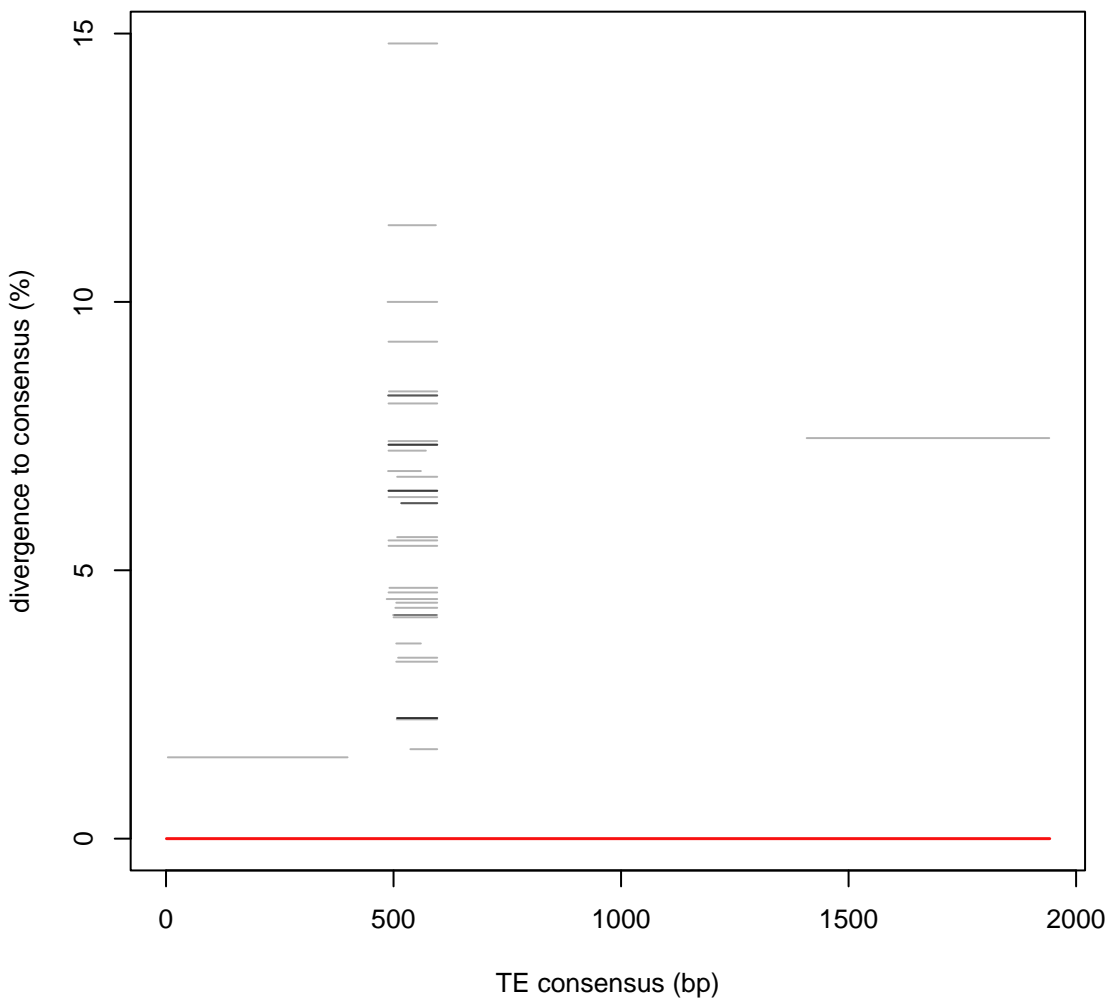
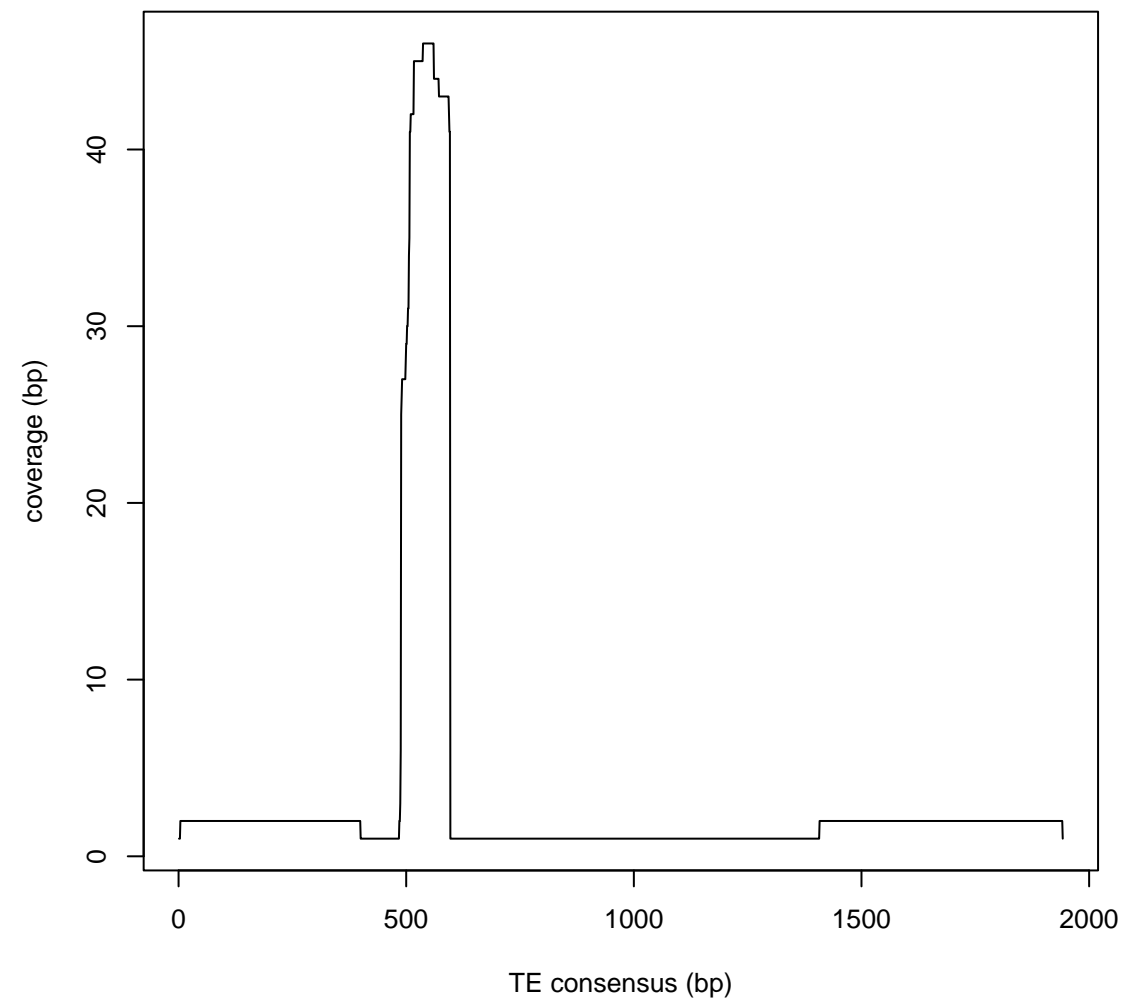


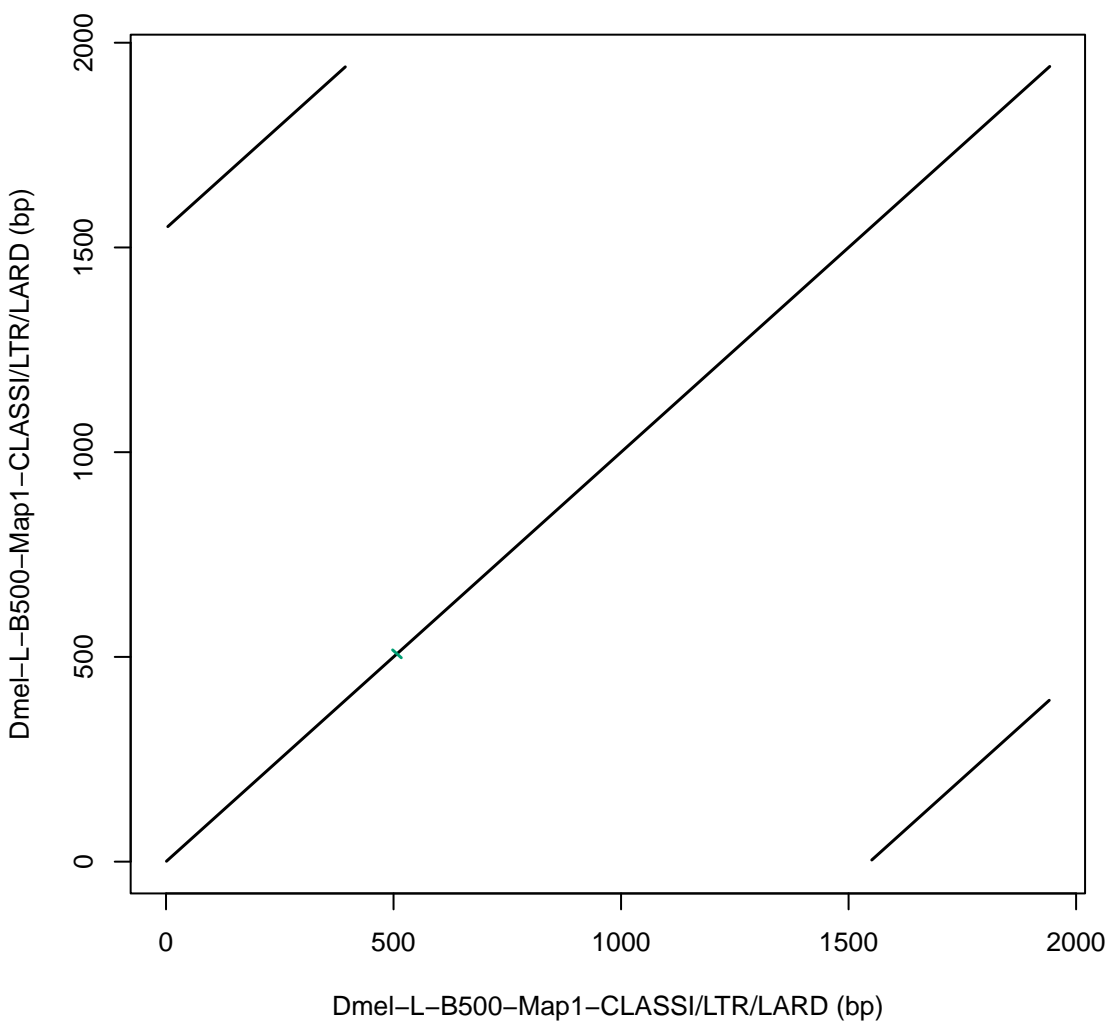
TE: Dmel-L-B500-Map1-CLASSI/LTR/LARD  
consensus size: 1942bp; fragments: 48; full length: 1 (>=1747.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

