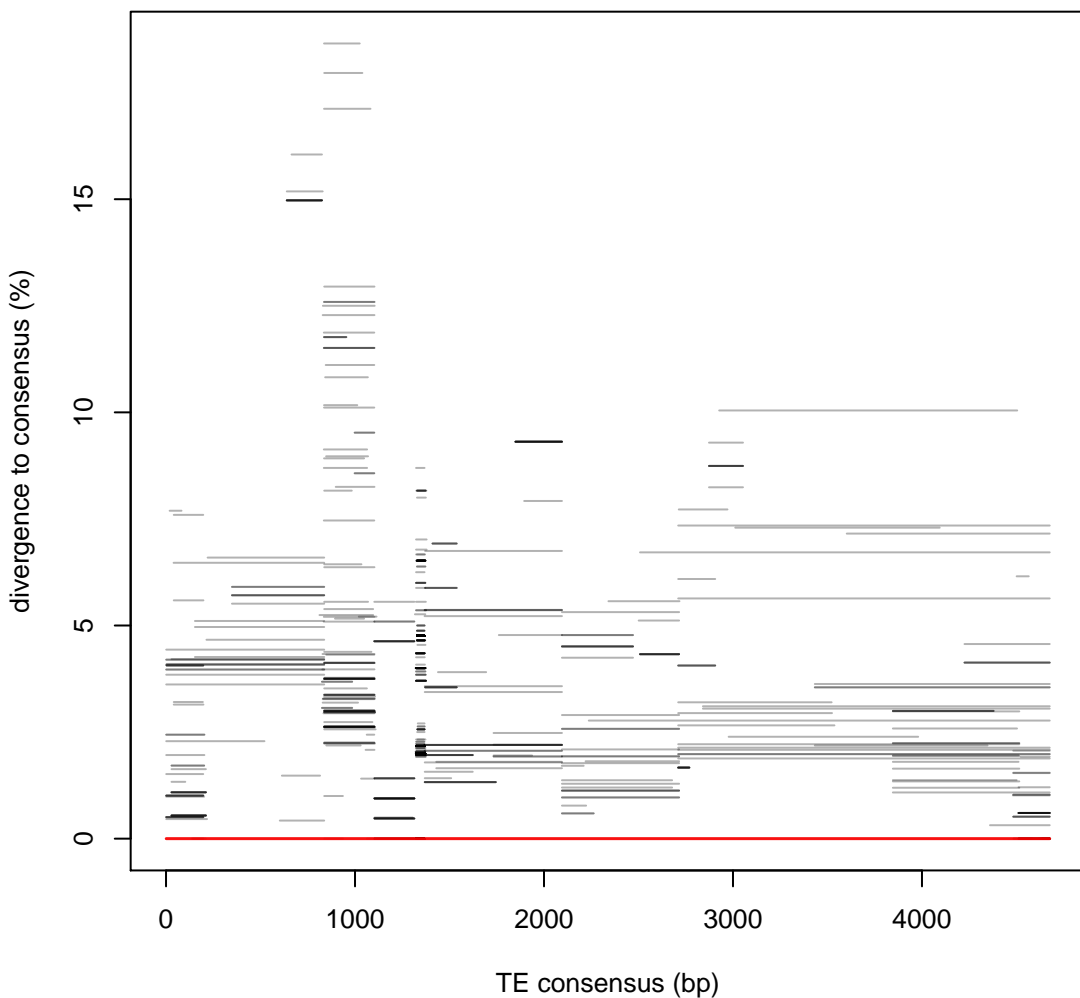
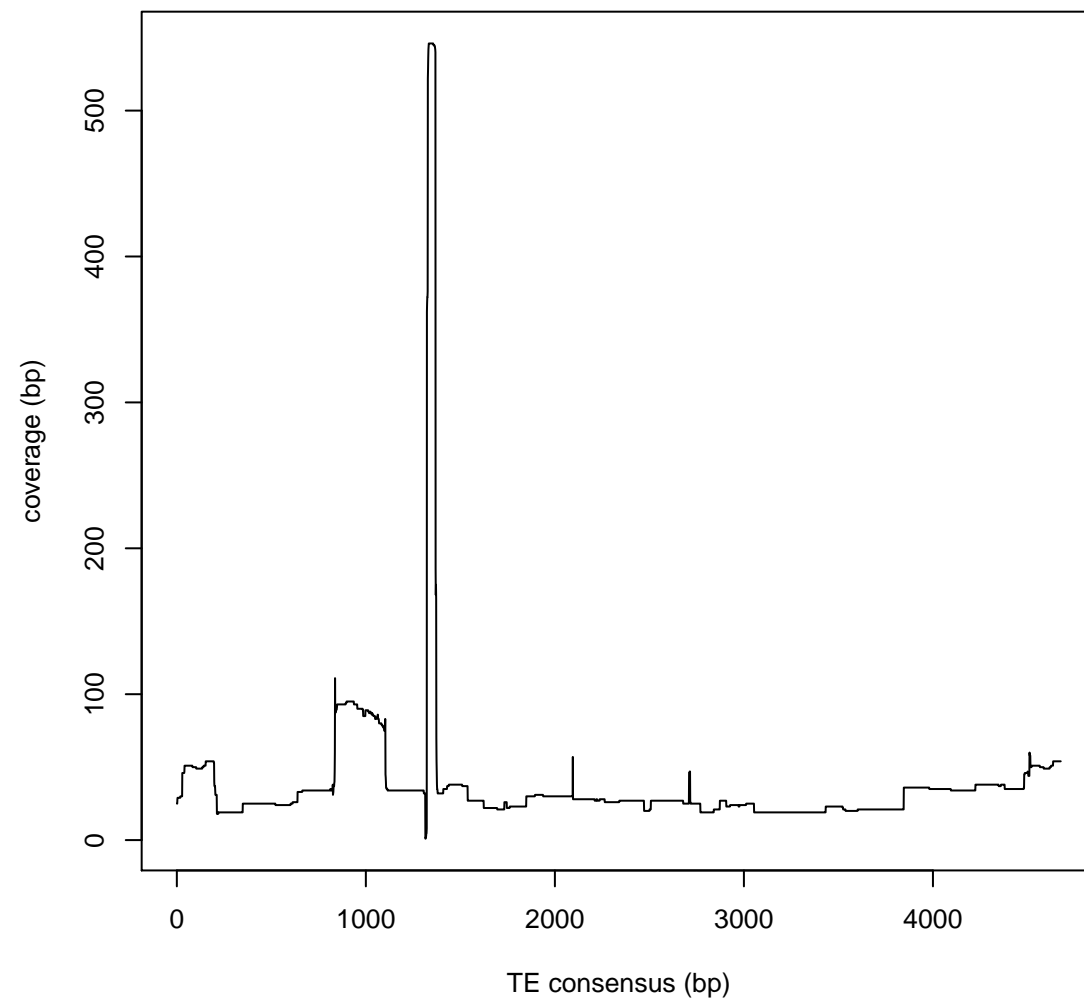


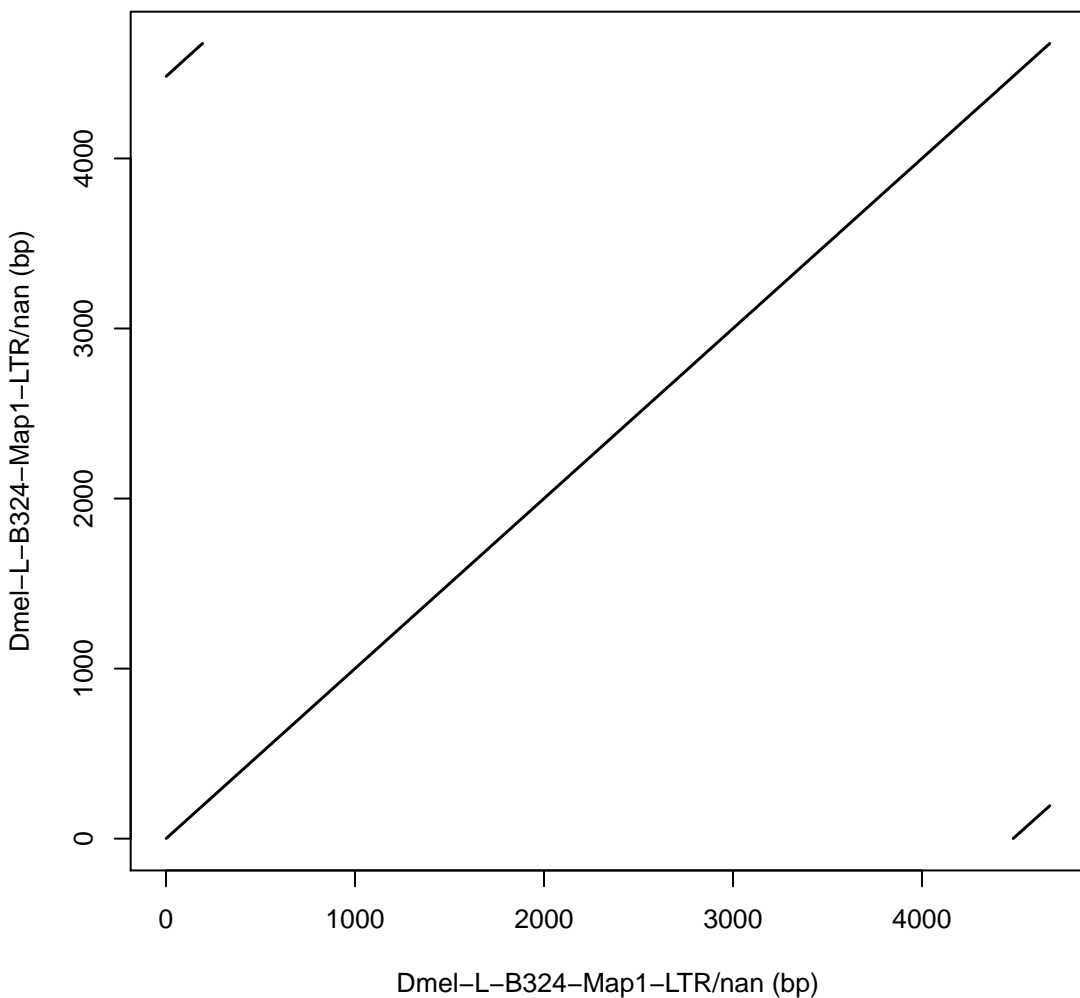
TE: Dmel-L-B324-Map1-LTR/nan
consensus size: 4676bp; fragments: 935; full length: 1 (>=4208.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

