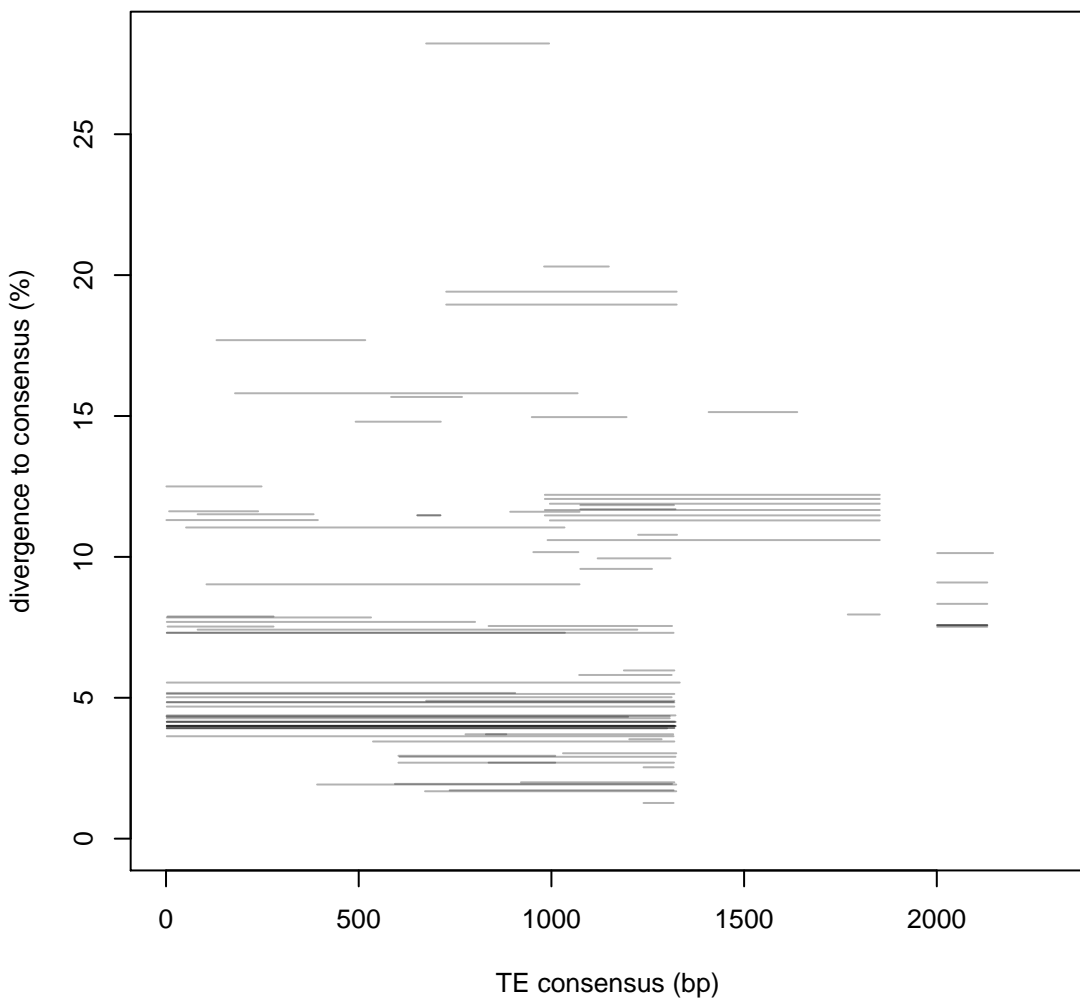
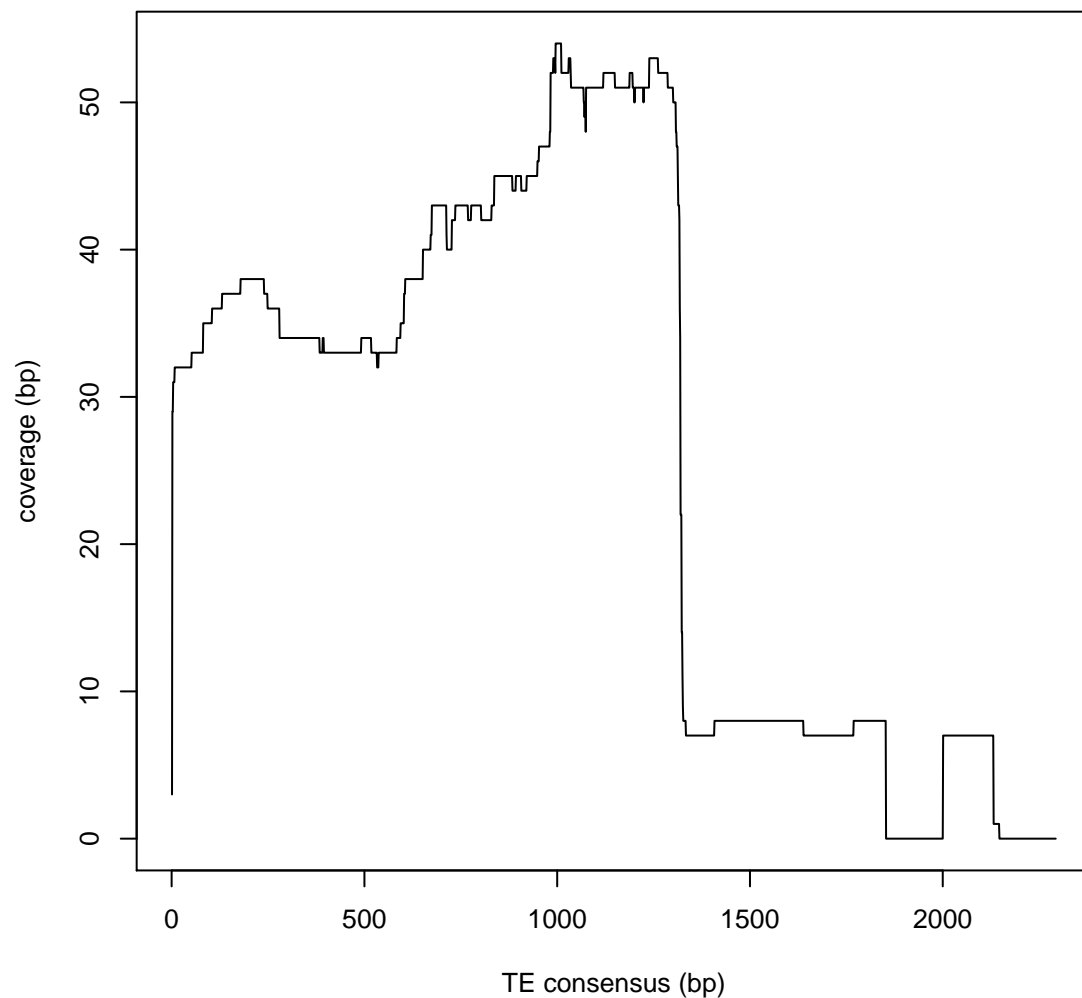


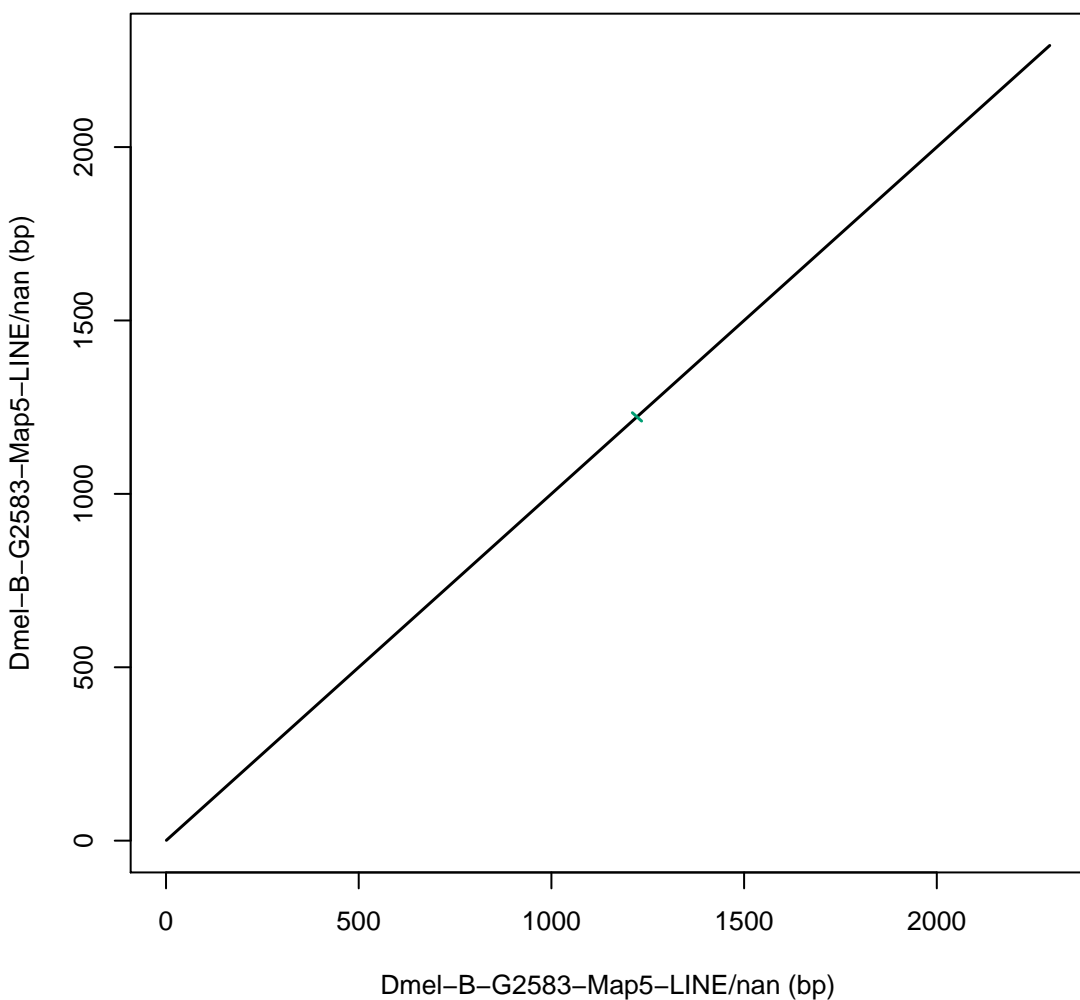
TE: Dmel-B-G2583-Map5-LINE/nan  
consensus size: 2293bp; fragments: 90; full length: 0 (>=2063.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

