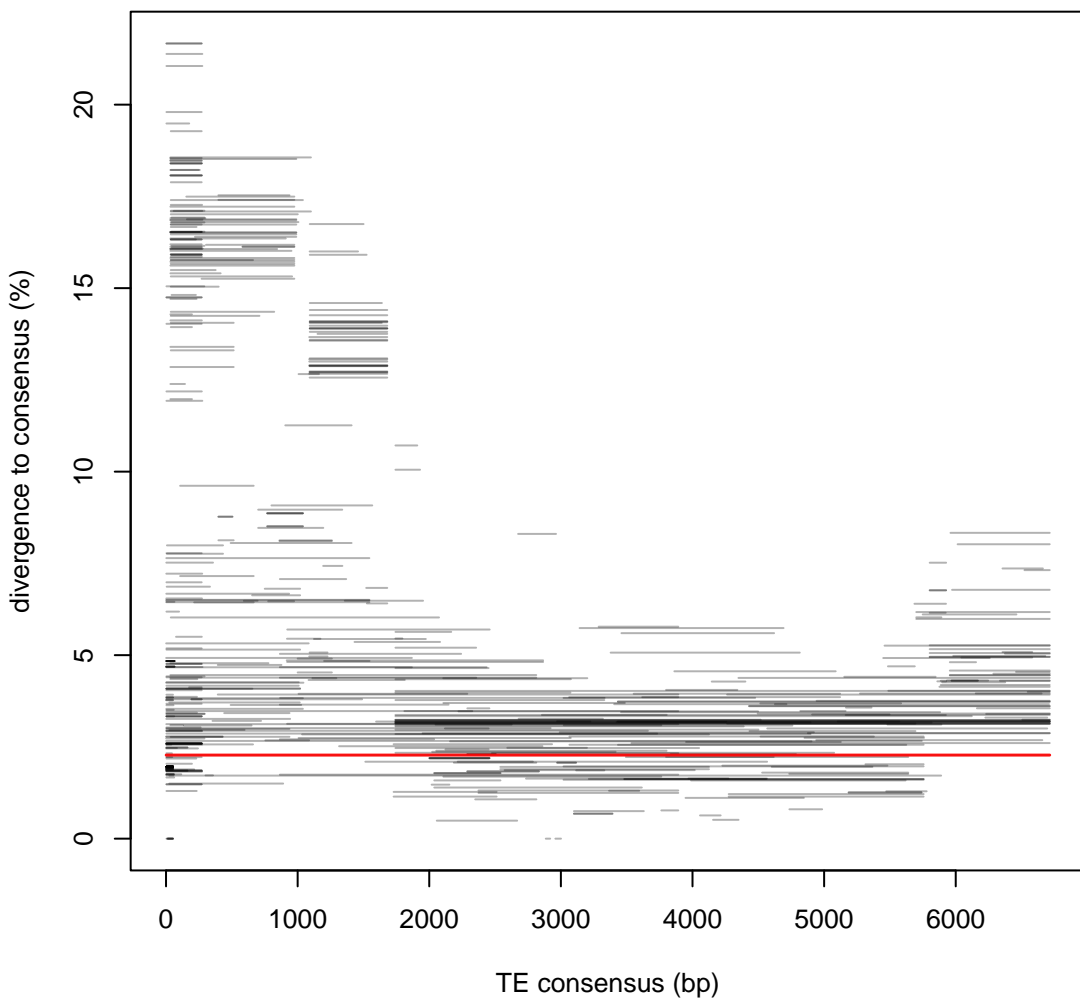
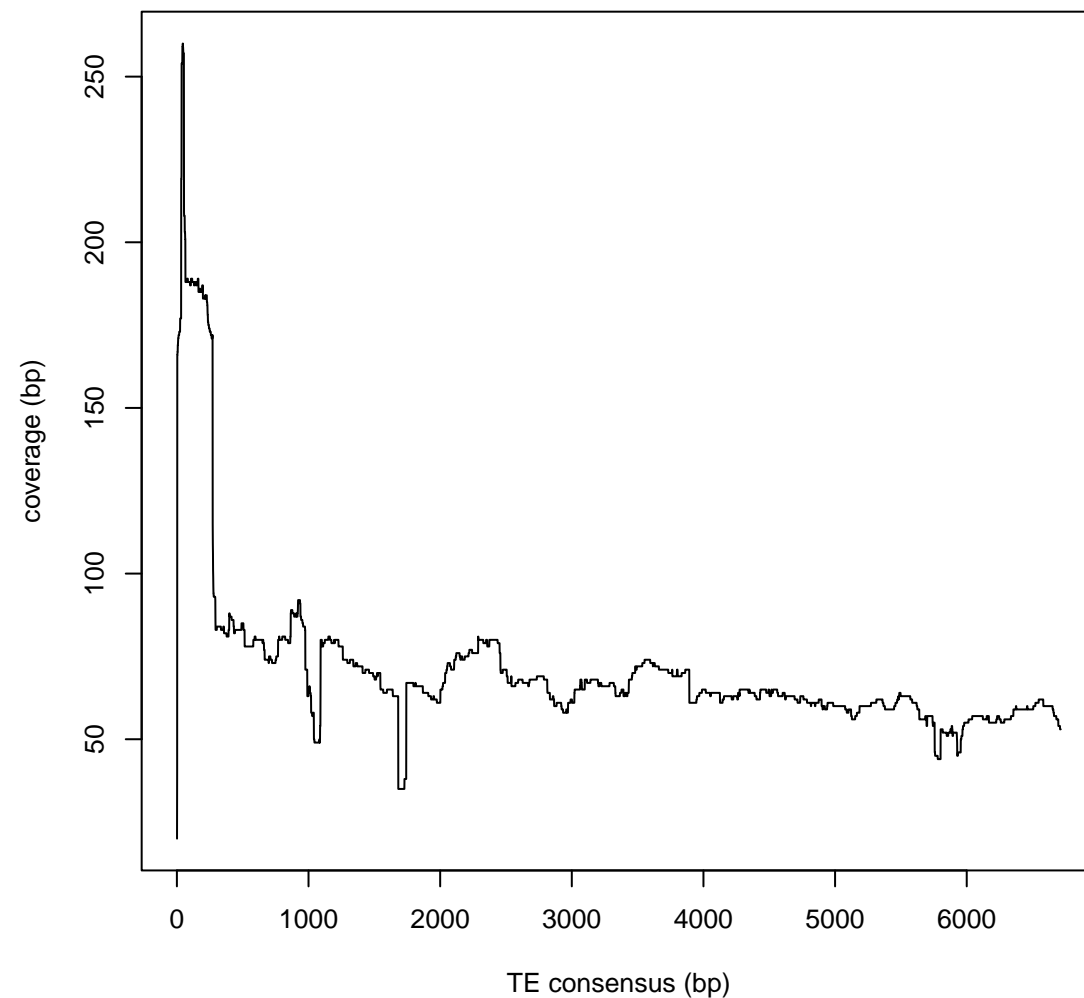


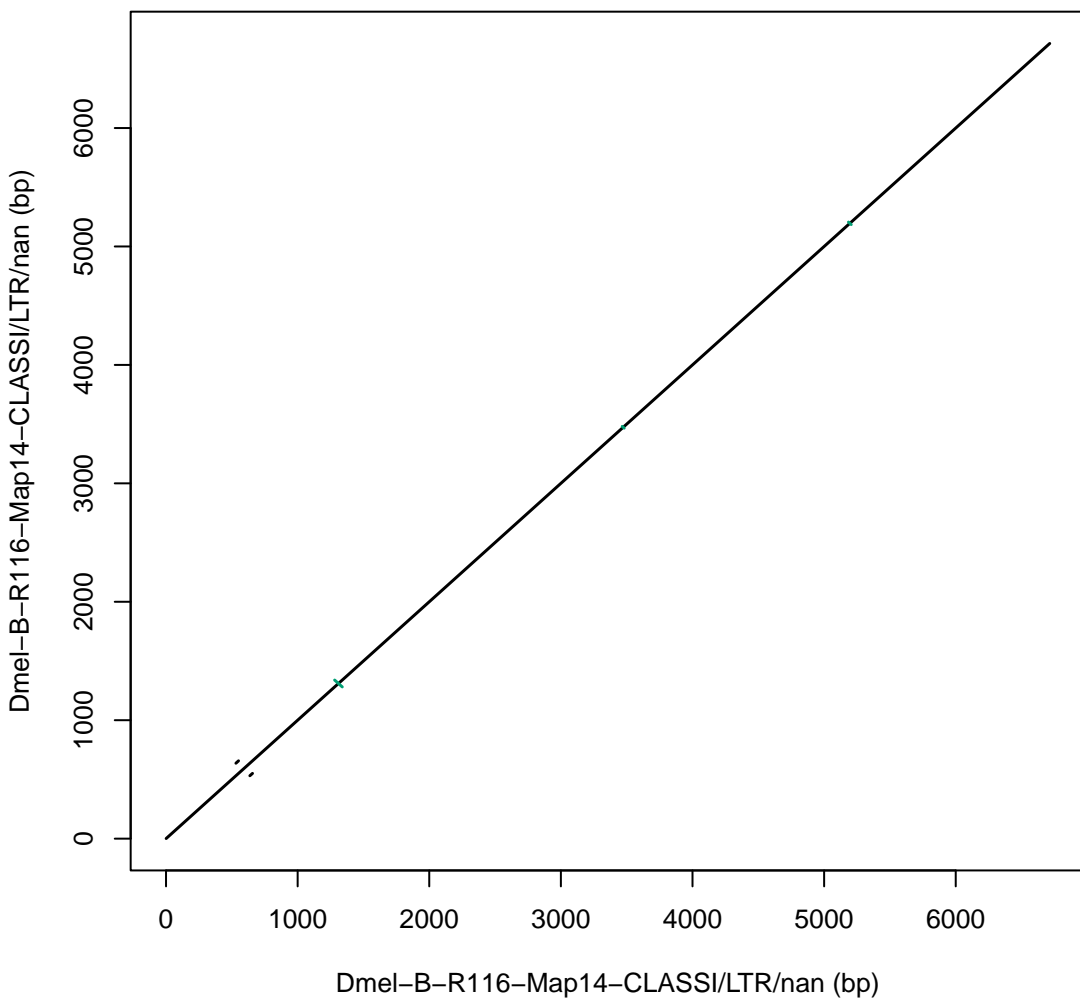
TE: Dmel-B-R116-Map14-CLASSI/LTR/nan  
consensus size: 6714bp; fragments: 572; full length: 1 (>=6042.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

