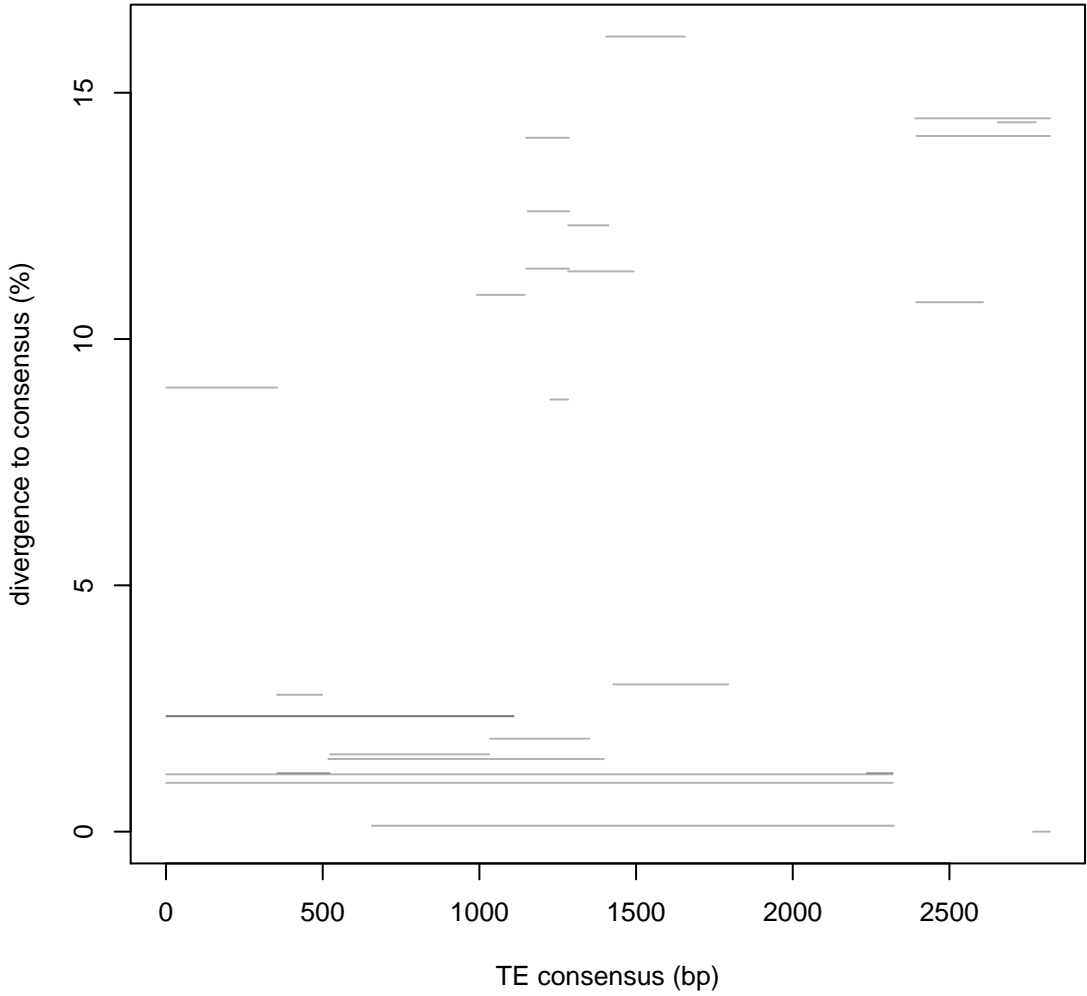
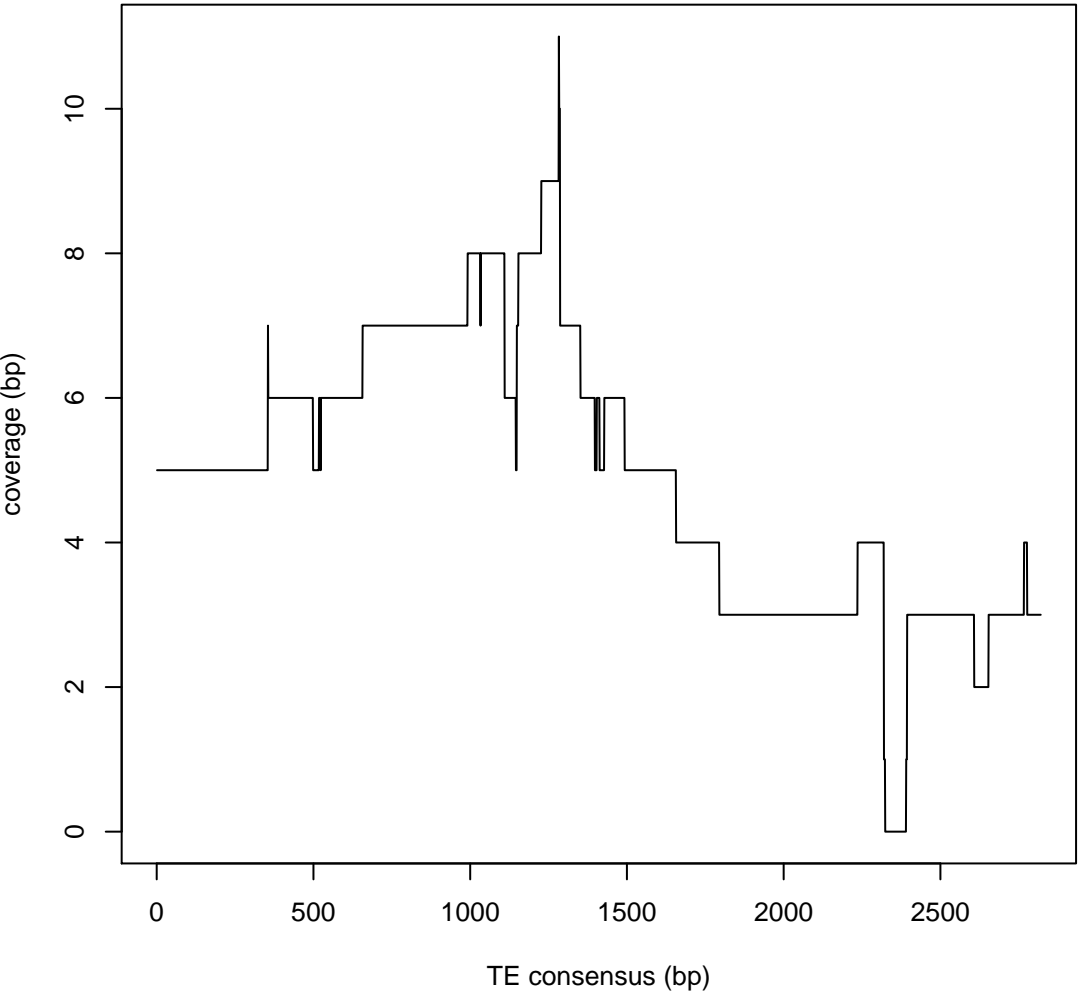


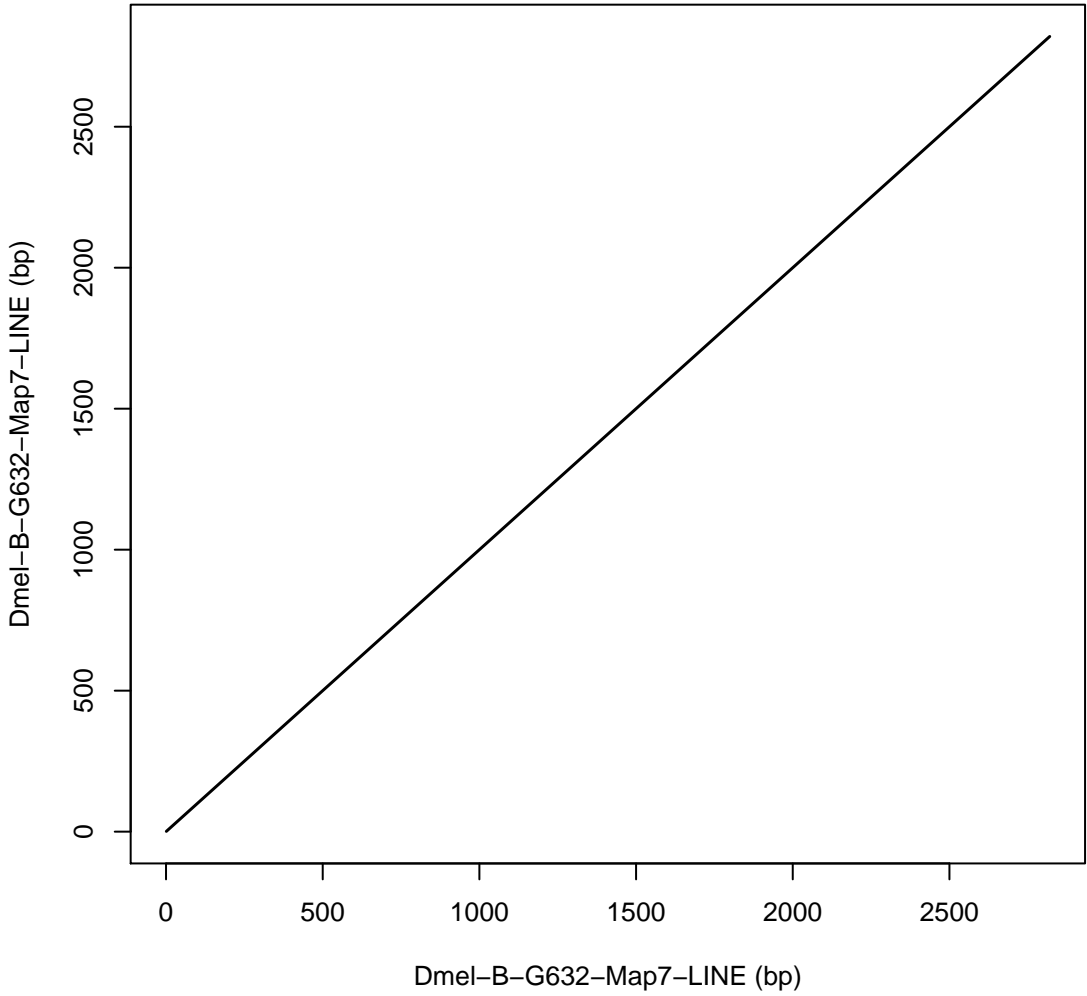
TE: Dmel-B-G632-Map7-LINE  
consensus size: 2820bp; fragments: 26; full length: 0 (>=2538bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

