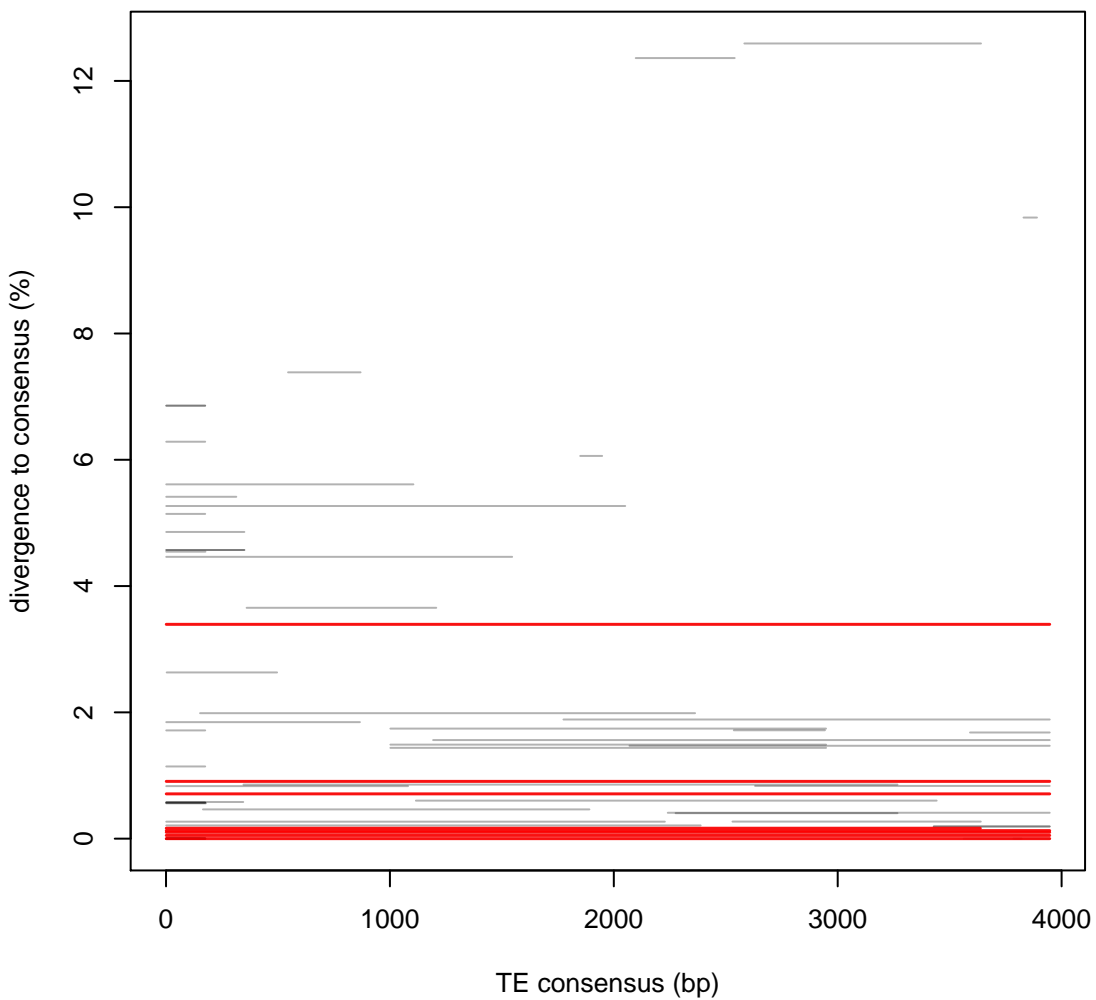
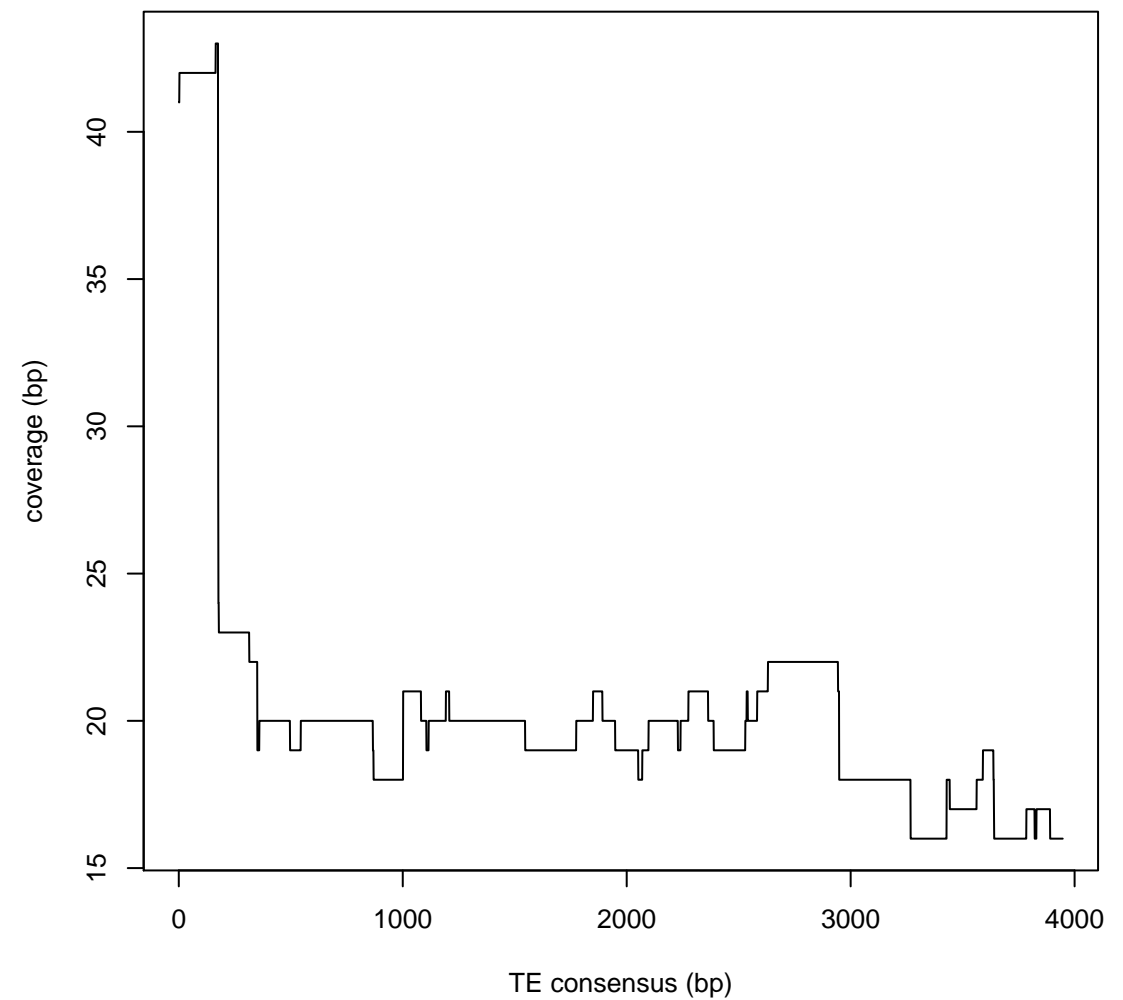


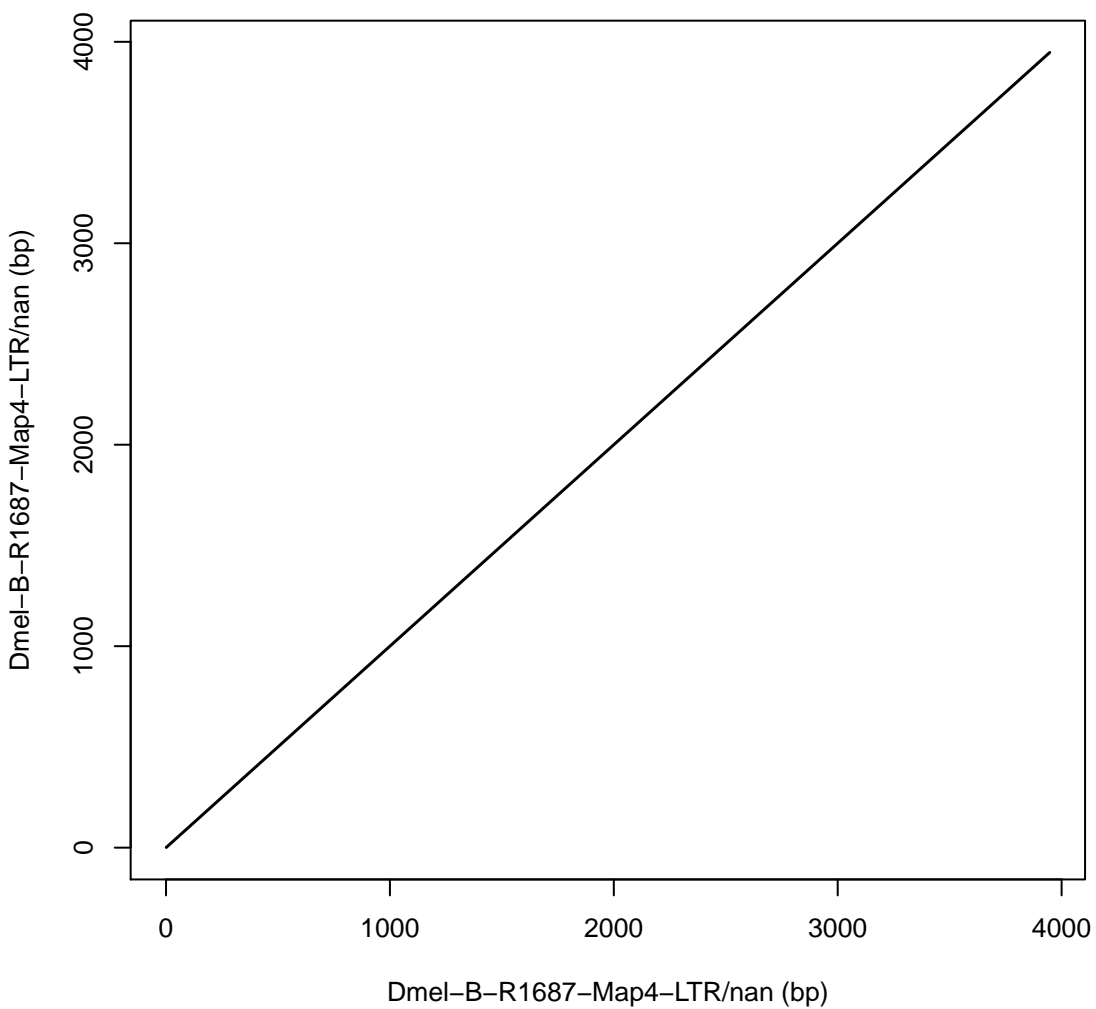
TE: Dmel-B-R1687-Map4-LTR/nan  
consensus size: 3947bp; fragments: 68; full length: 8 (>=3552.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

