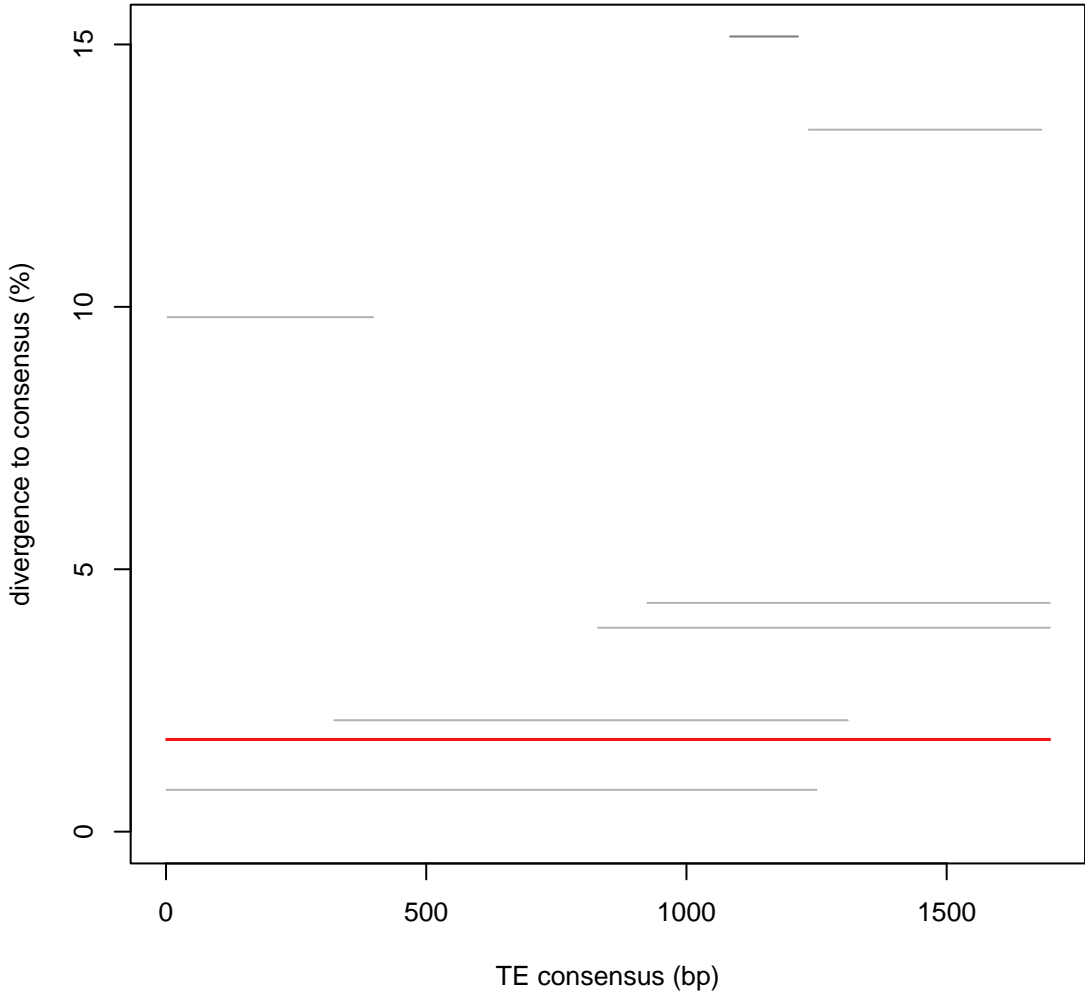
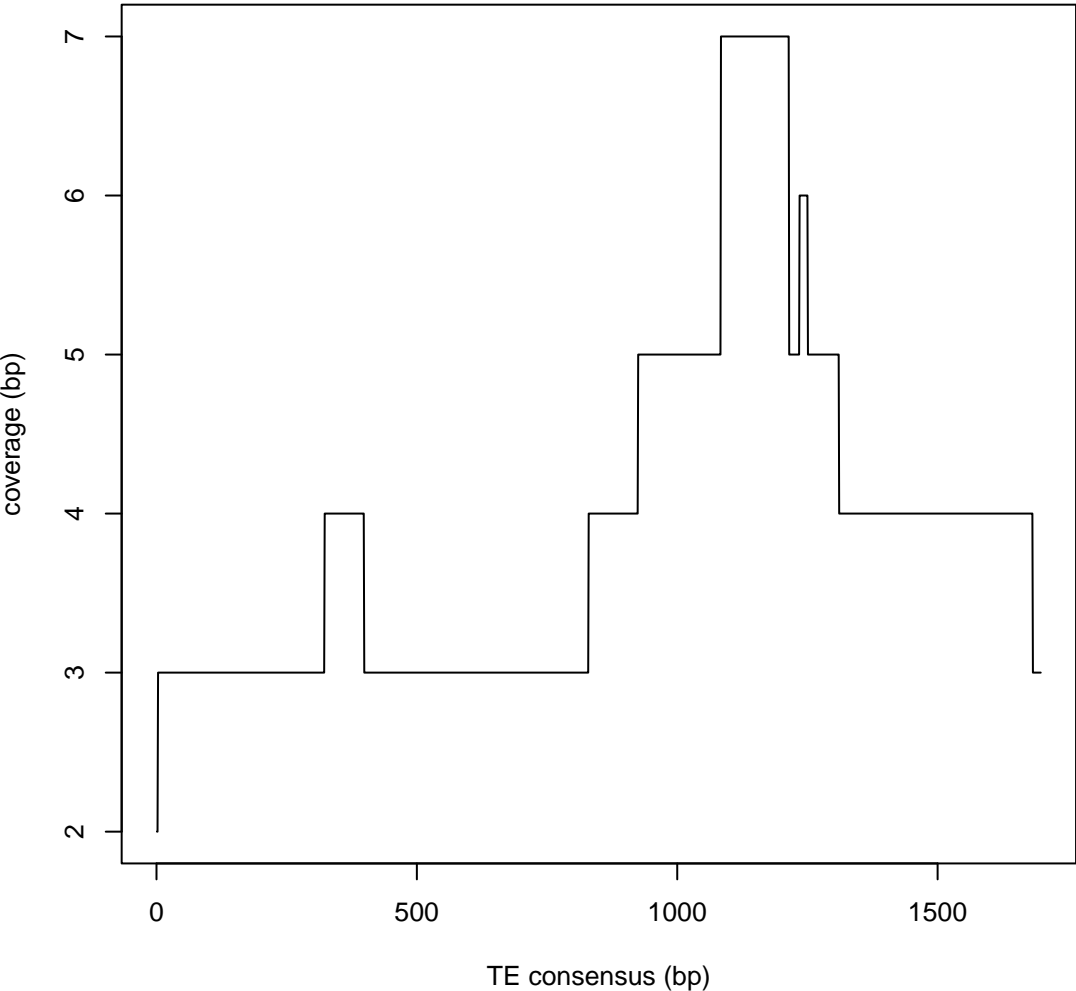


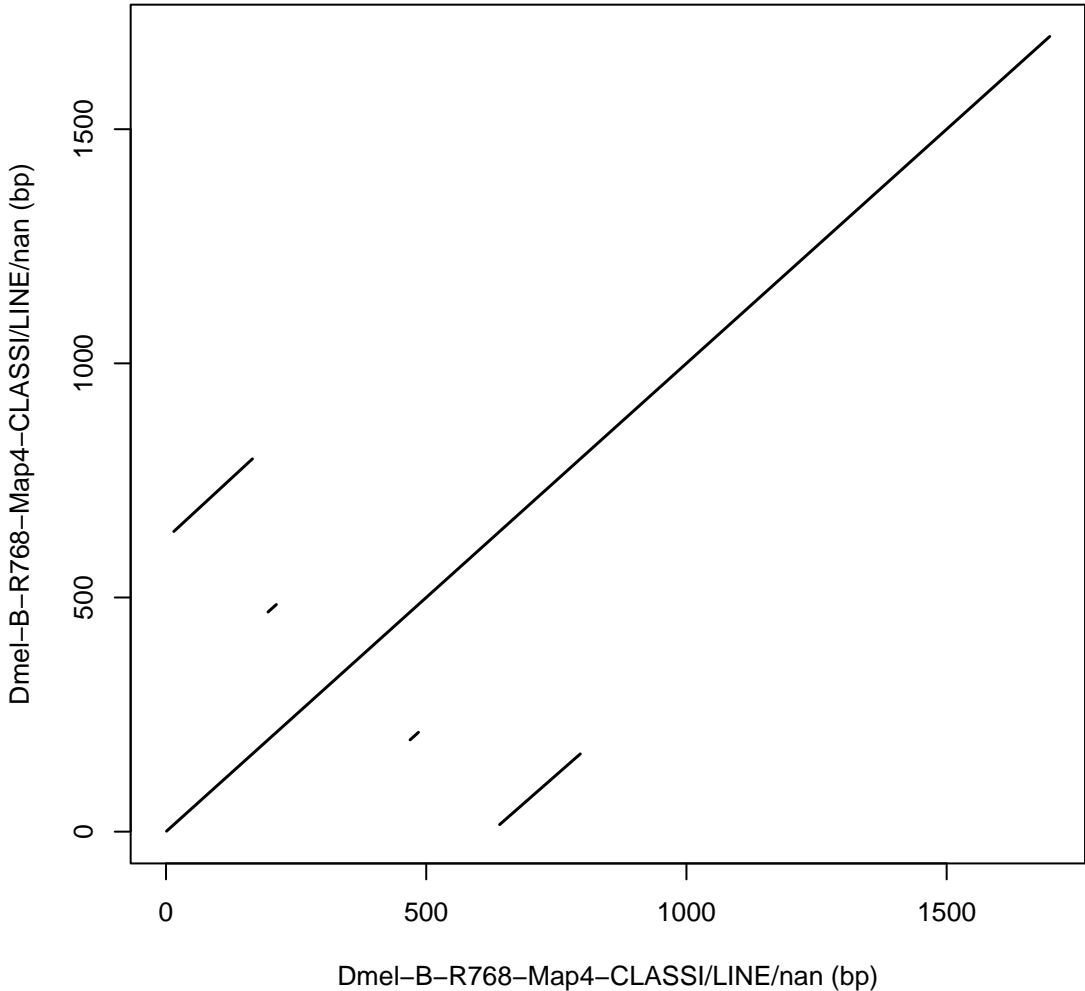
TE: Dmel-B-R768-Map4-CLASSI/LINE/nan  
consensus size: 1698bp; fragments: 9; full length: 1 (>=1528.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

