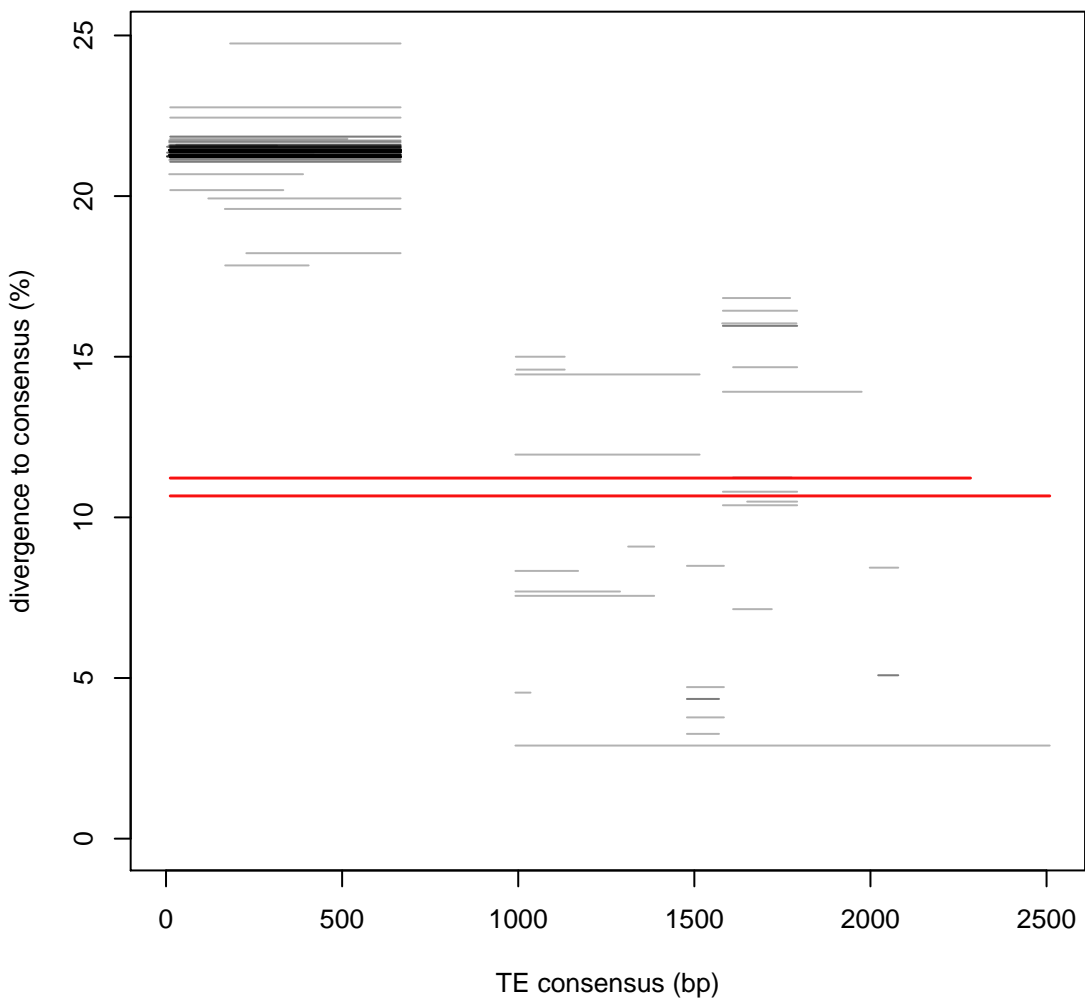
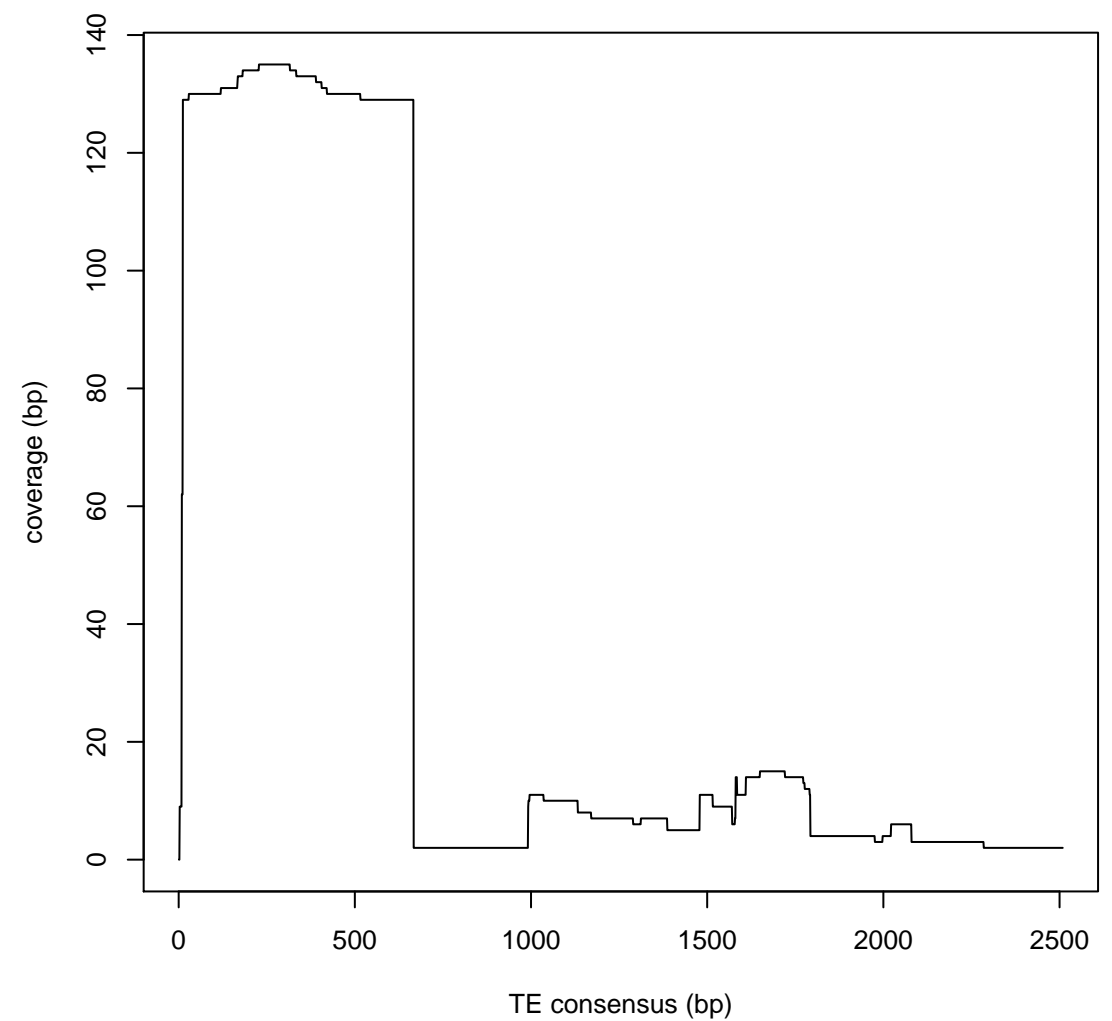


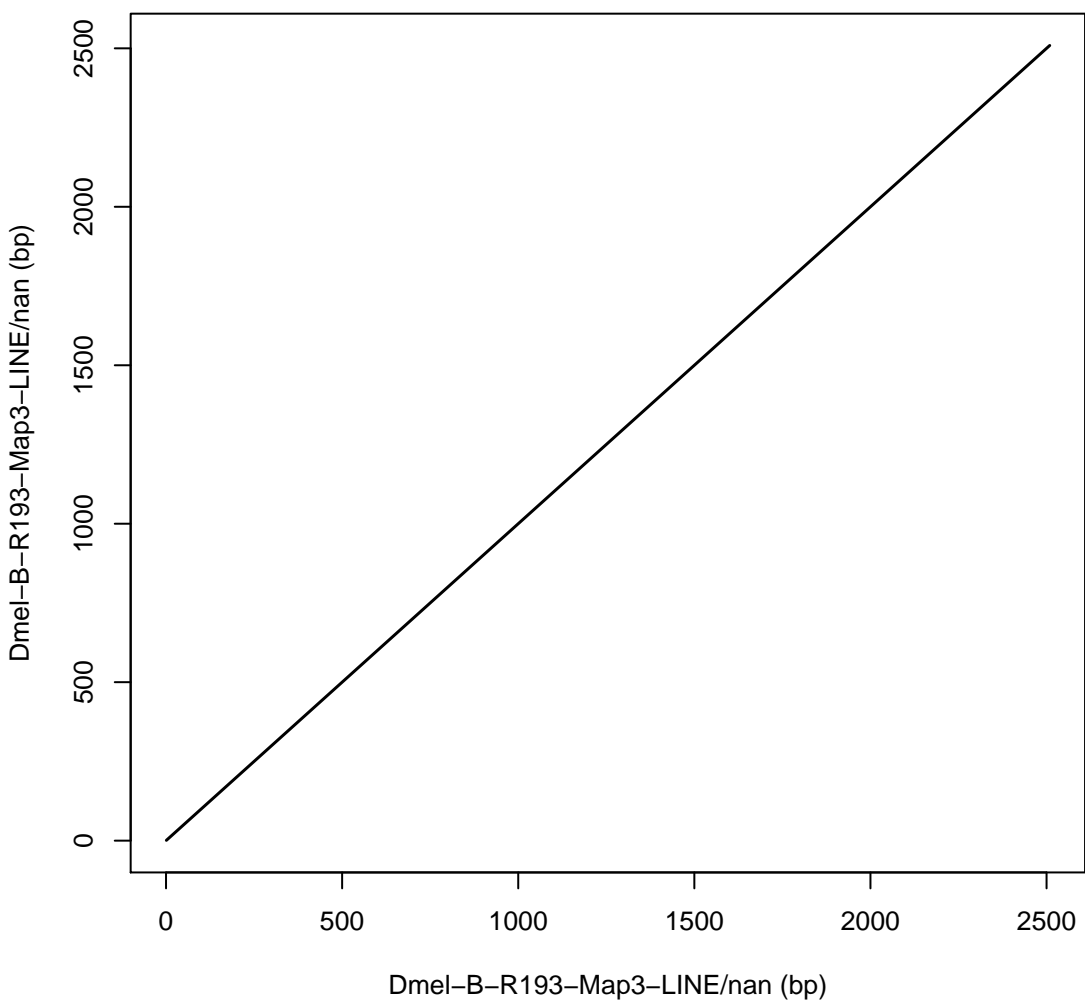
TE: Dmel-B-R193-Map3-LINE/nan  
consensus size: 2509bp; fragments: 166; full length: 2 (>=2258.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

