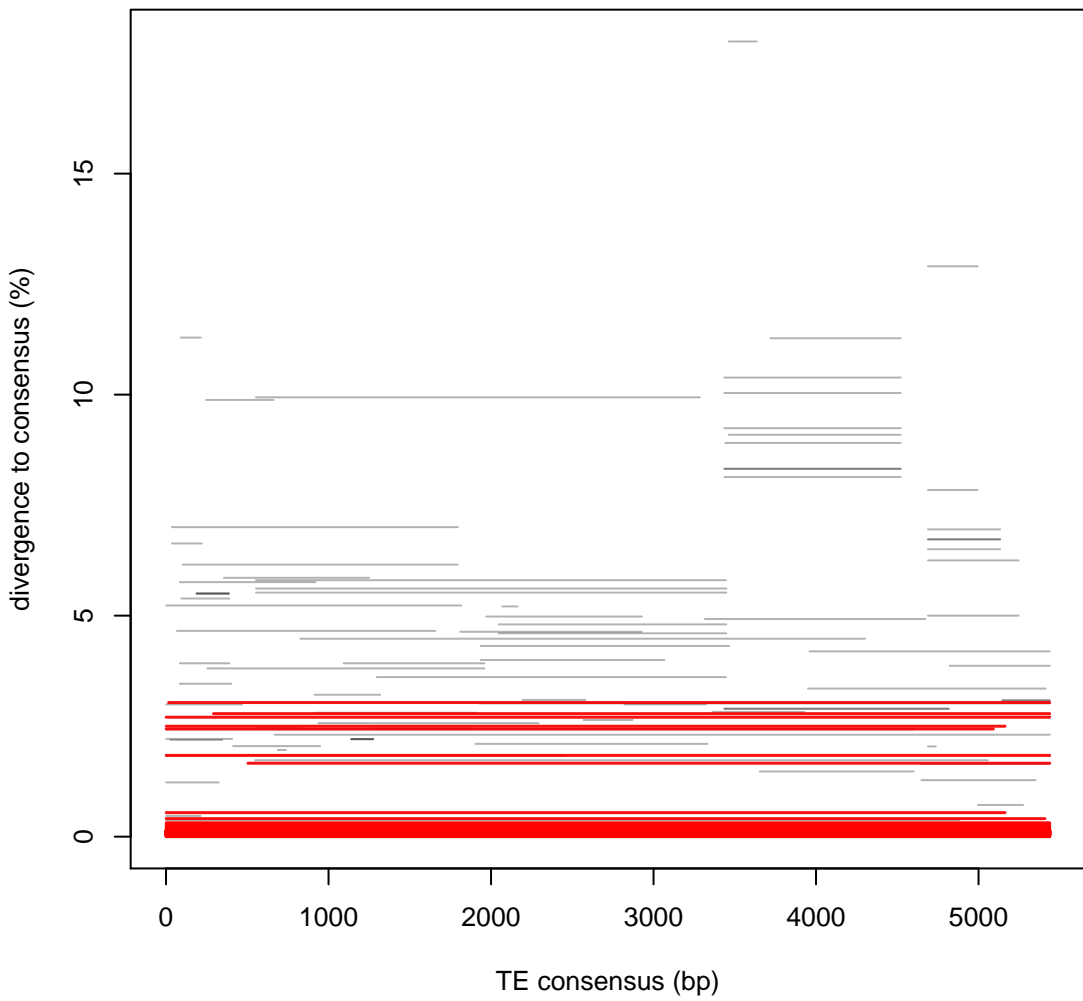
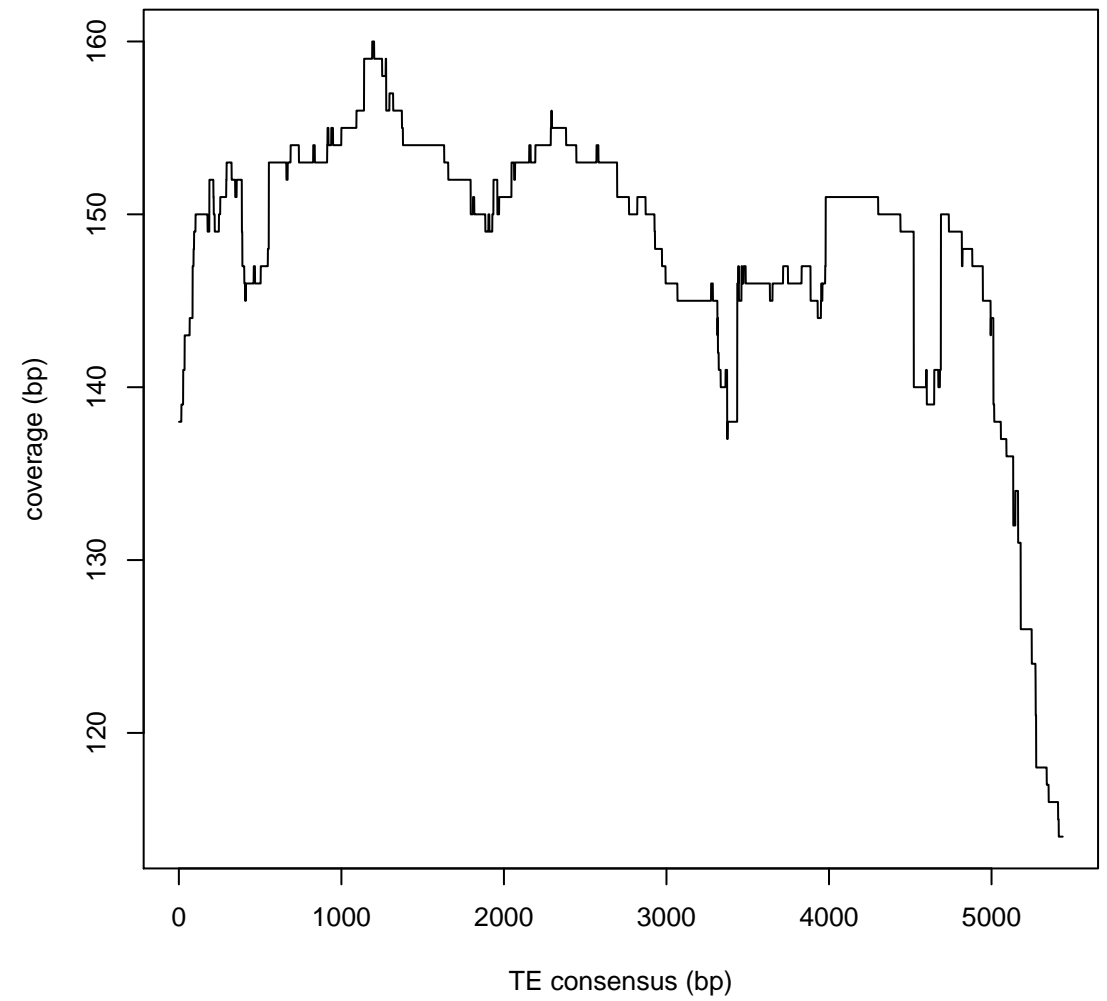


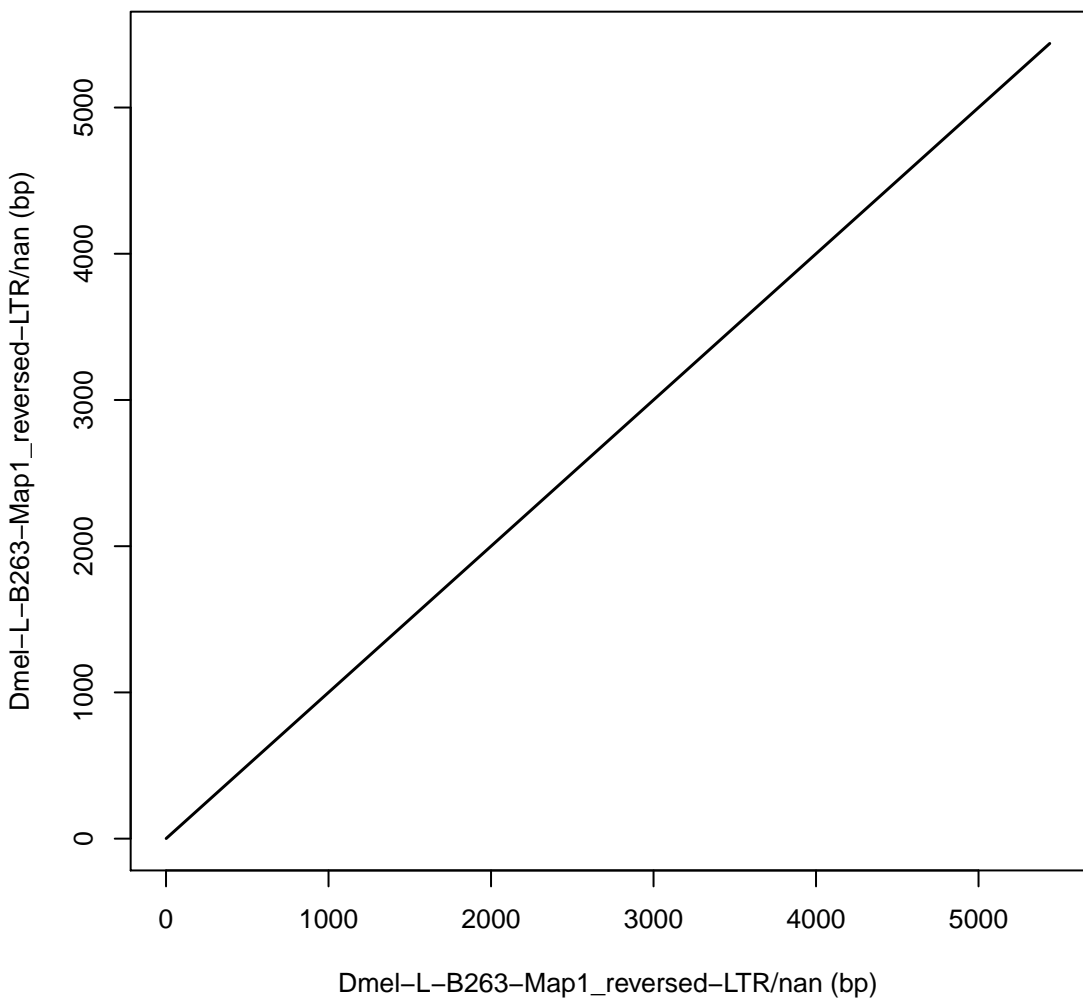
TE: Dmel-L-B263-Map1_reversed-LTR/nan
consensus size: 5438bp; fragments: 244; full length: 111 (>=4894.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

