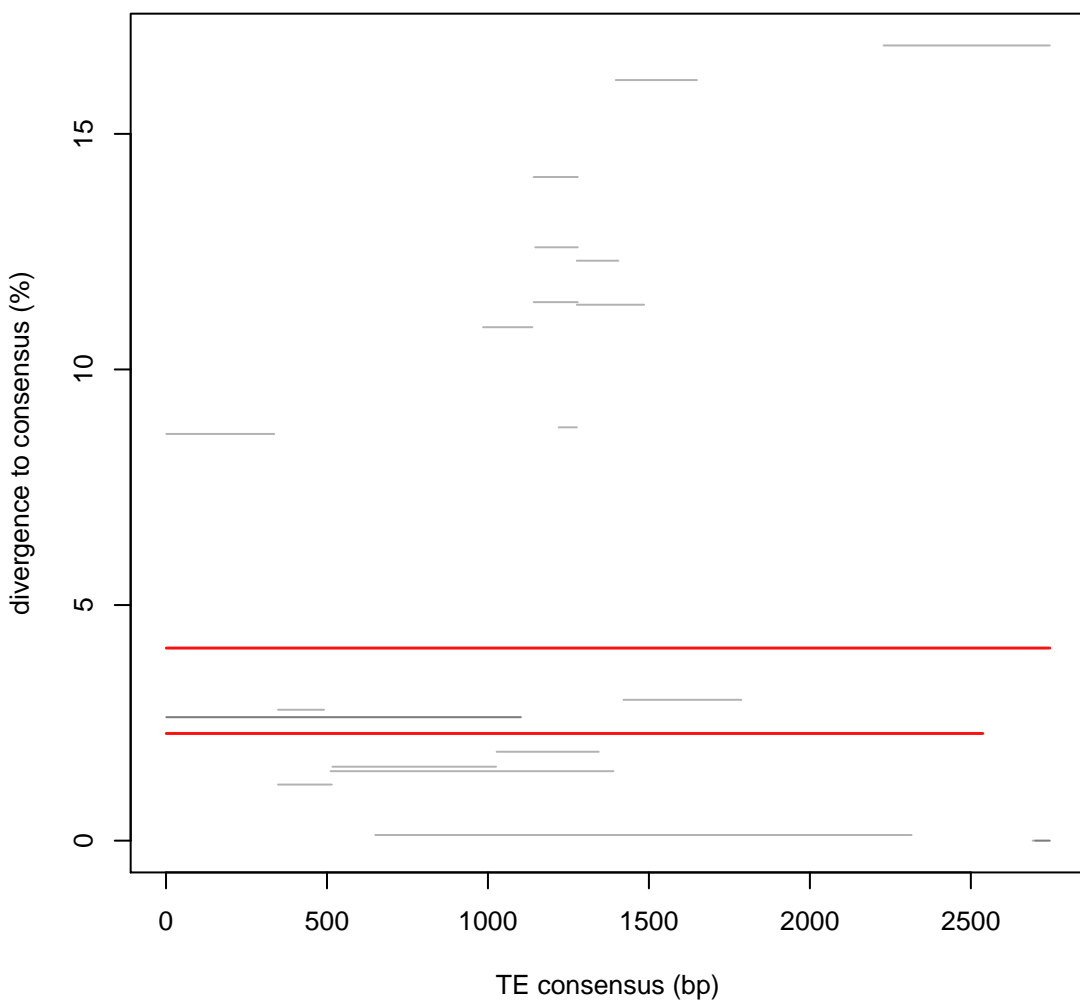
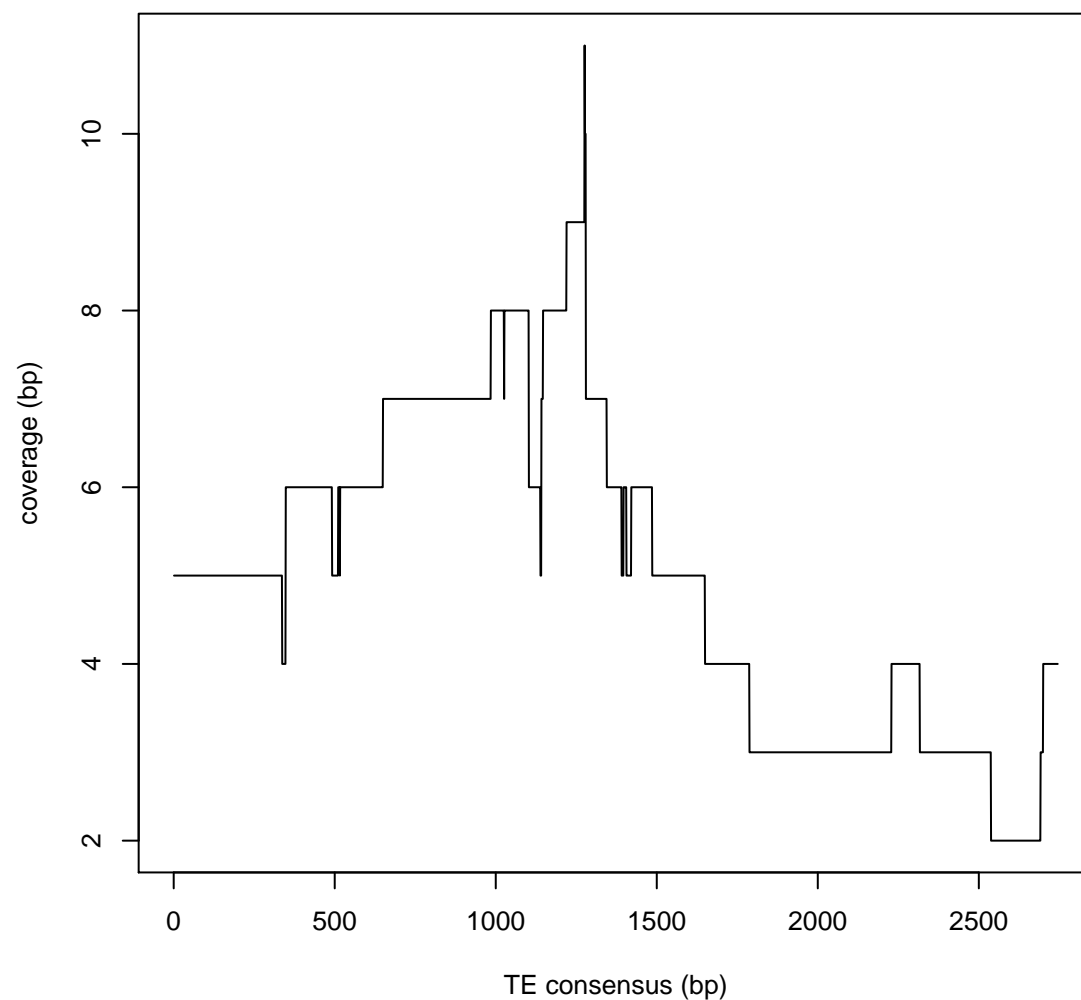


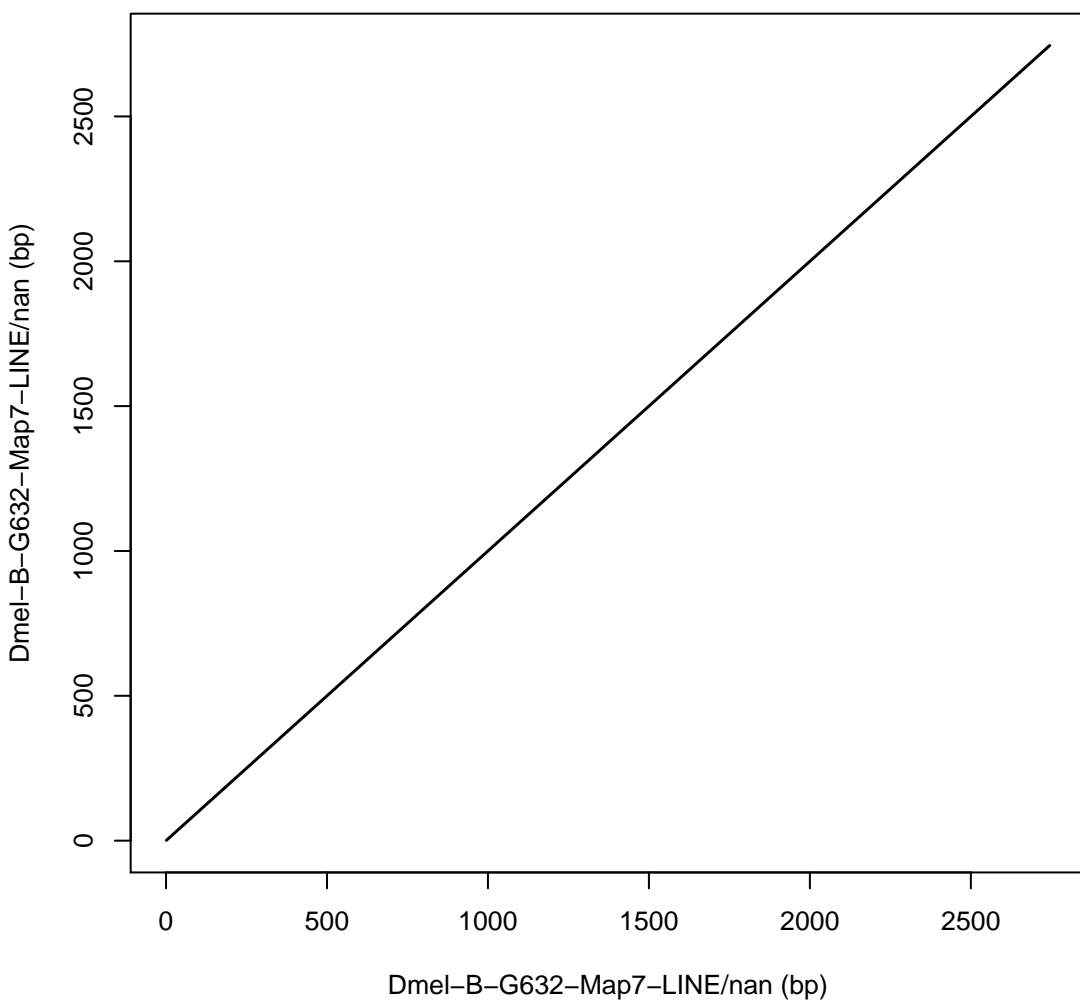
TE: Dmel-B-G632-Map7-LINE/nan  
consensus size: 2745bp; fragments: 23; full length: 2 ( $\geq 2470.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

