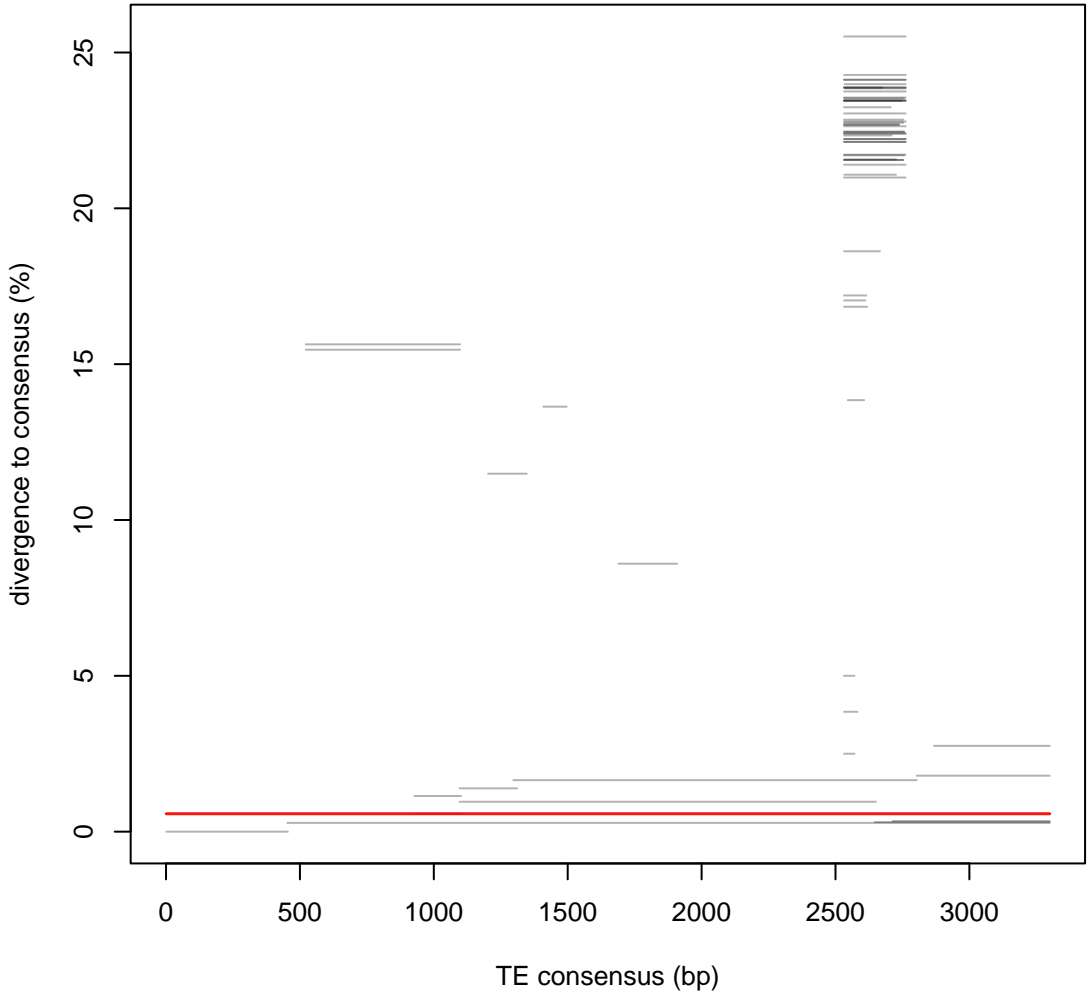
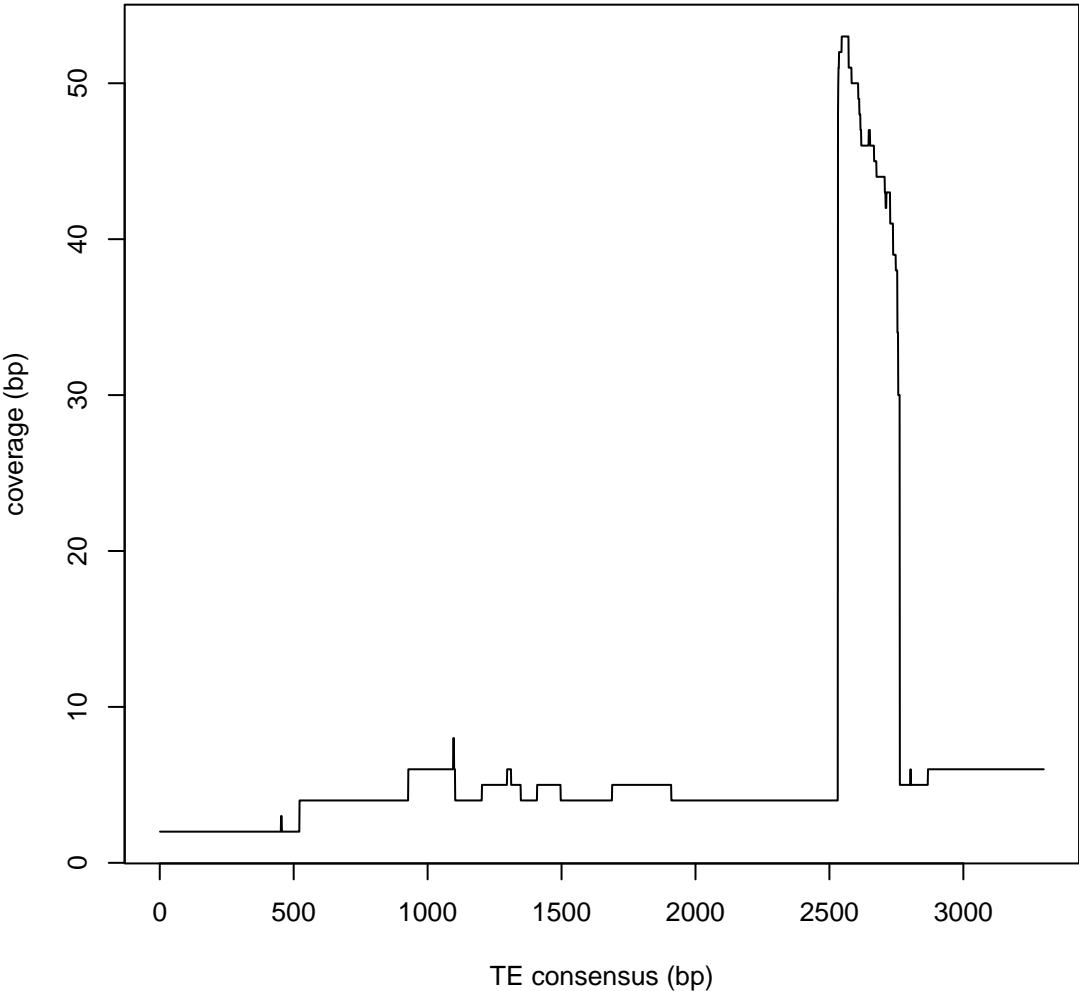


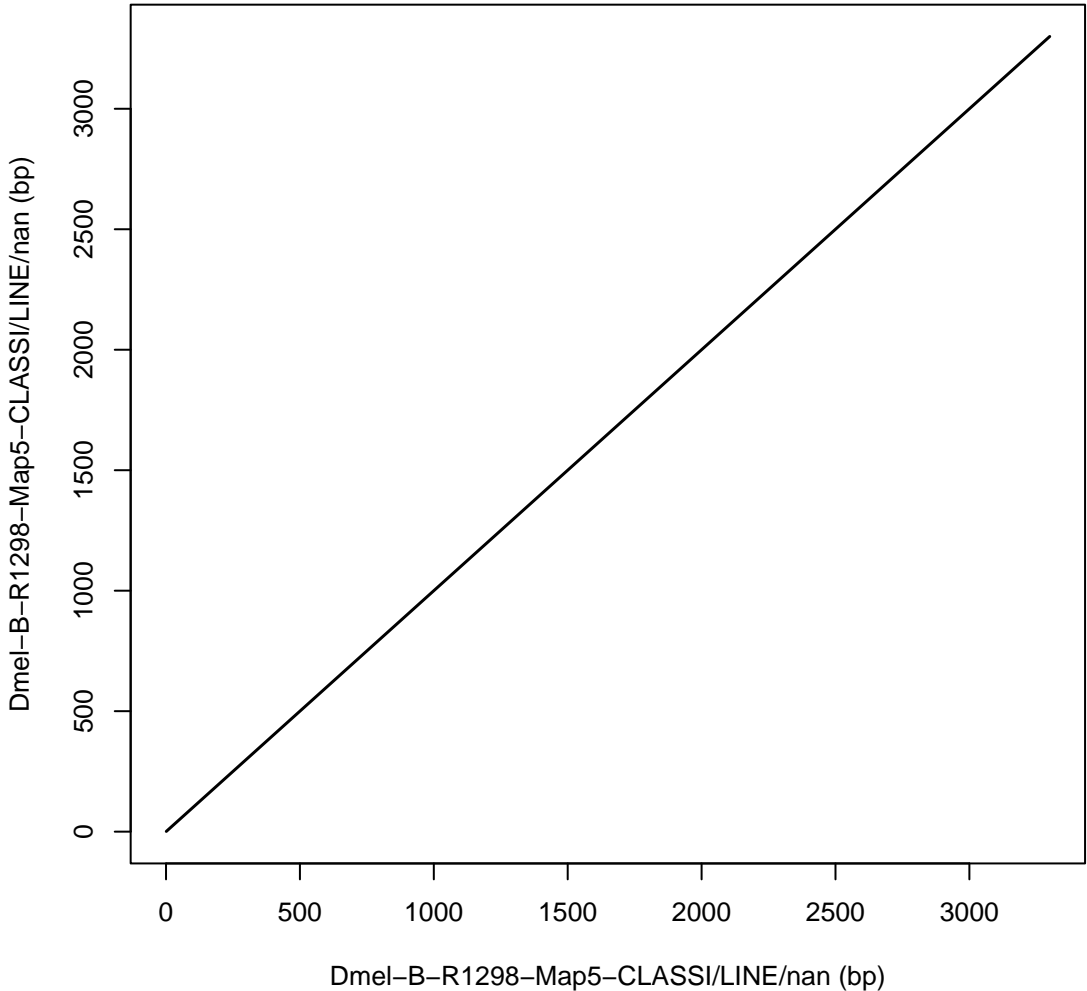
TE: Dmel-B-R1298-Map5-CLASSI/LINE/nan  
consensus size: 3300bp; fragments: 66; full length: 1 (>=2970bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

