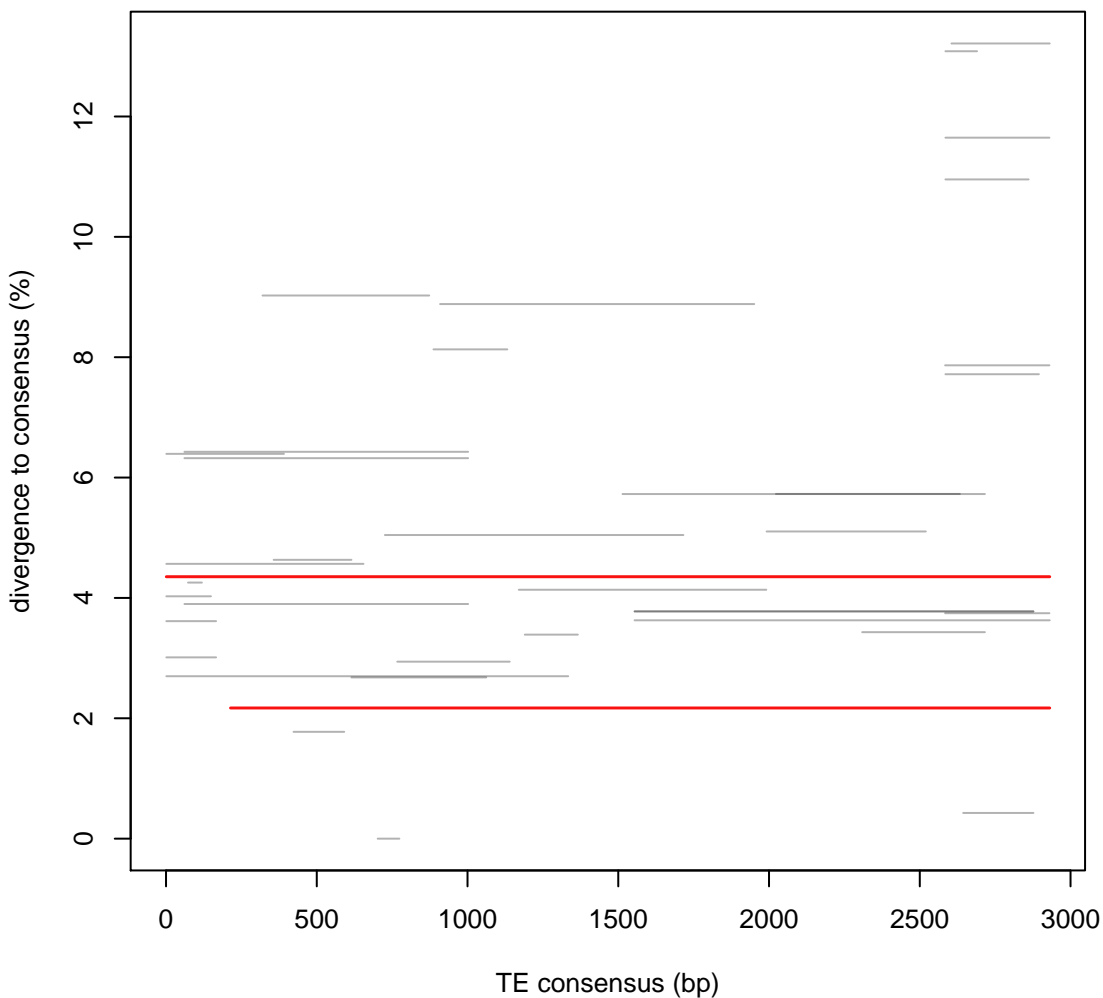
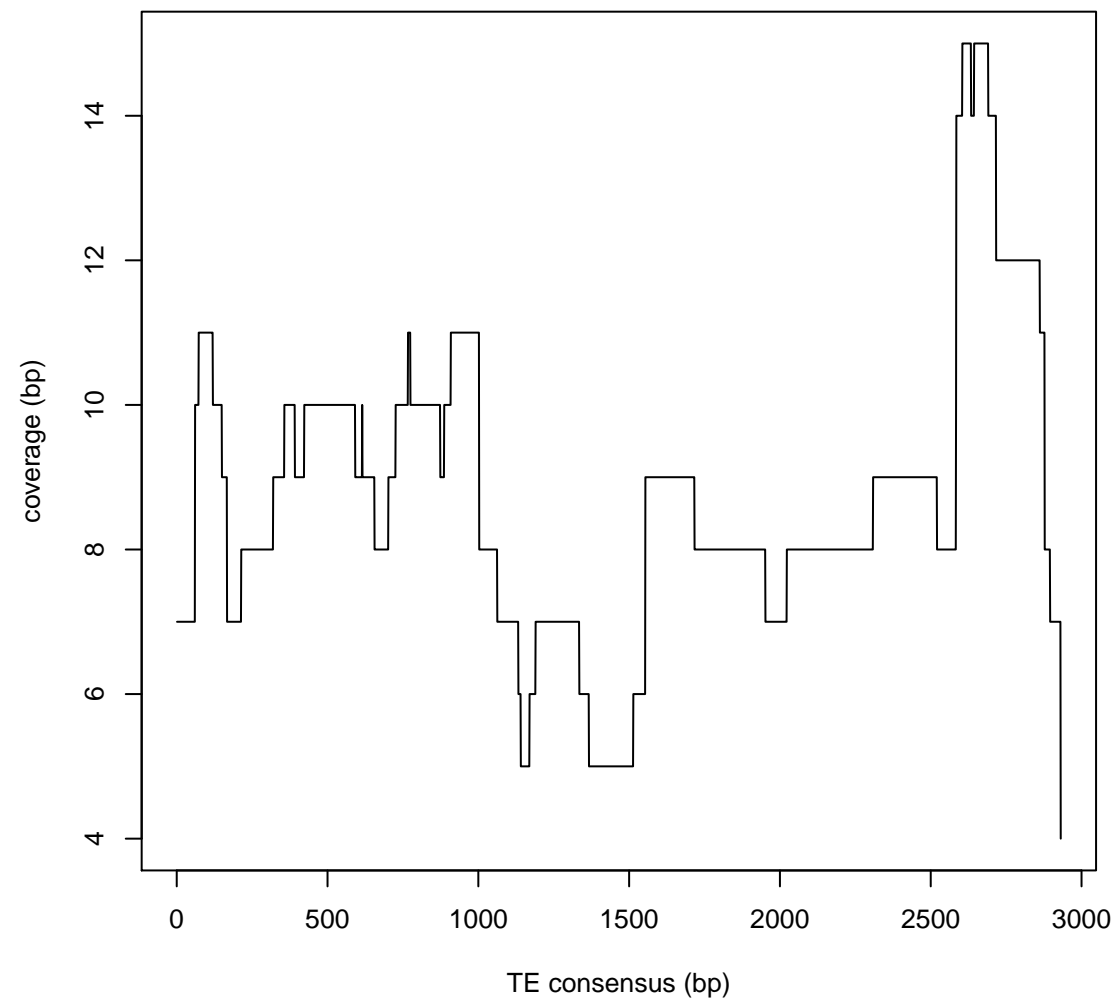


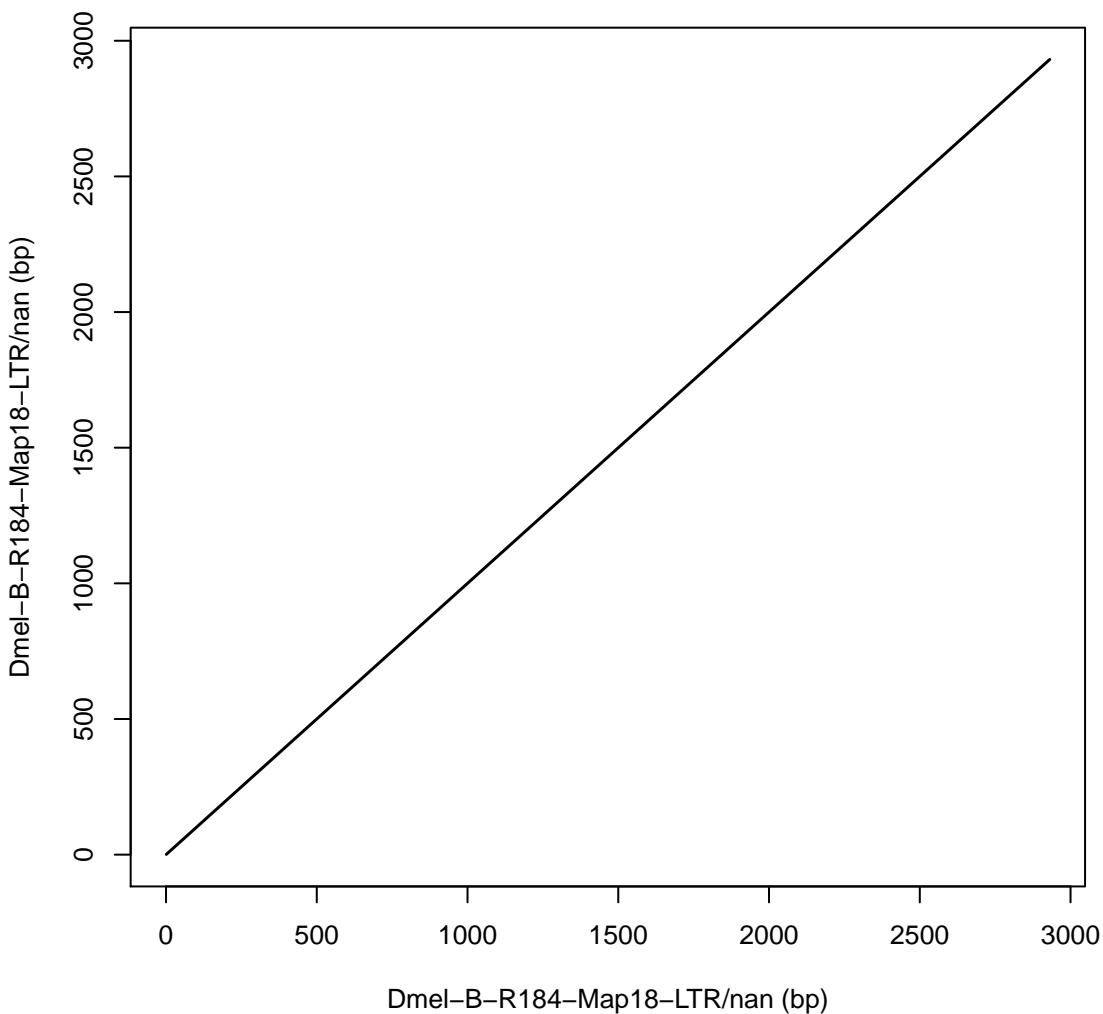
TE: Dmel-B-R184-Map18-LTR/nan  
consensus size: 2931bp; fragments: 38; full length: 2 (>=2637.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

