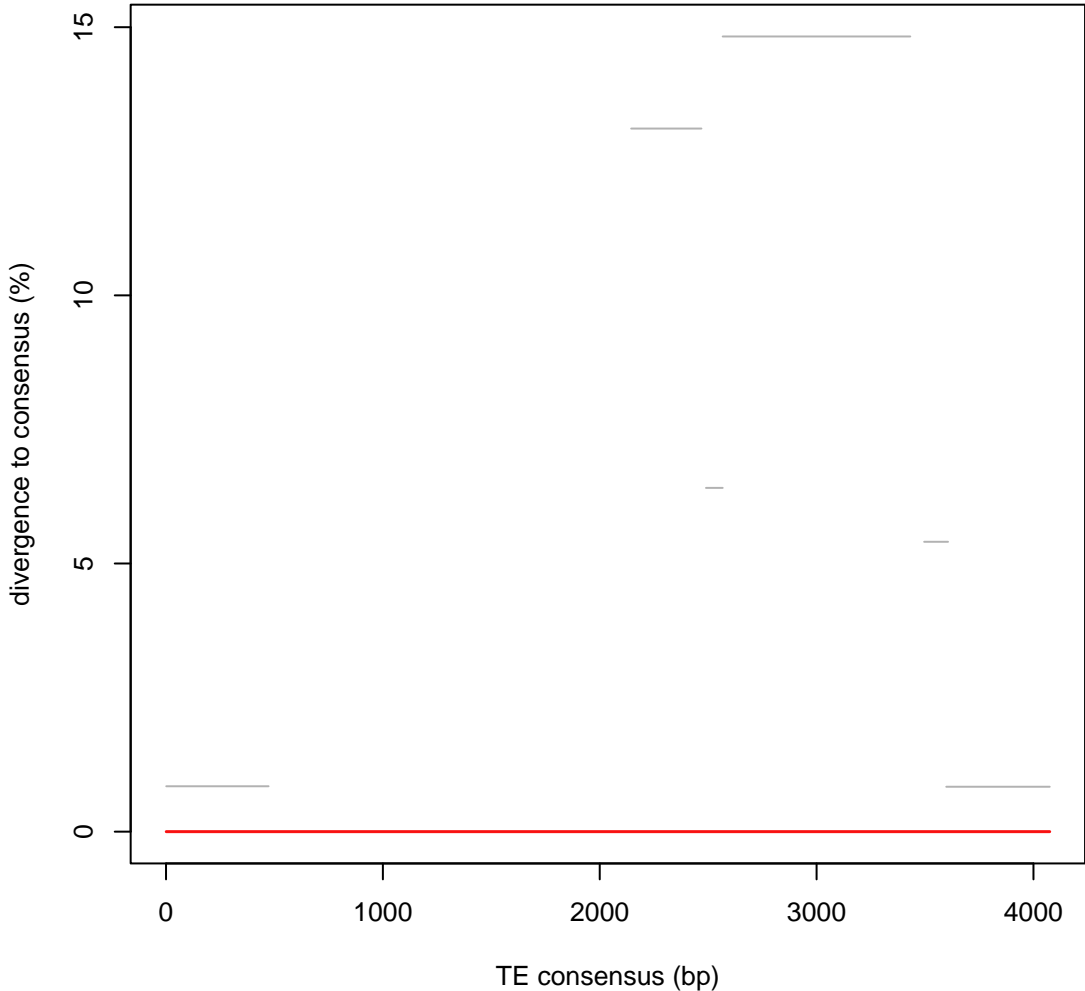
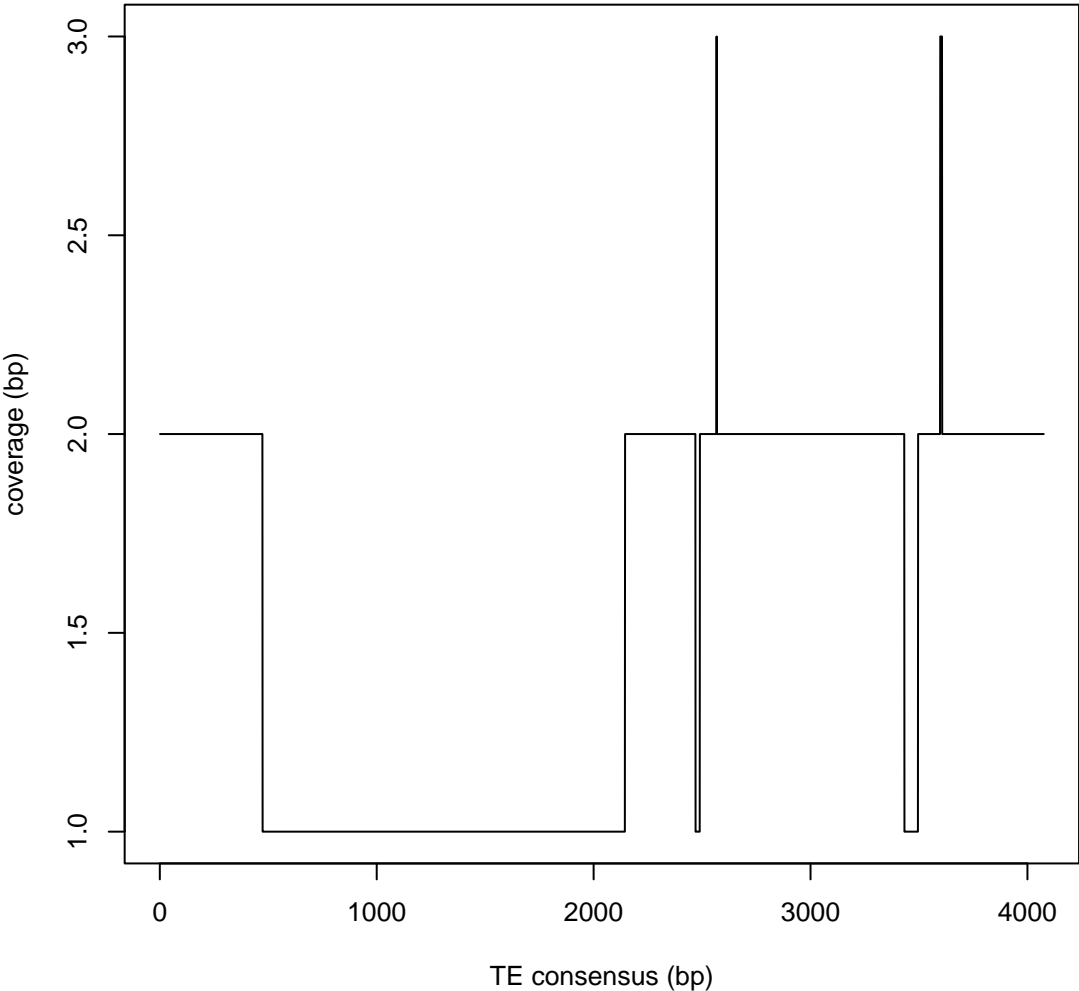


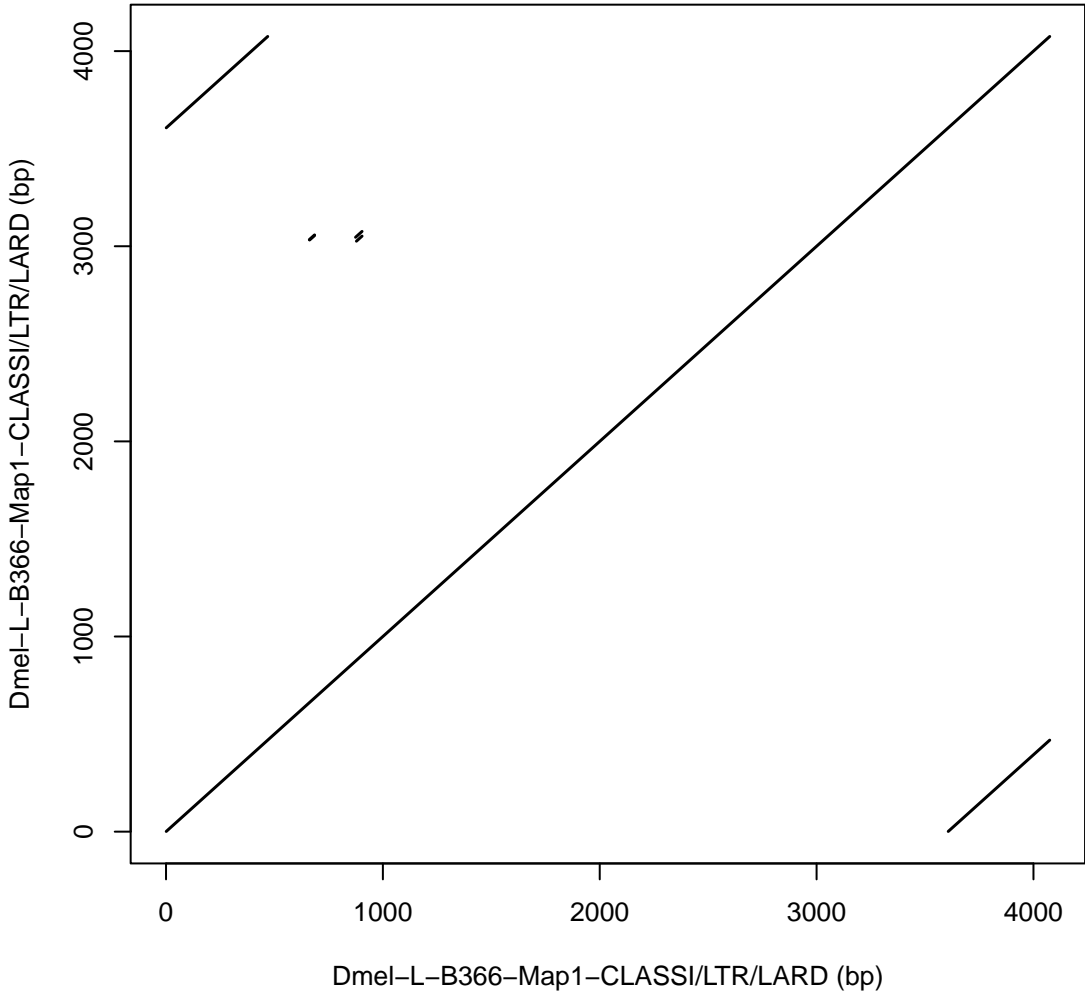
TE: Dmel-L-B366-Map1-CLASSI/LTR/LARD
consensus size: 4075bp; fragments: 7; full length: 1 (>=3667.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

