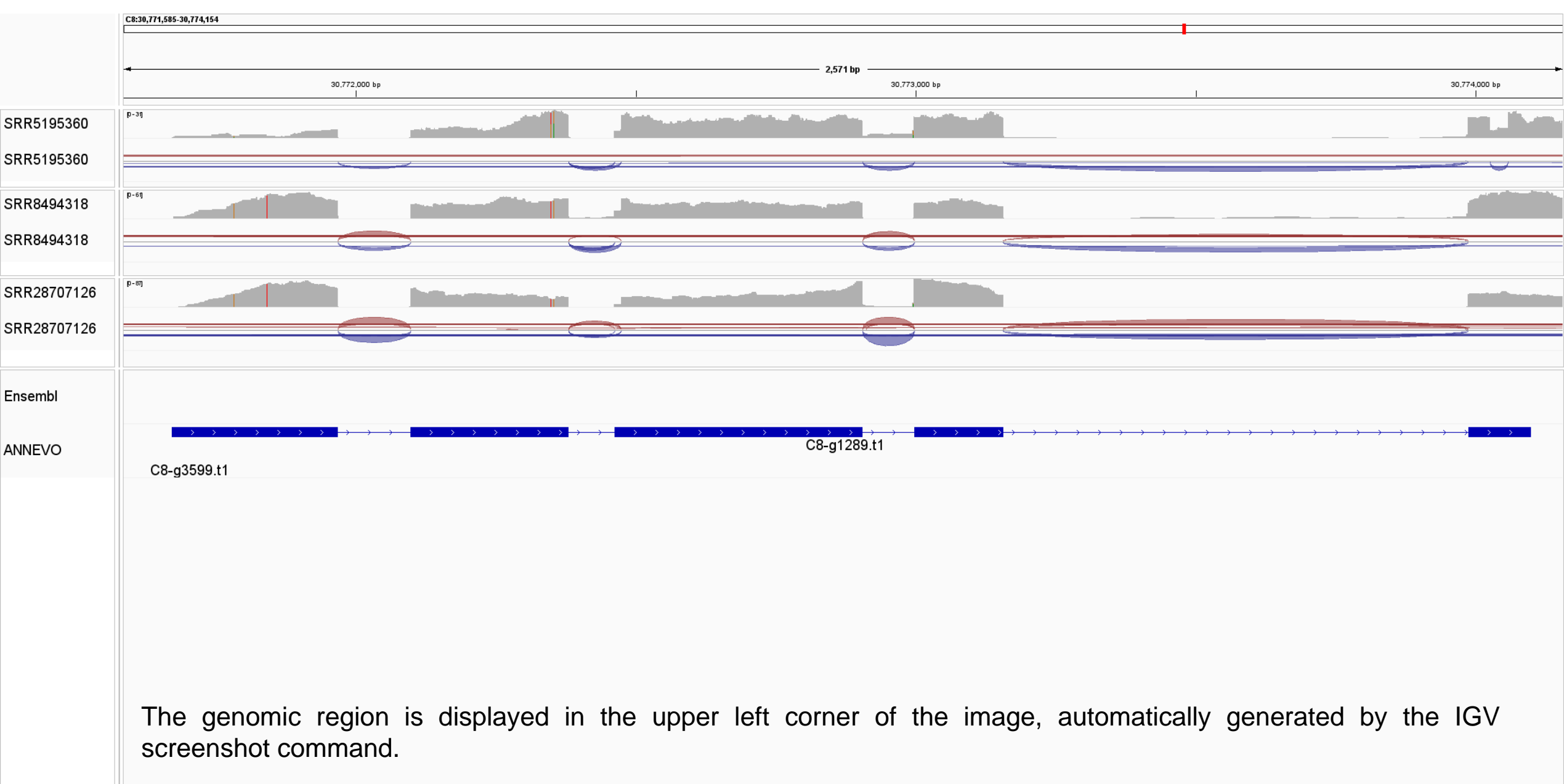




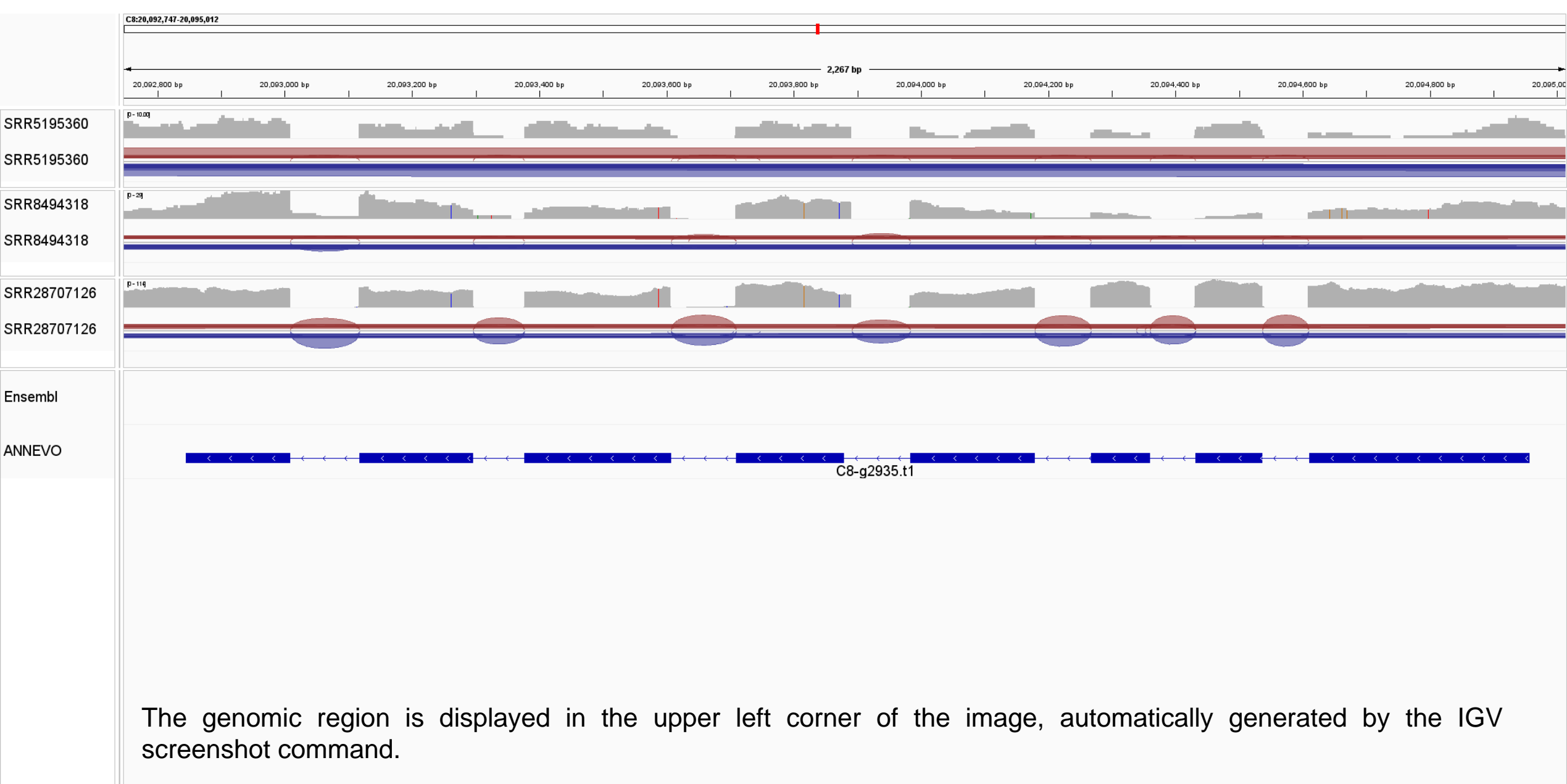


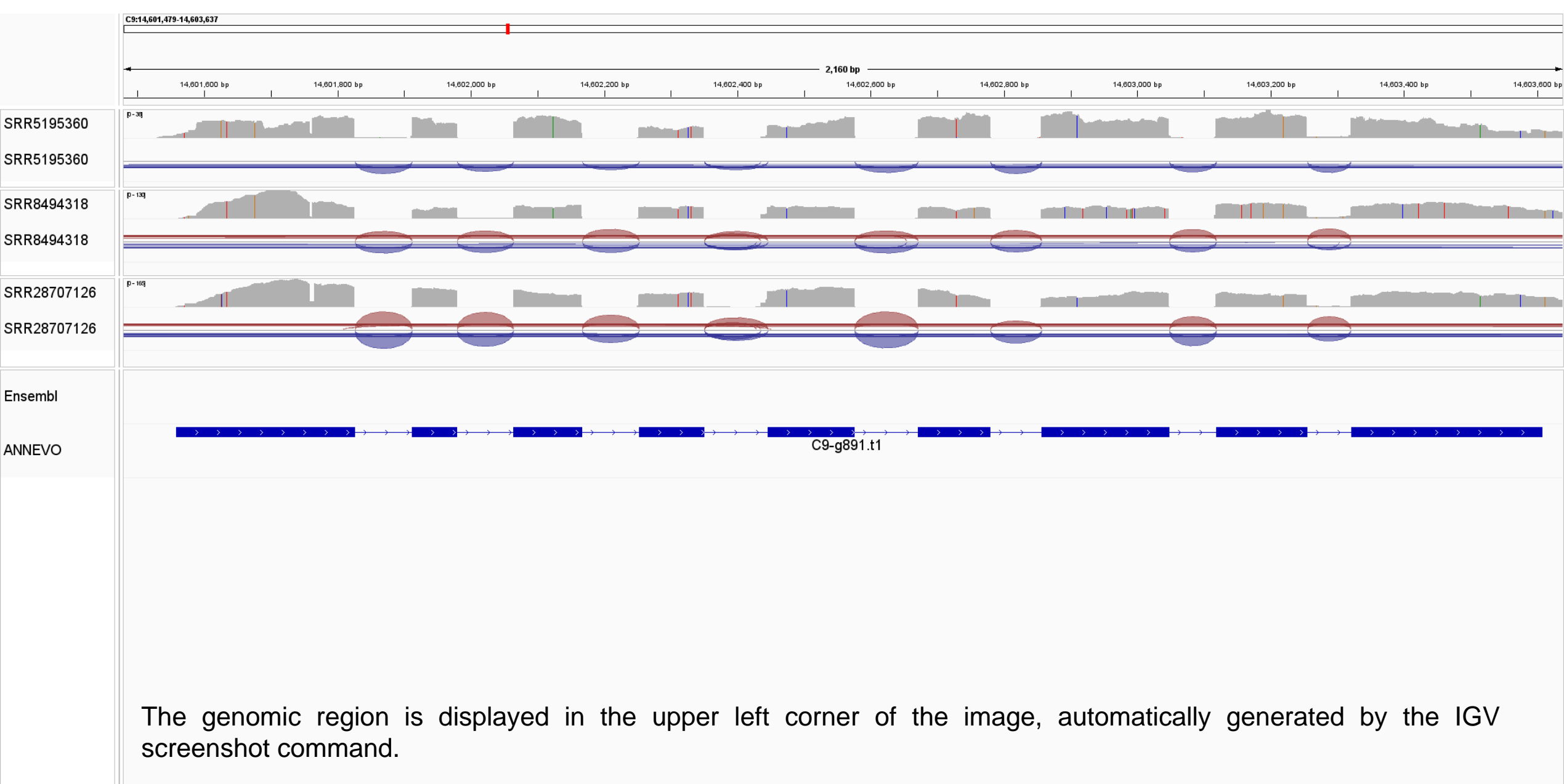


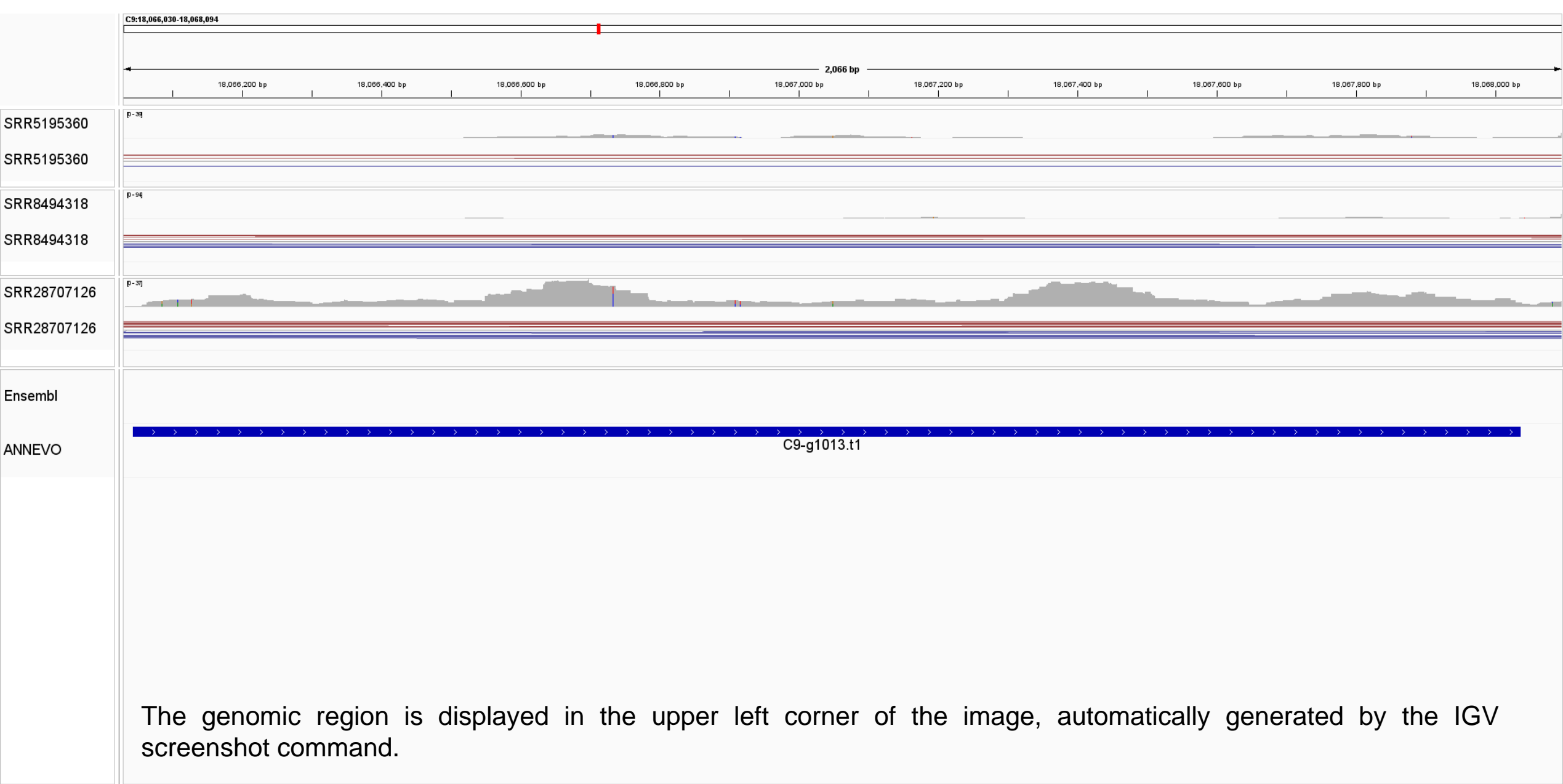










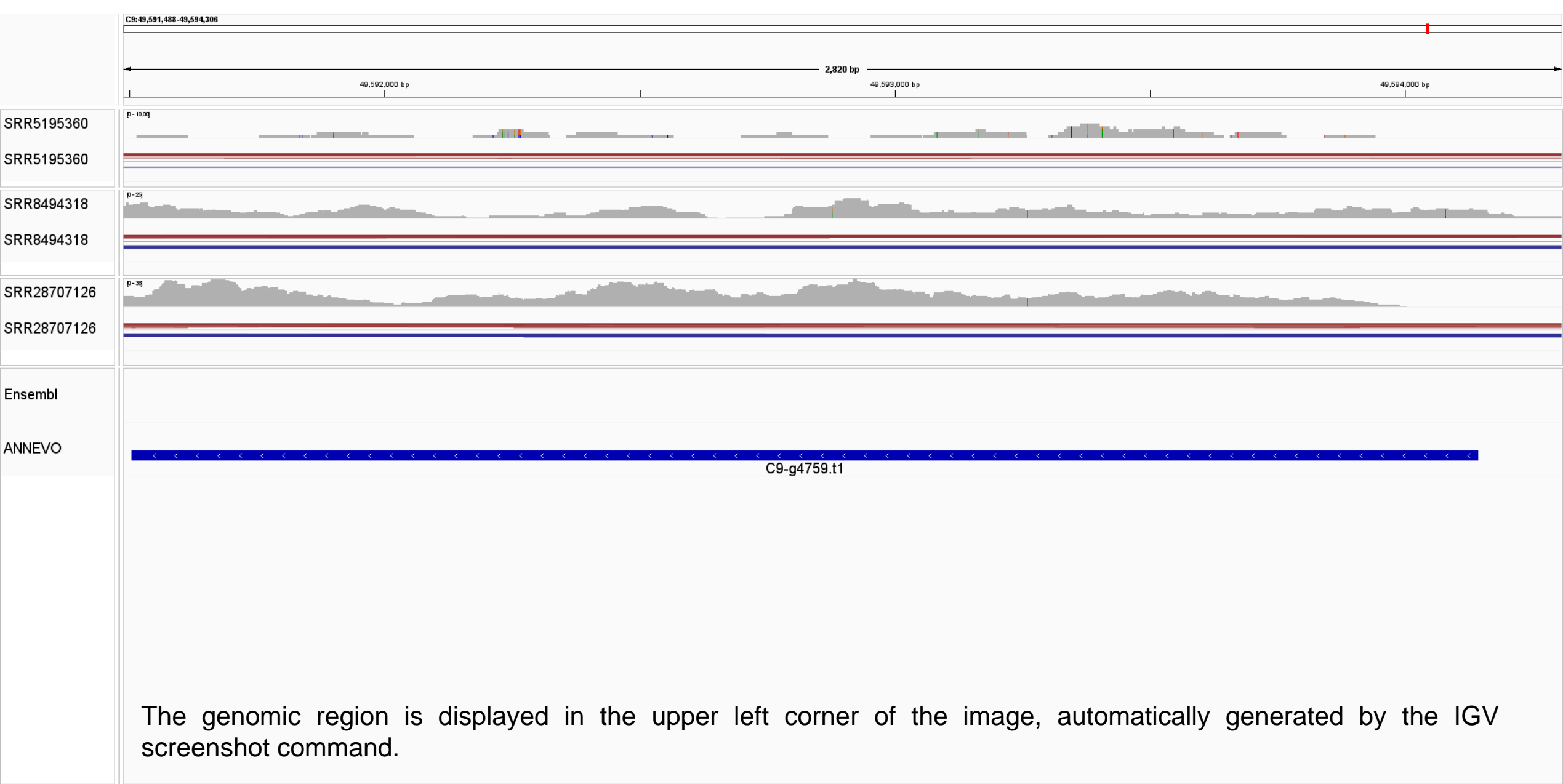


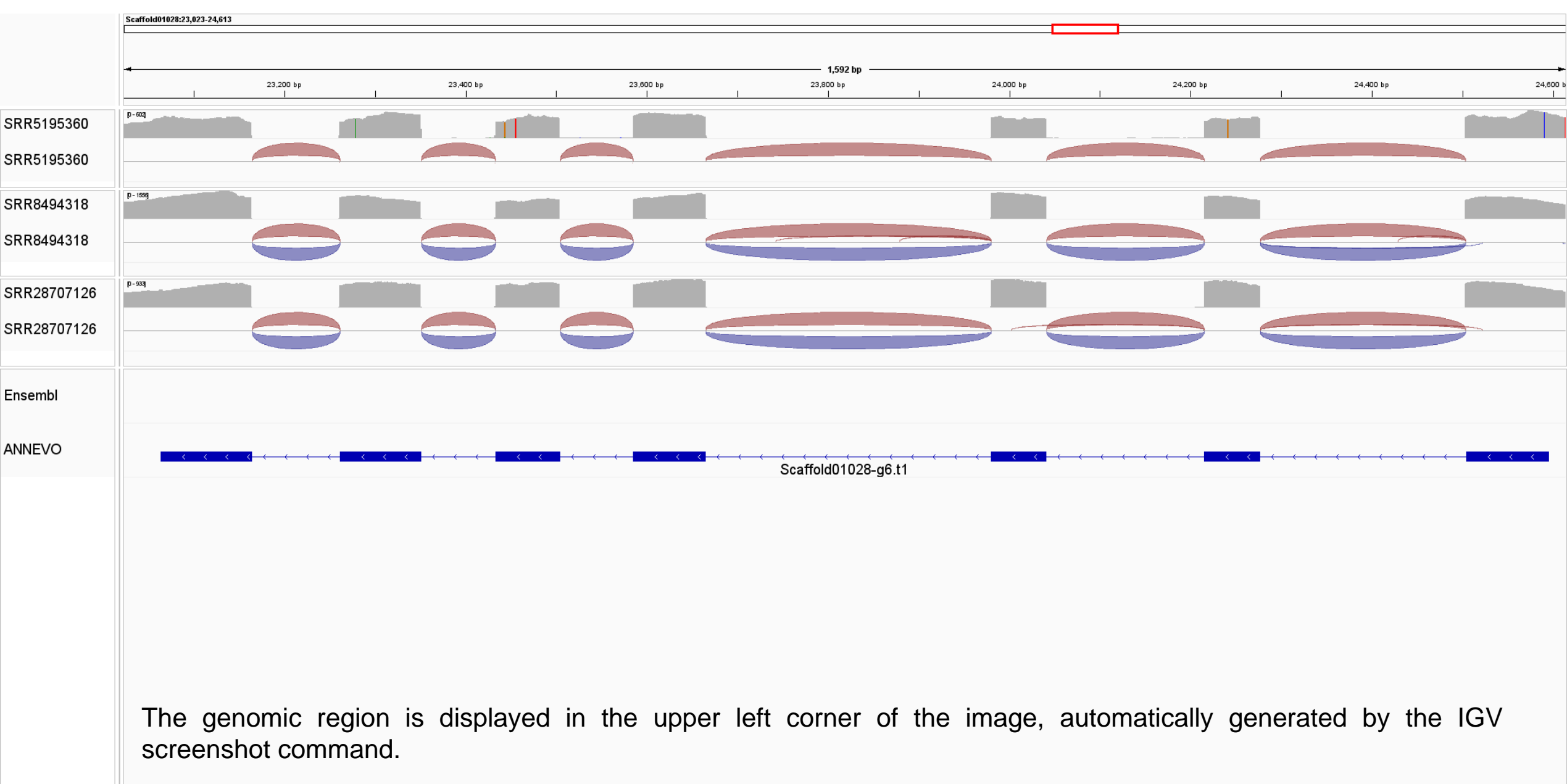






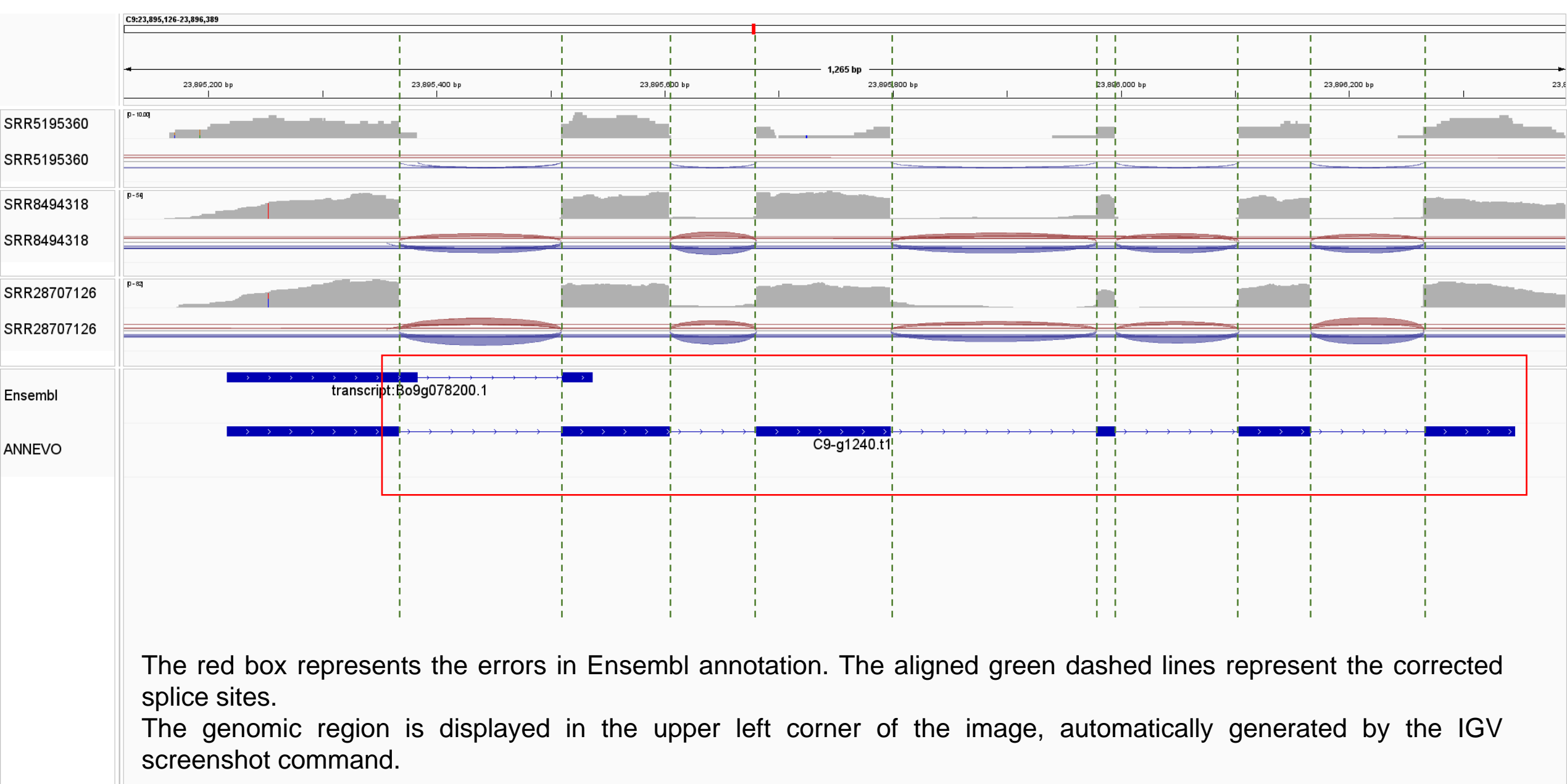


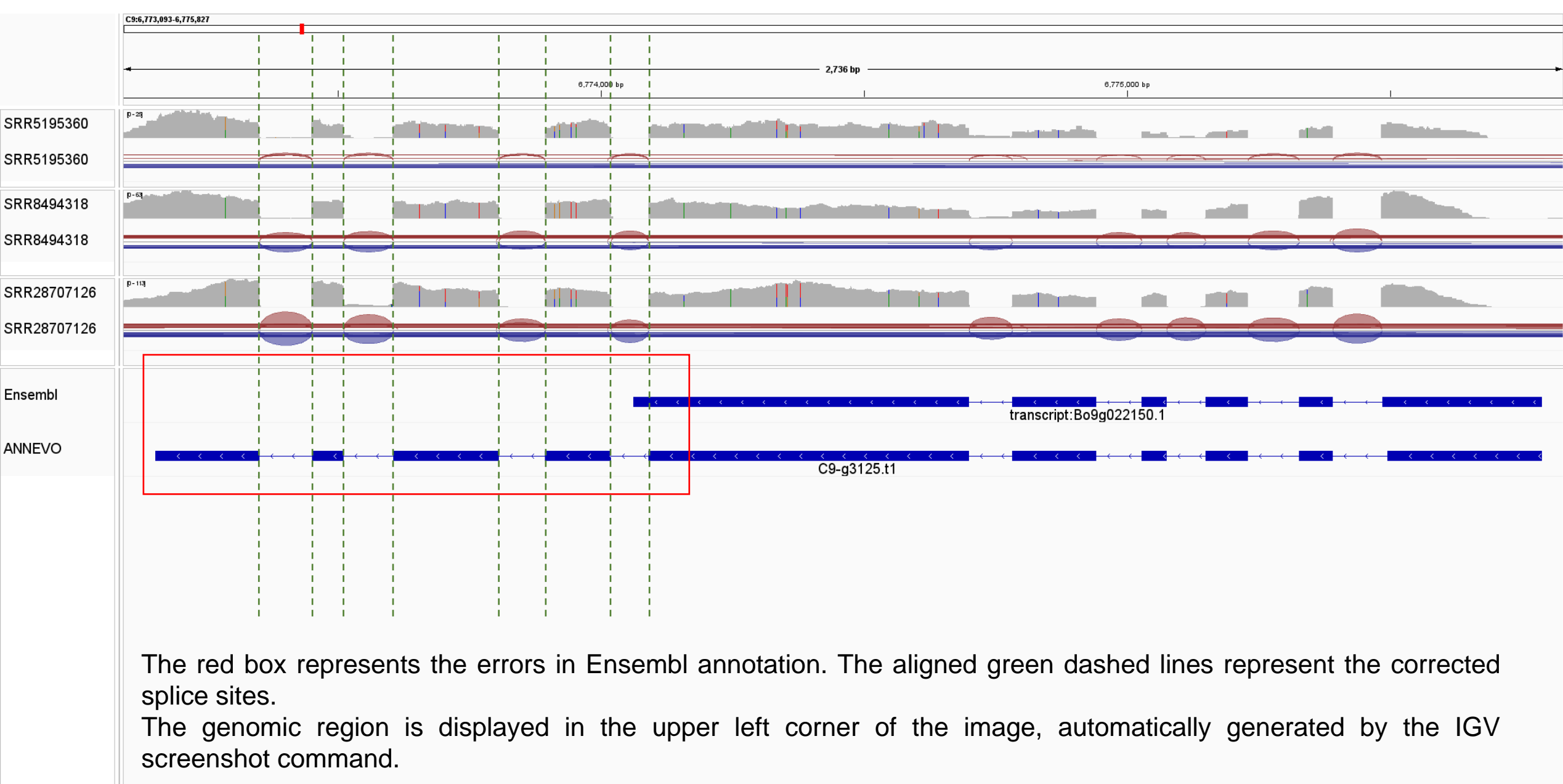




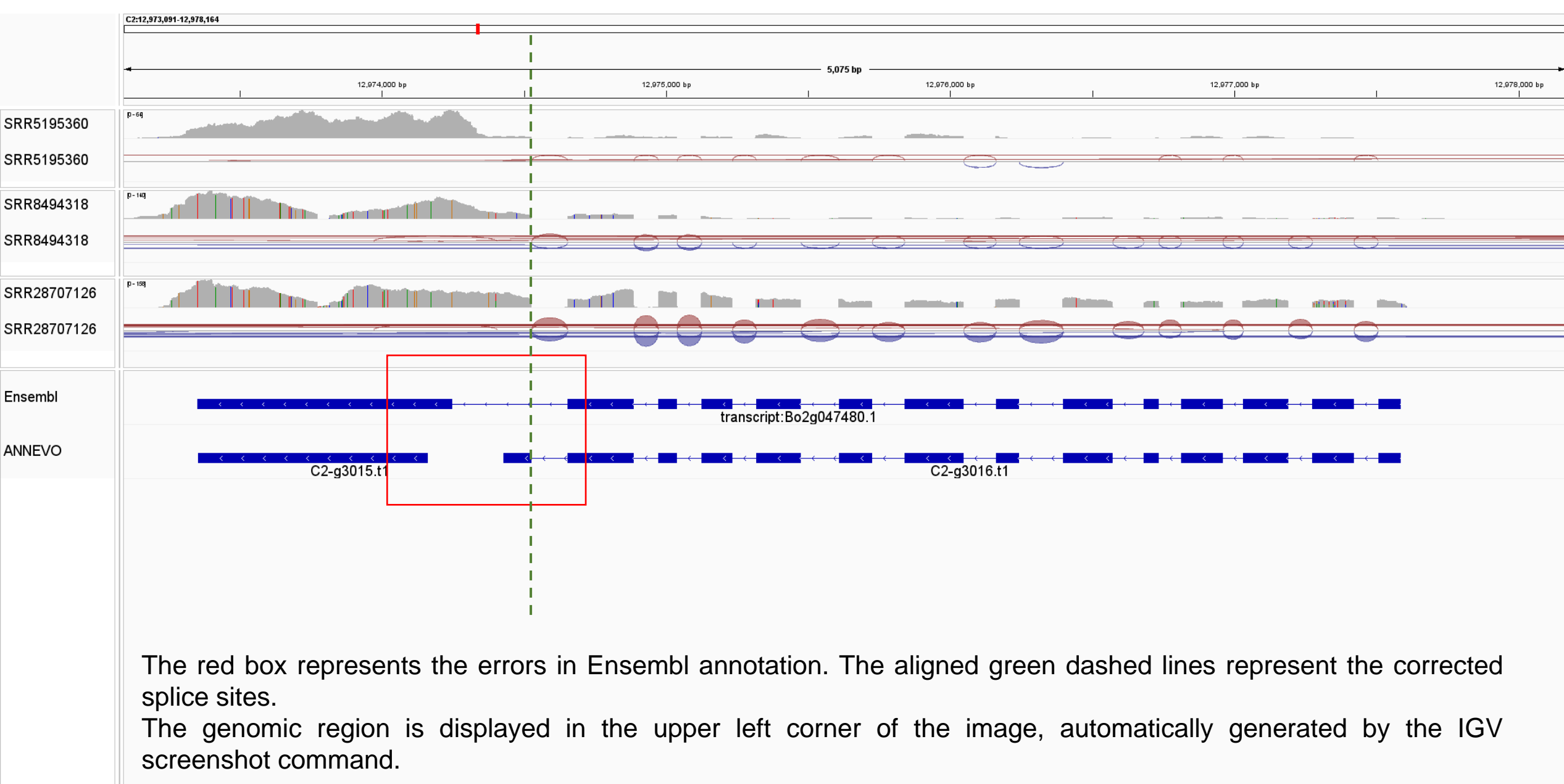
Incomplete Gene in Ensembl

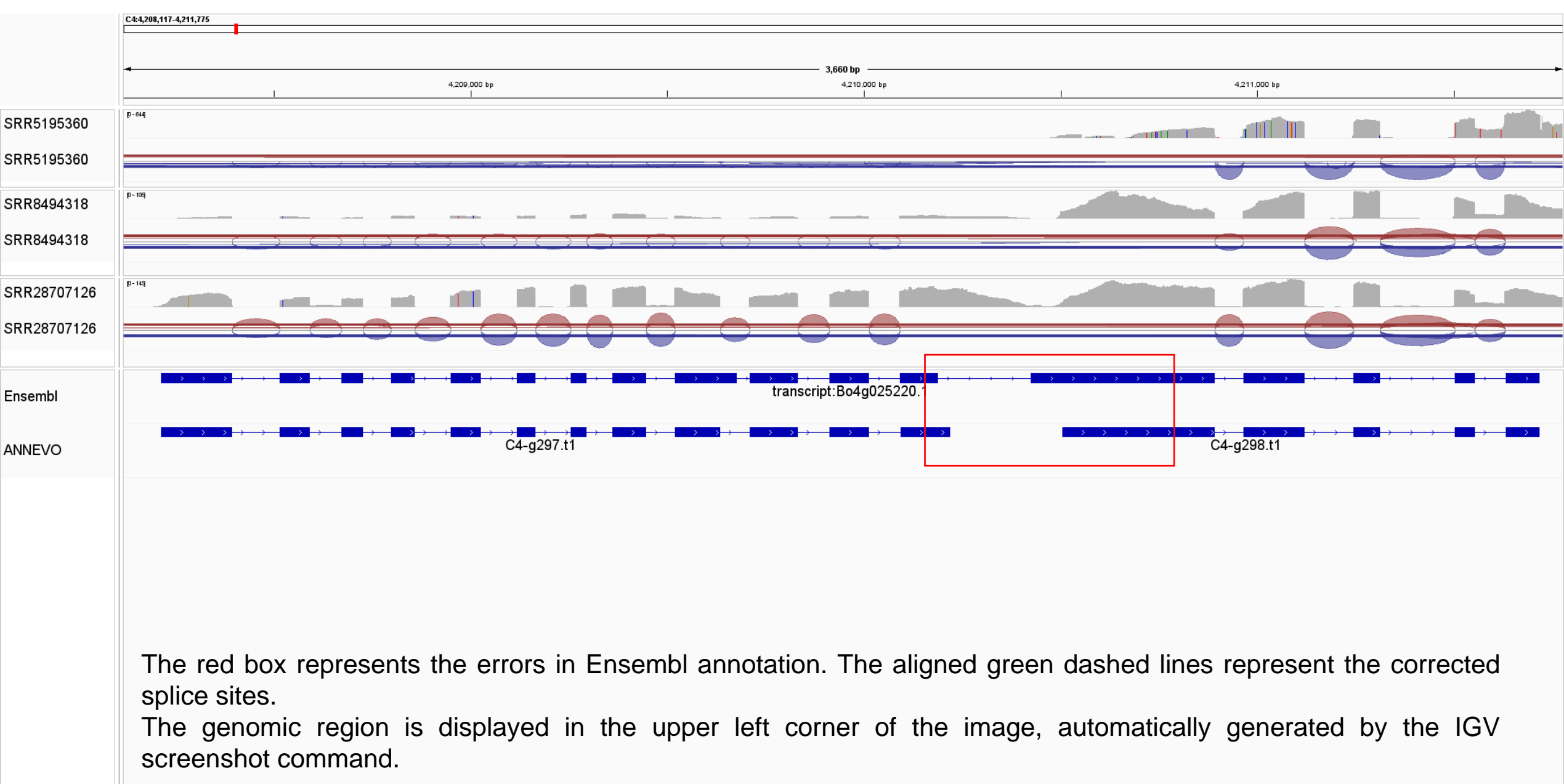


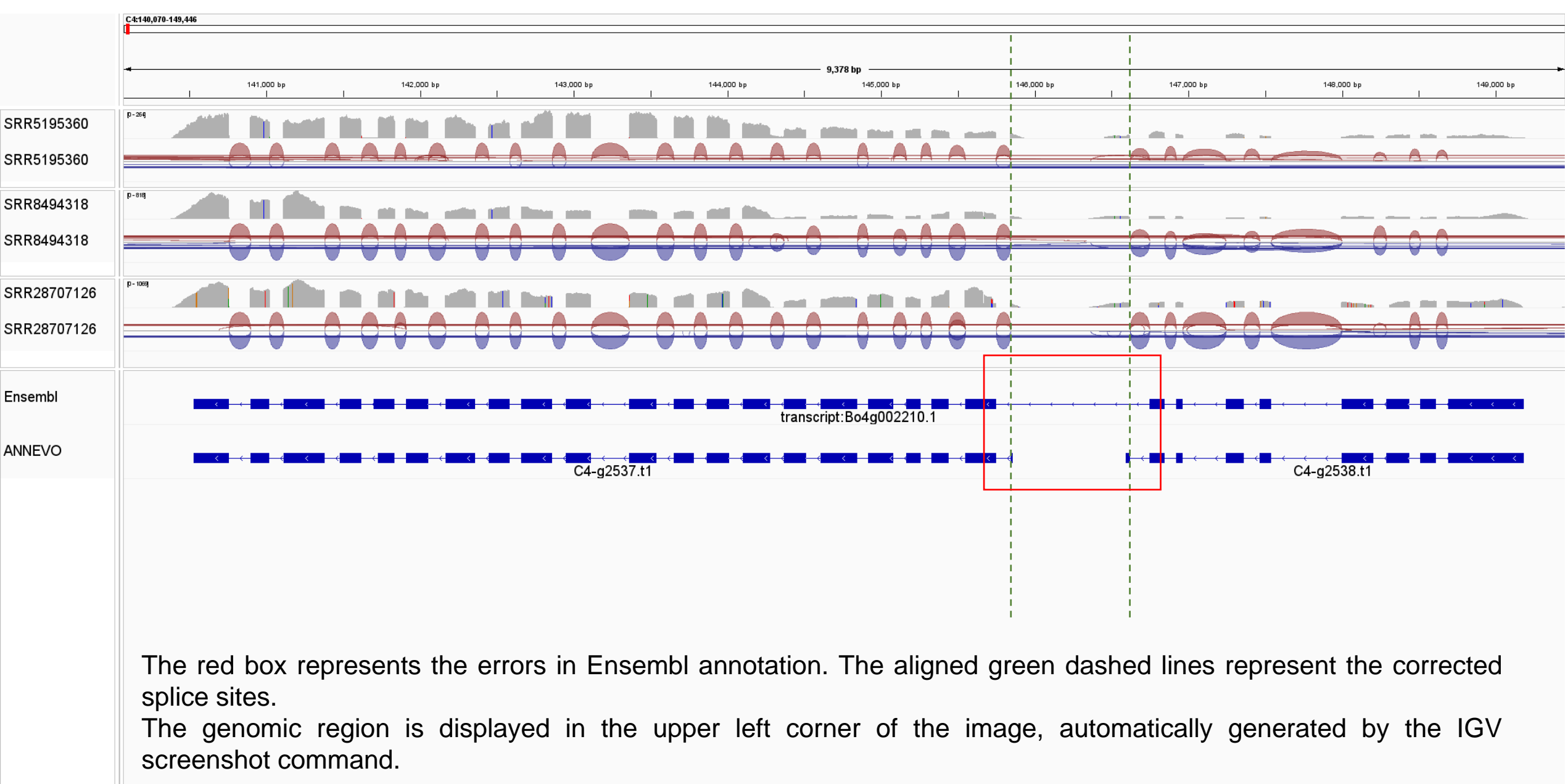




Erroneous Gene Fusion in Ensembl

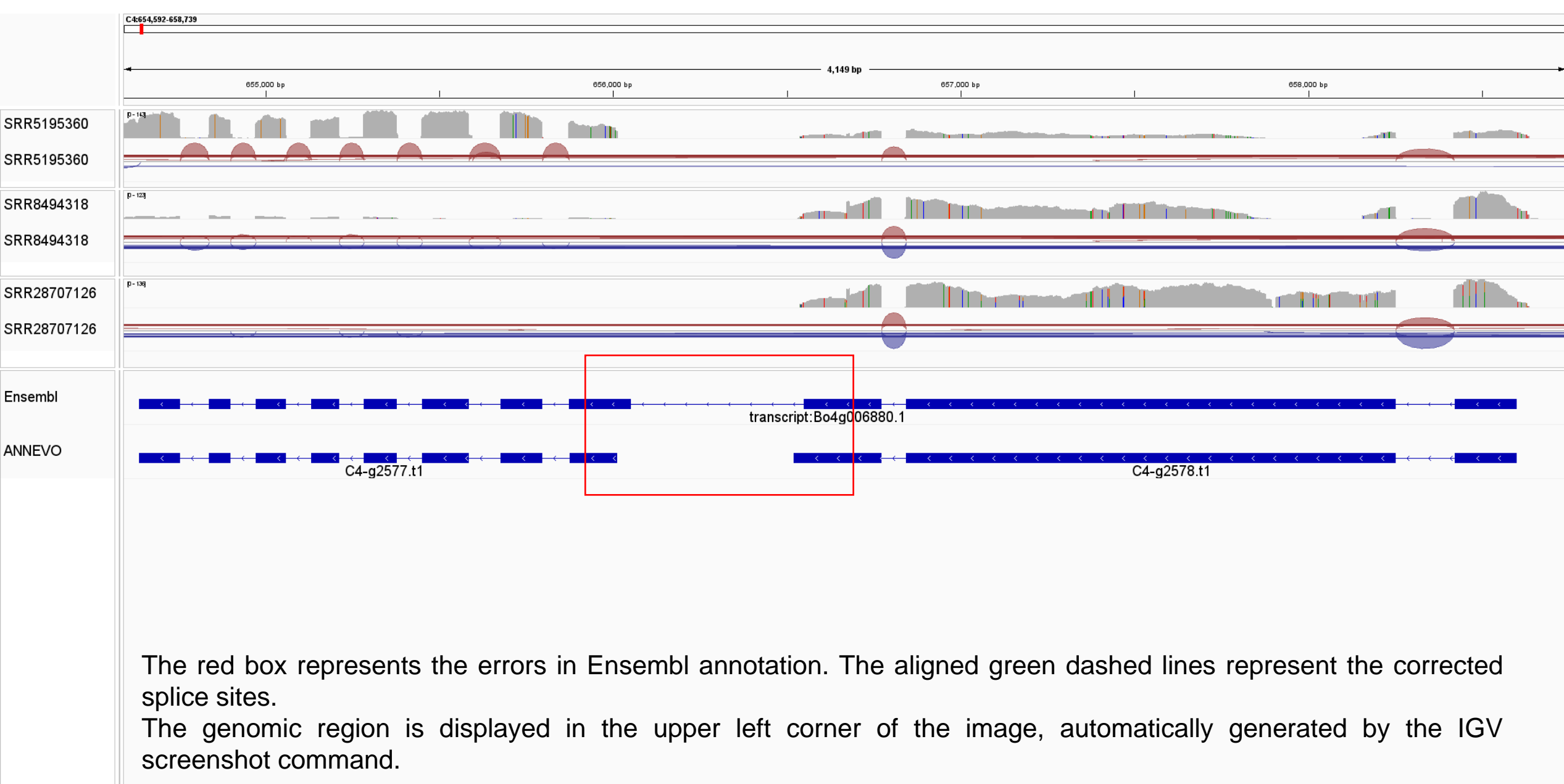


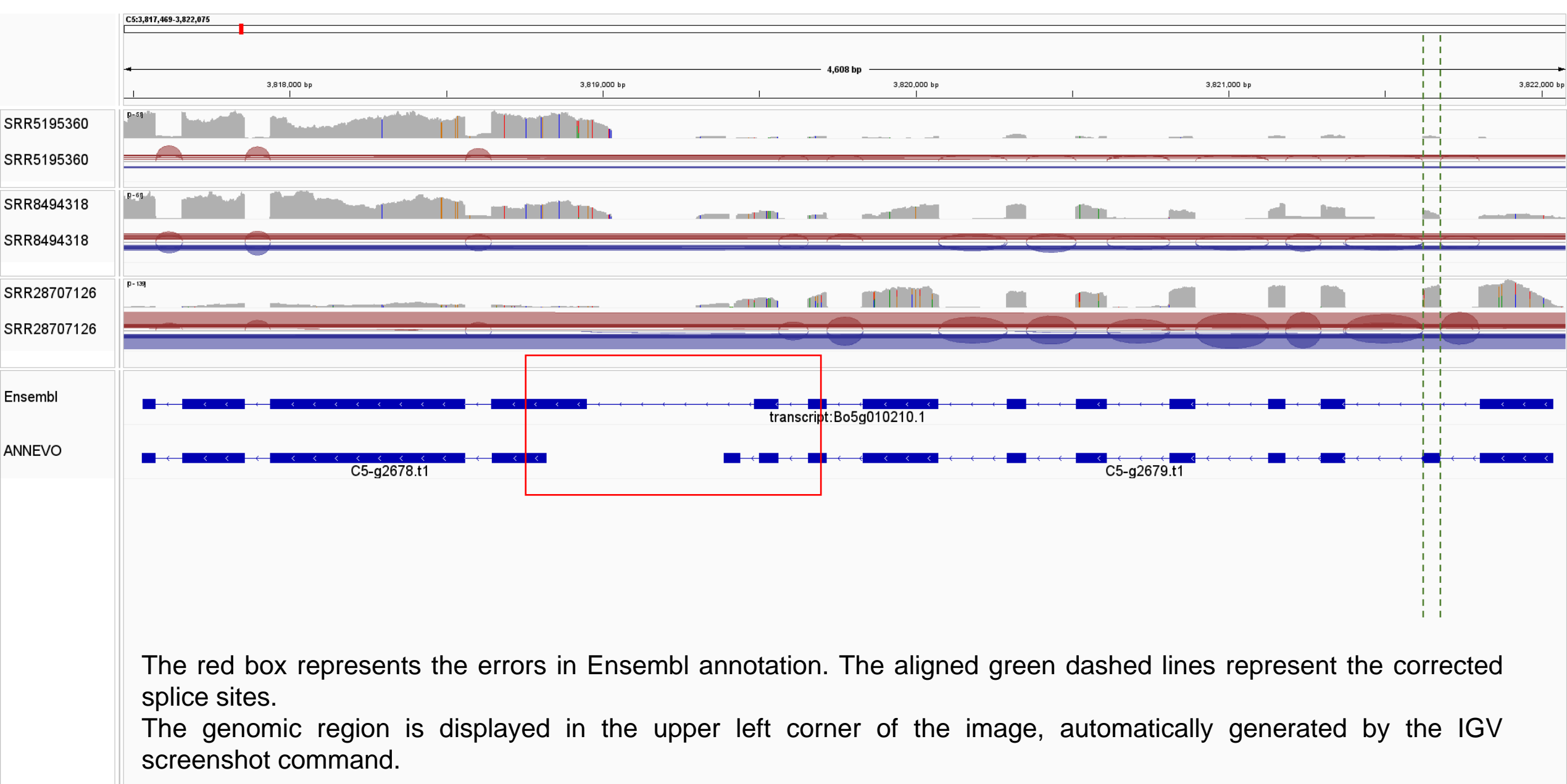


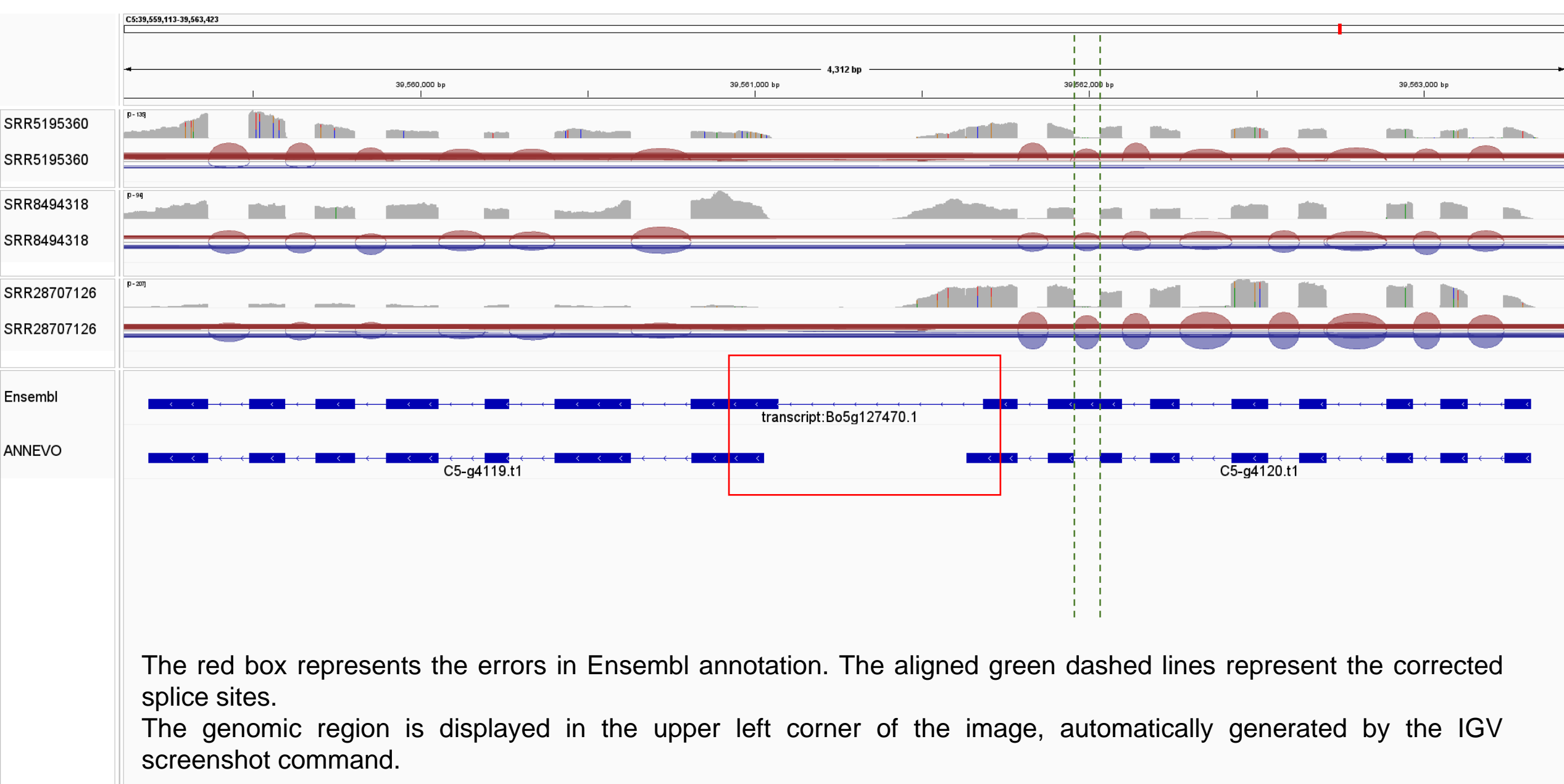


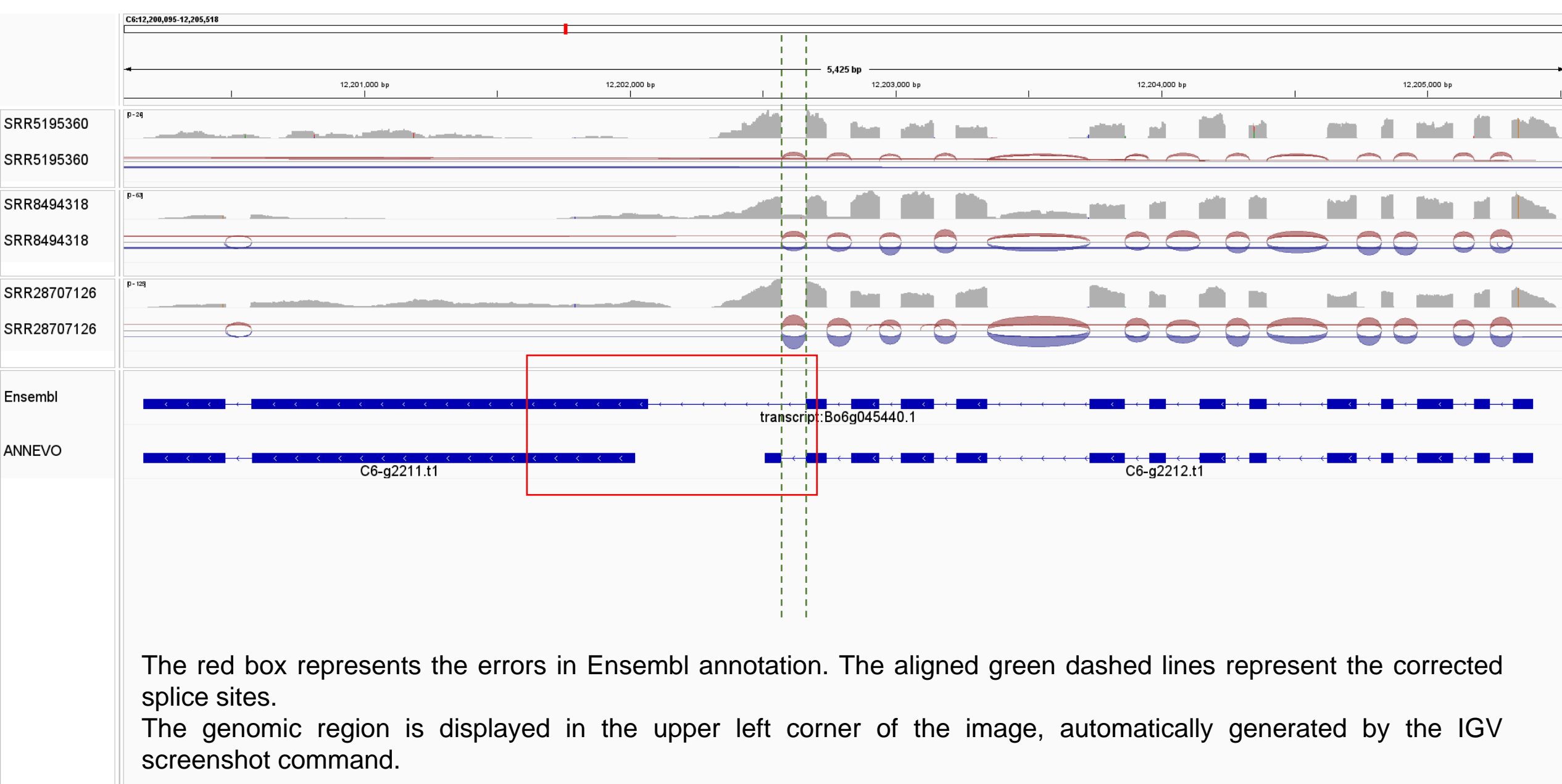
The red box represents the errors in Ensembl annotation. The aligned green dashed lines represent the corrected splice sites.

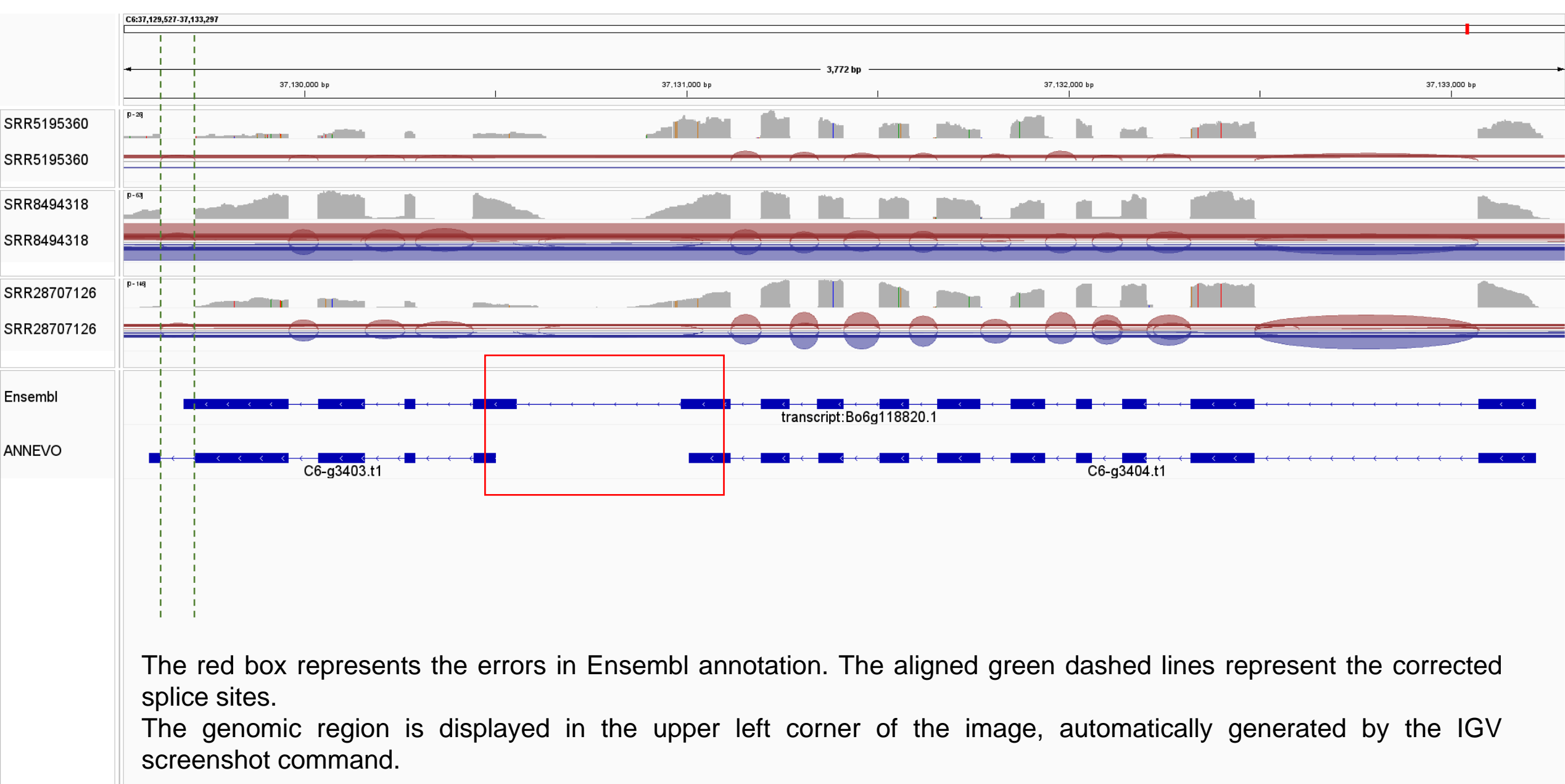
The genomic region is displayed in the upper left corner of the image, automatically generated by the IGV screenshot command.





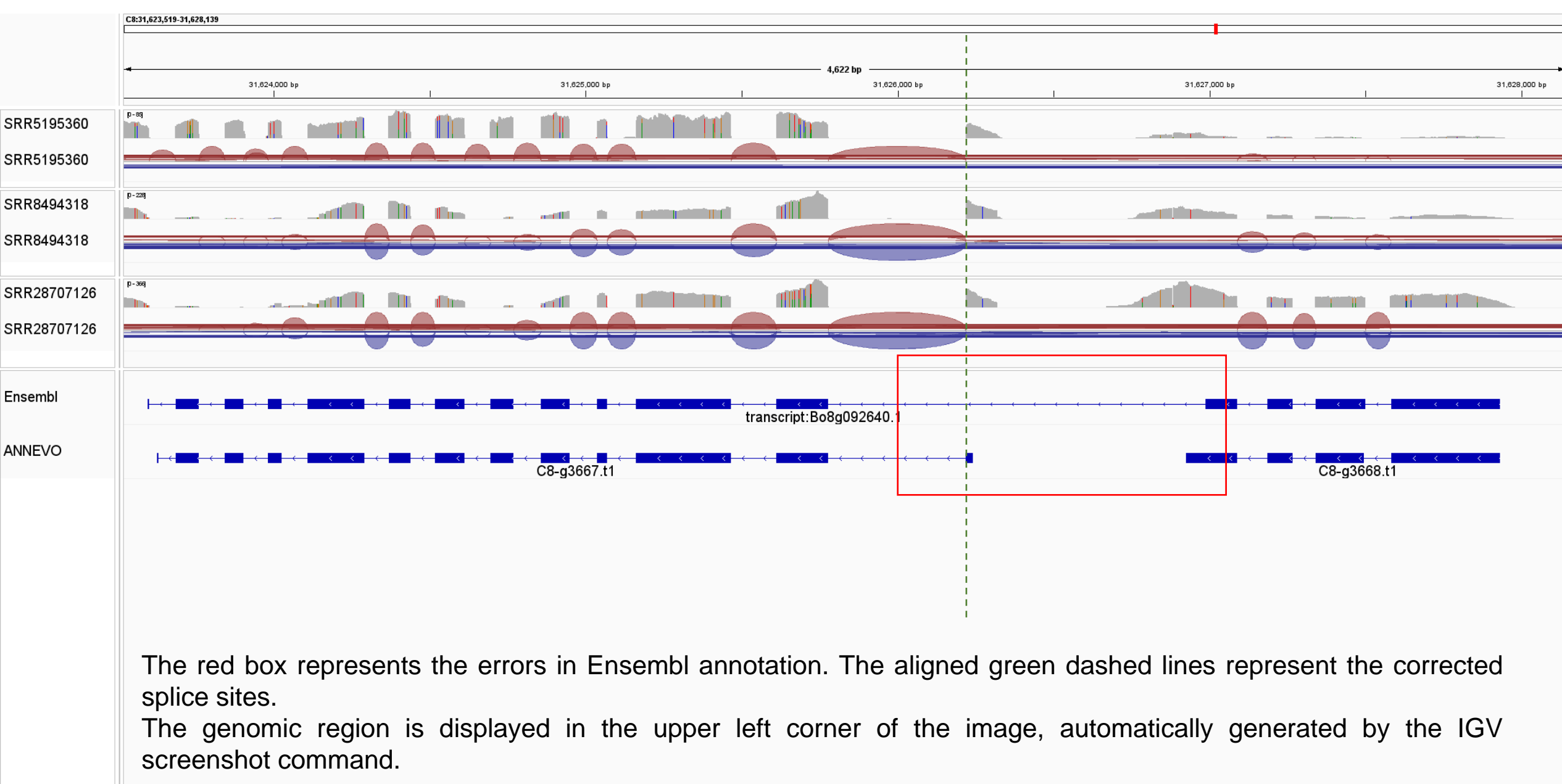


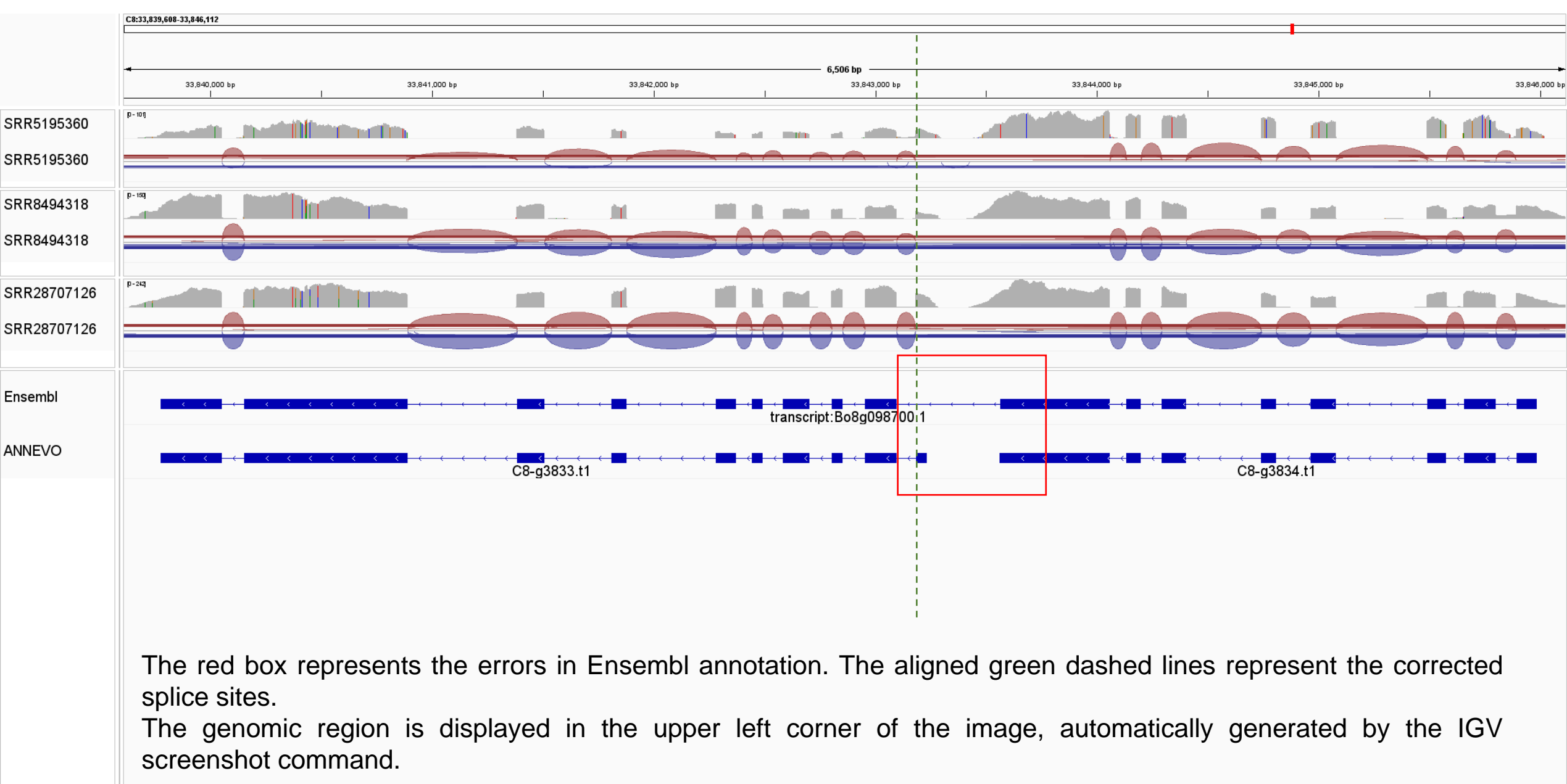




The red box represents the errors in Ensembl annotation. The aligned green dashed lines represent the corrected splice sites.

The genomic region is displayed in the upper left corner of the image, automatically generated by the IGV screenshot command.





Erroneous Gene Break in Ensembl

