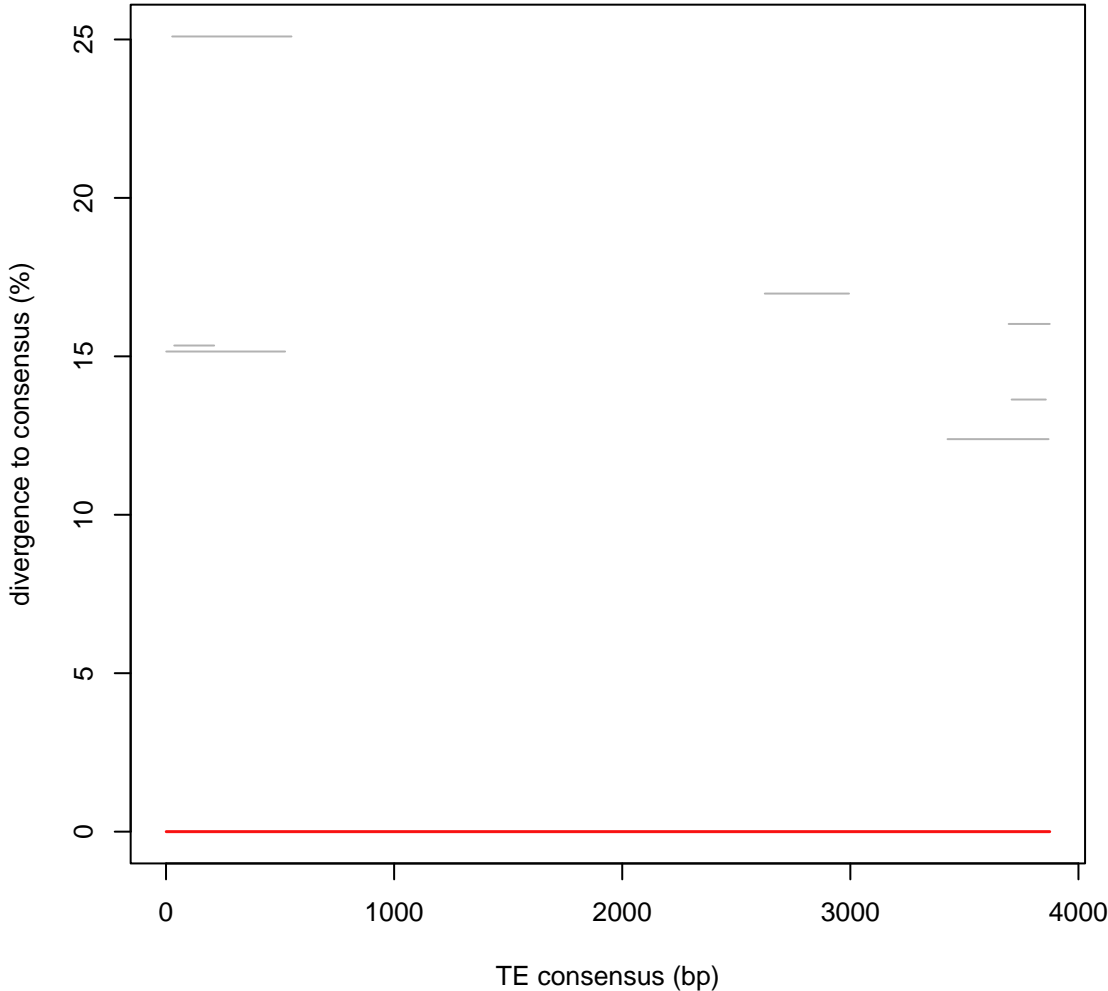
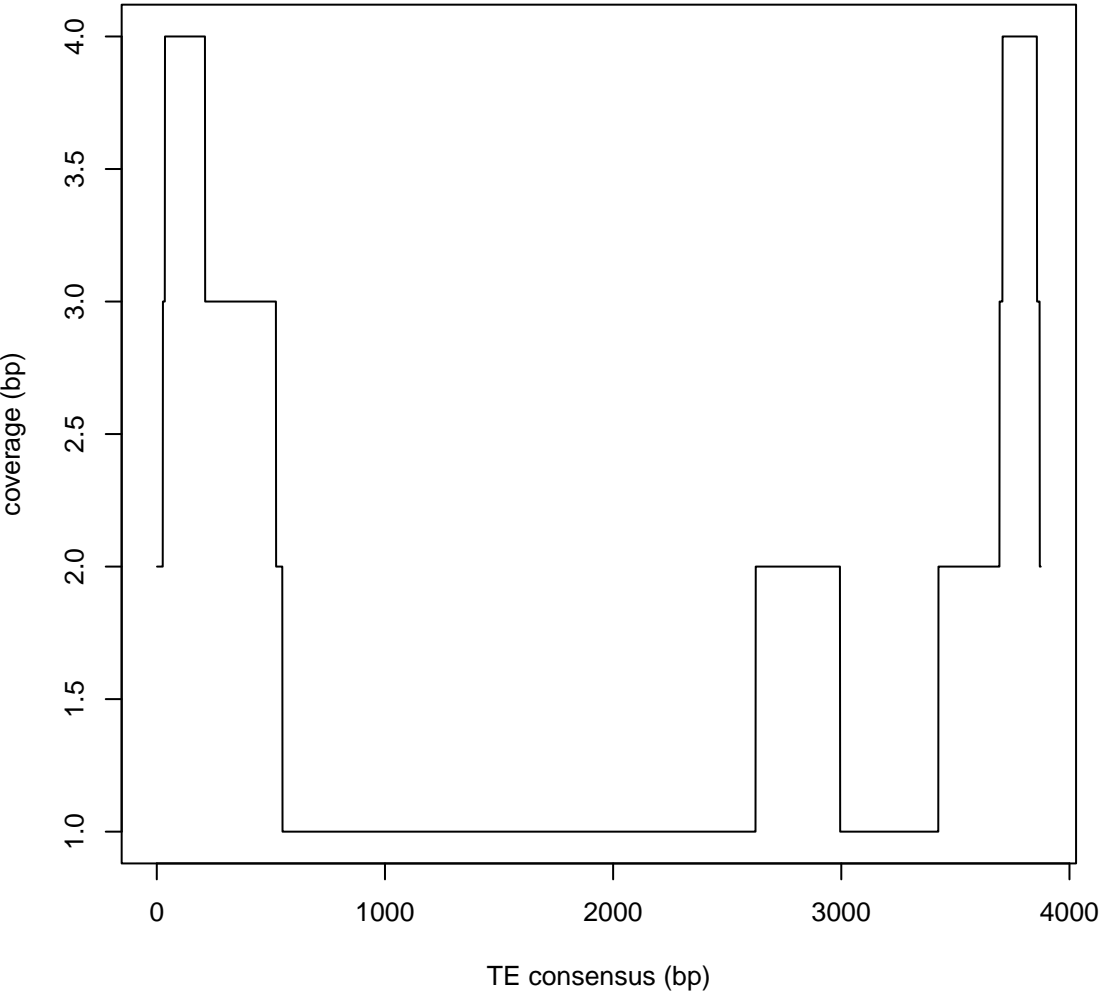


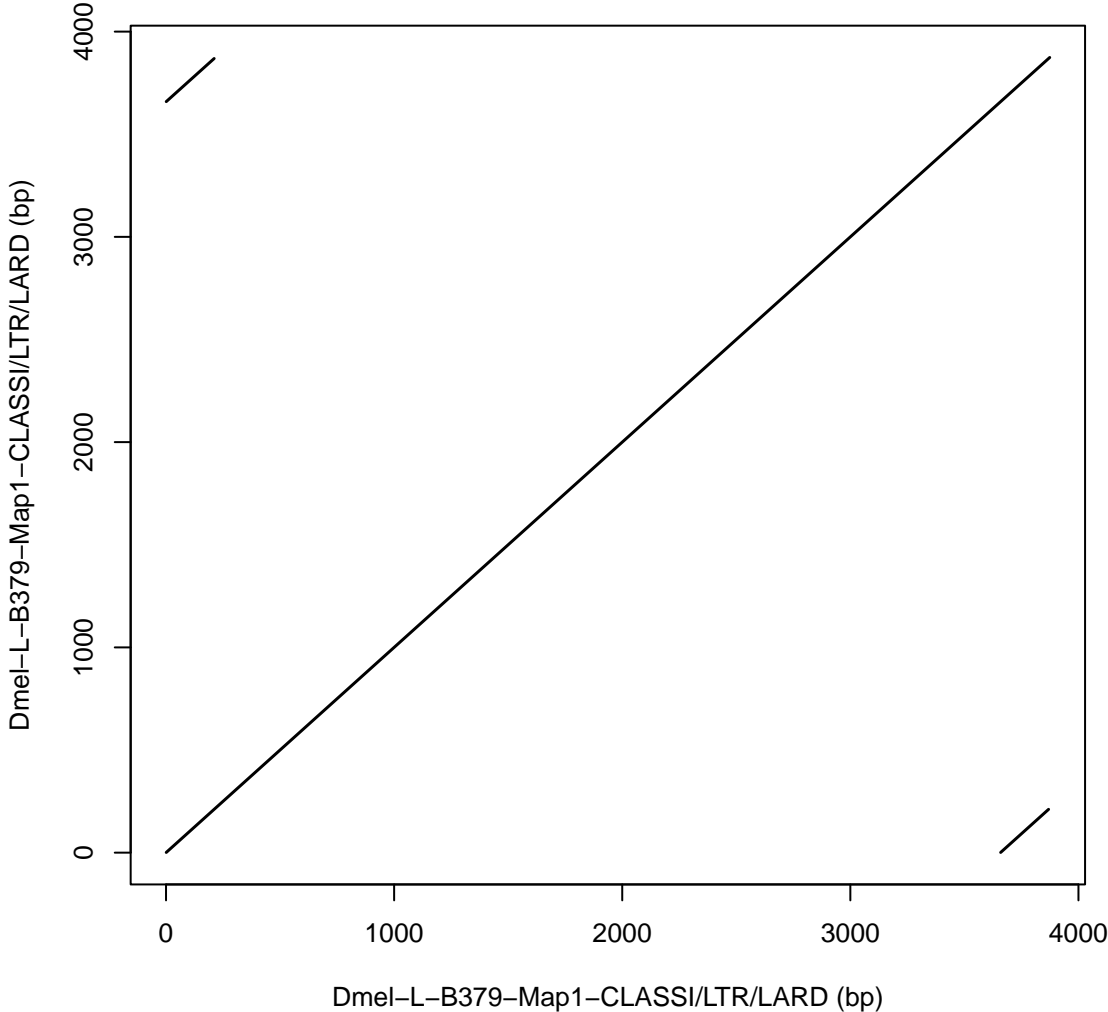
TE: Dmel-L-B379-Map1-CLASSI/LTR/LARD
consensus size: 3874bp; fragments: 8; full length: 1 (>=3486.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

