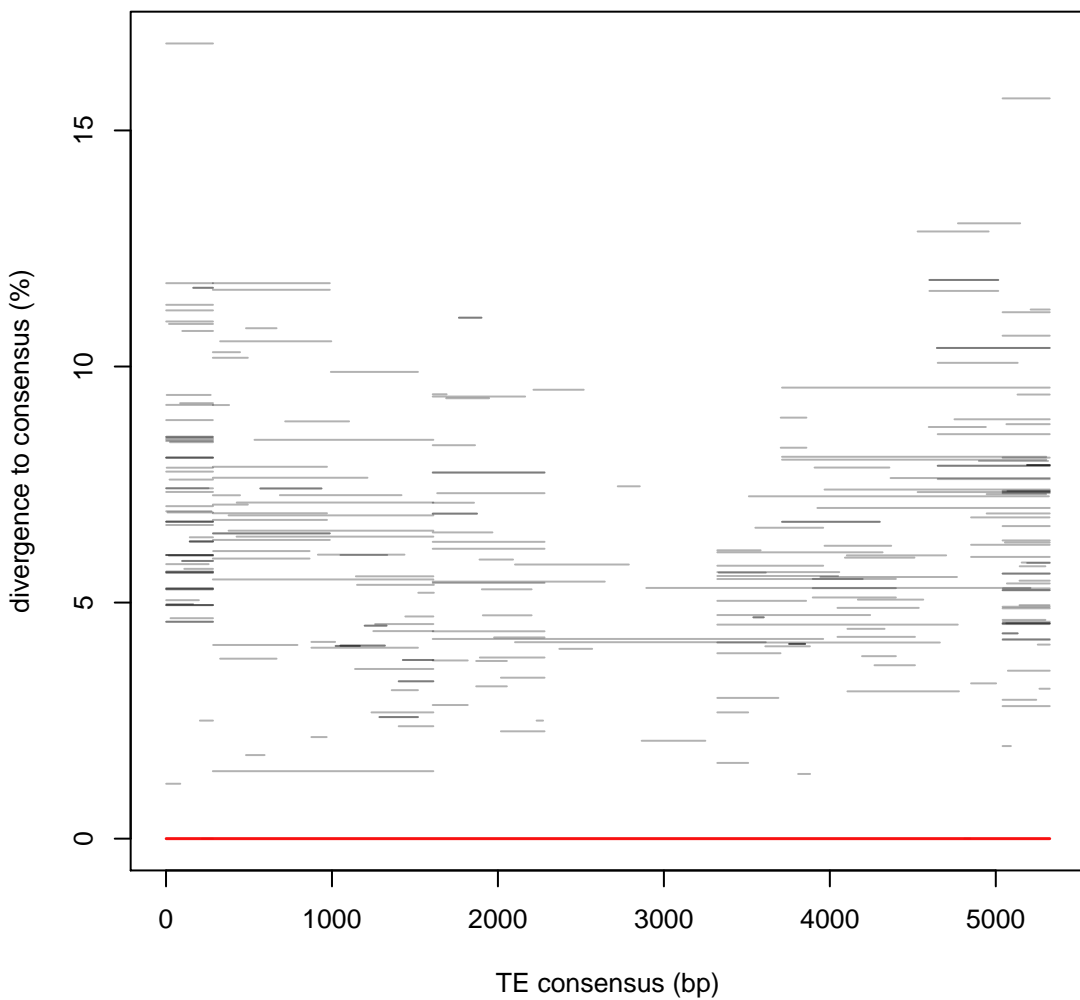
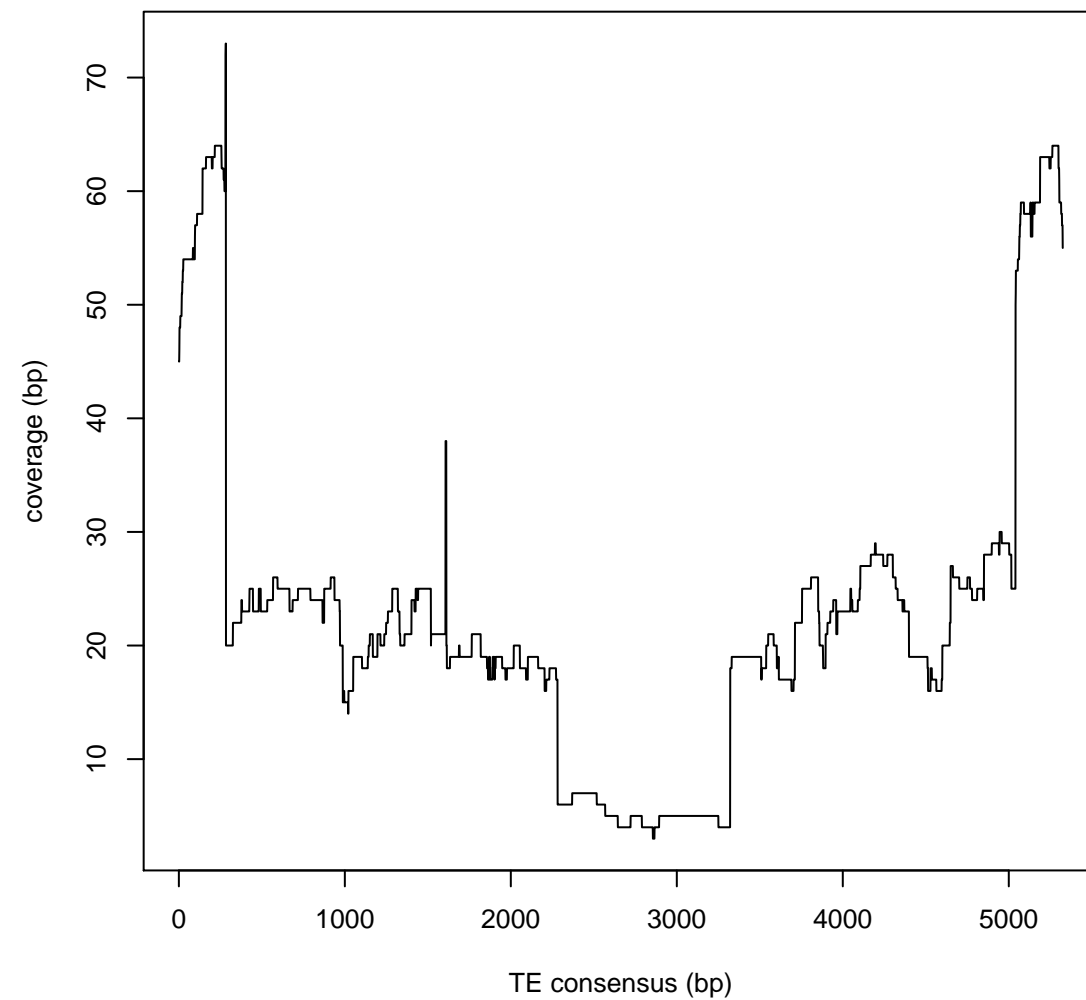


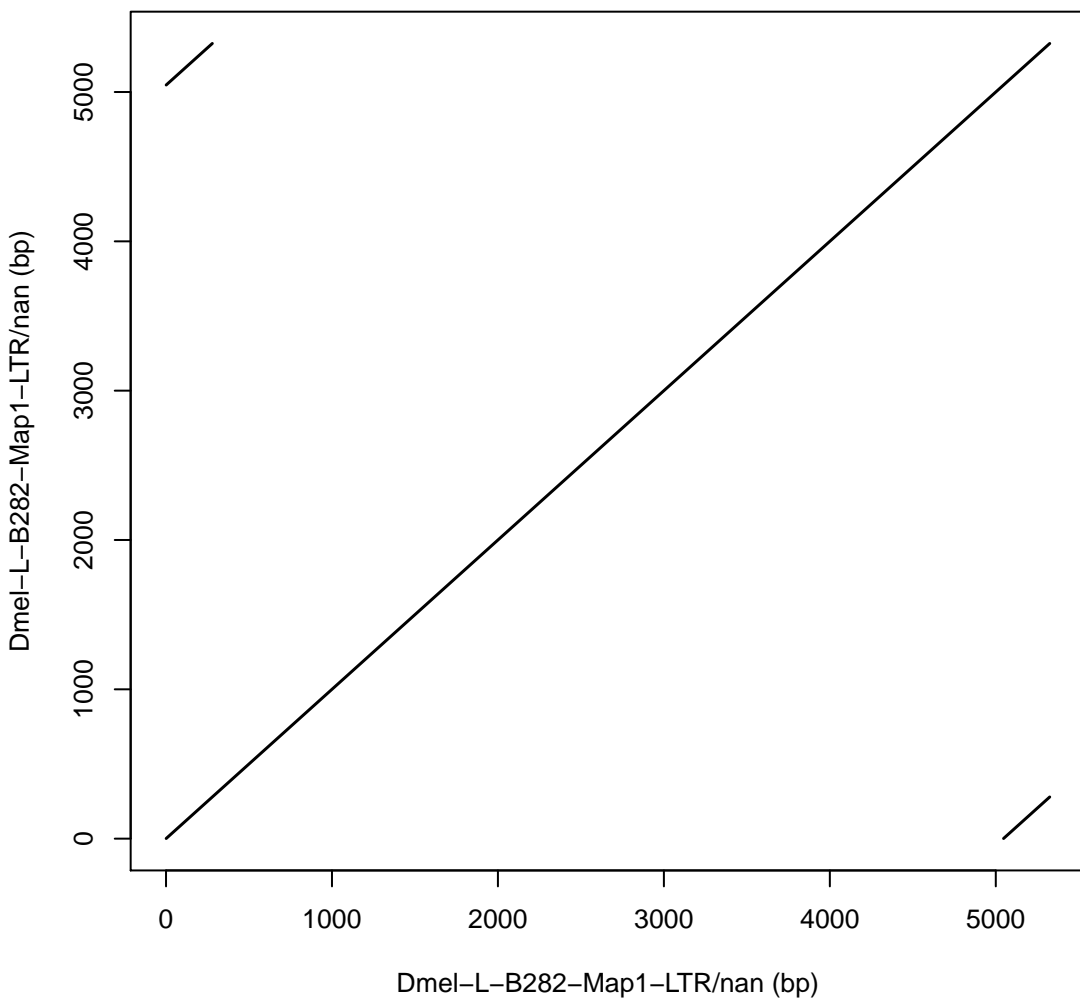
TE: Dmel-L-B282-Map1-LTR/nan
consensus size: 5325bp; fragments: 283; full length: 1 (>=4792.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

