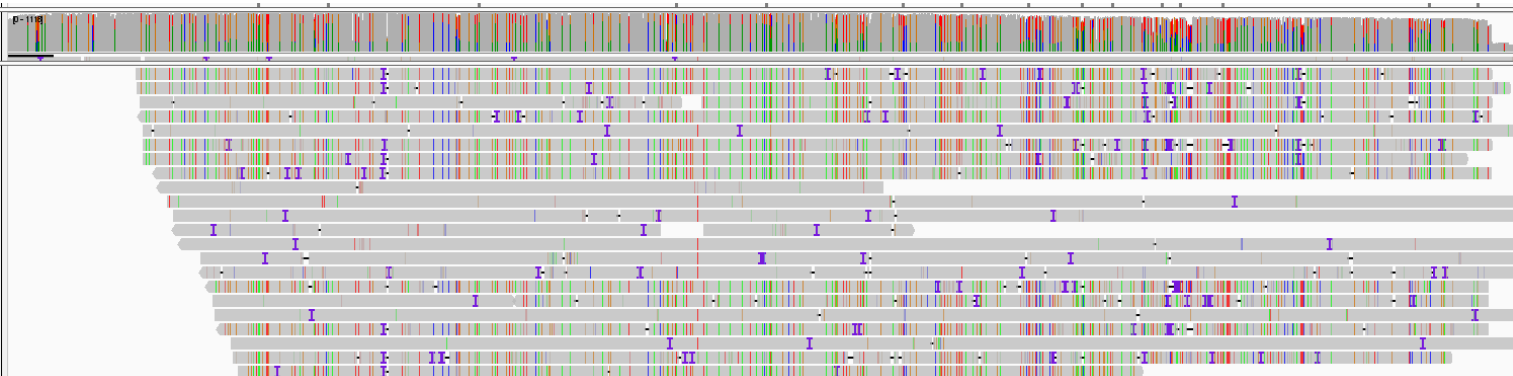


IGV screenshots of possible gene duplication

A



B



Supplementary Figure 7: Integrative Genomics Viewer screenshot of a genomic area that shows read with variant patterns of a possible gene duplication. Shown are A) a zoomed-in position of gene EFAU004_00073 that shows that most of the found variants indicate two different bases with around 50% ratio each and B) the full length of gene EFAU004_00073 that show, that all reads exhibit one of two possible variant patterns.