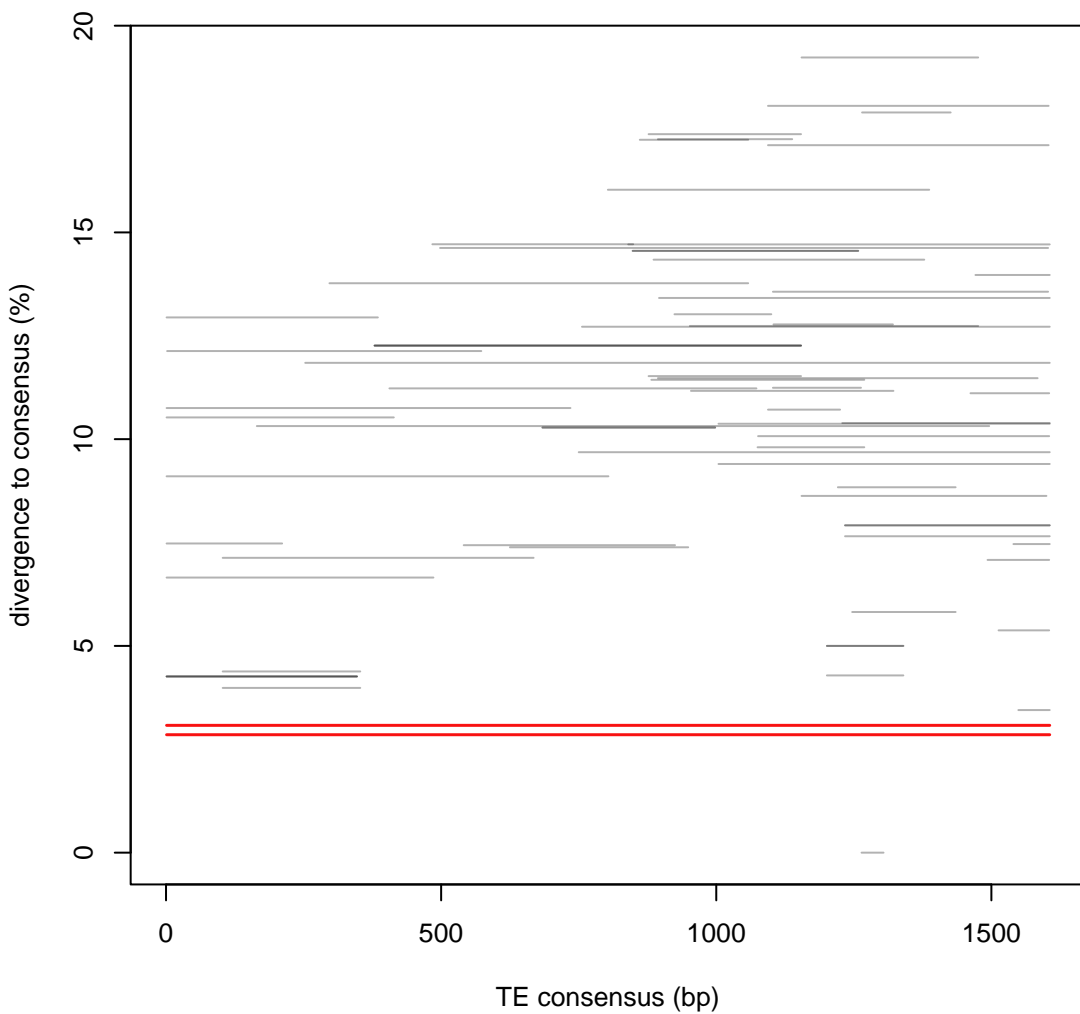
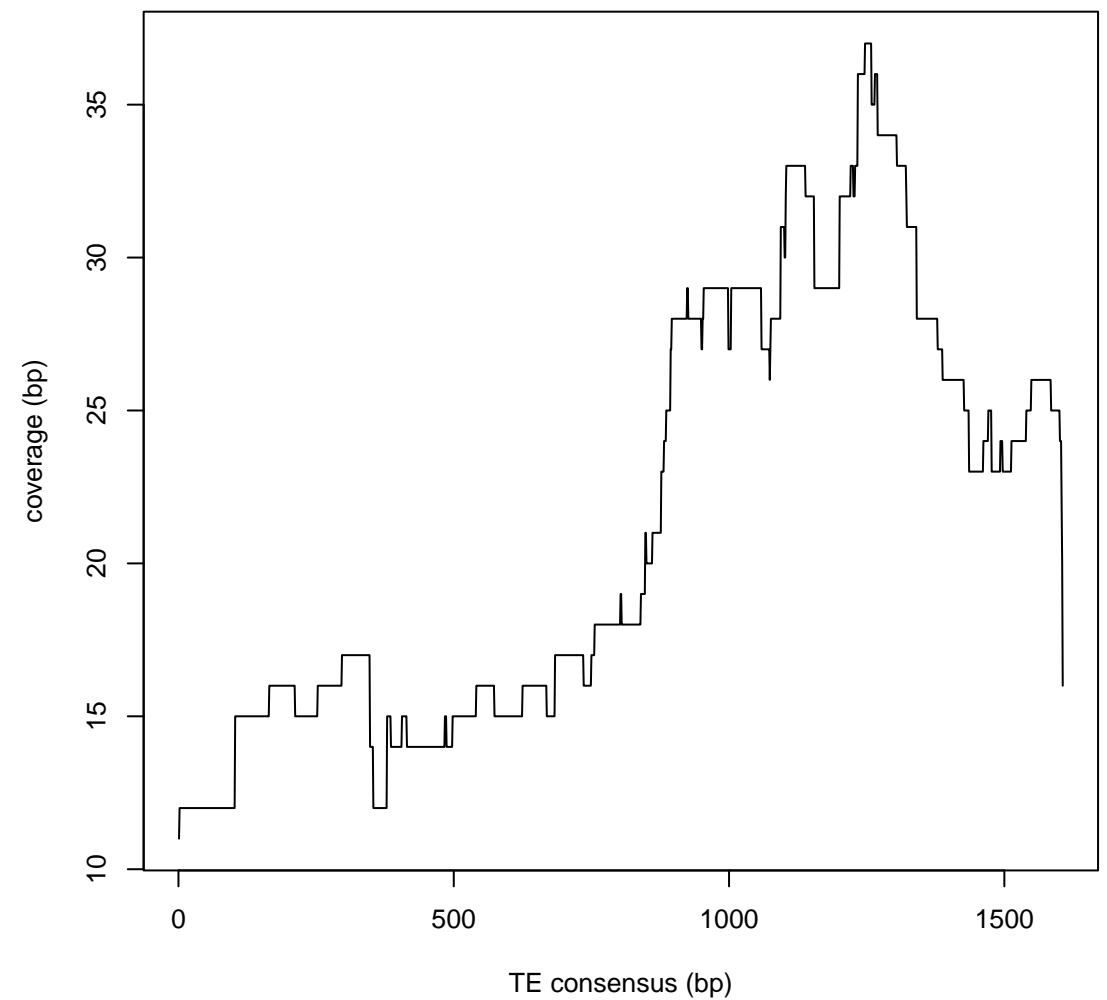


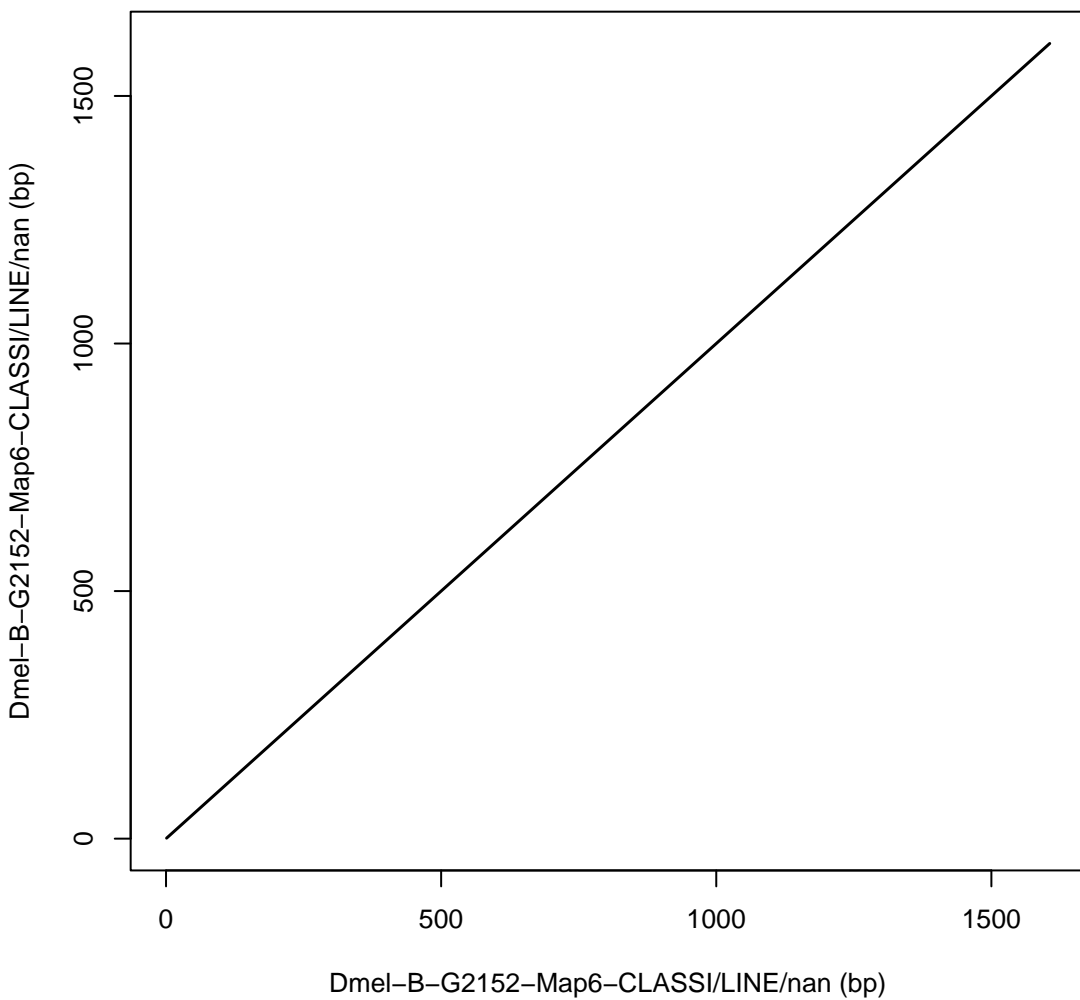
TE: Dmel-B-G2152-Map6-CLASSI/LINE/nan  
consensus size: 1606bp; fragments: 74; full length: 2 ( $\geq 1445.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

