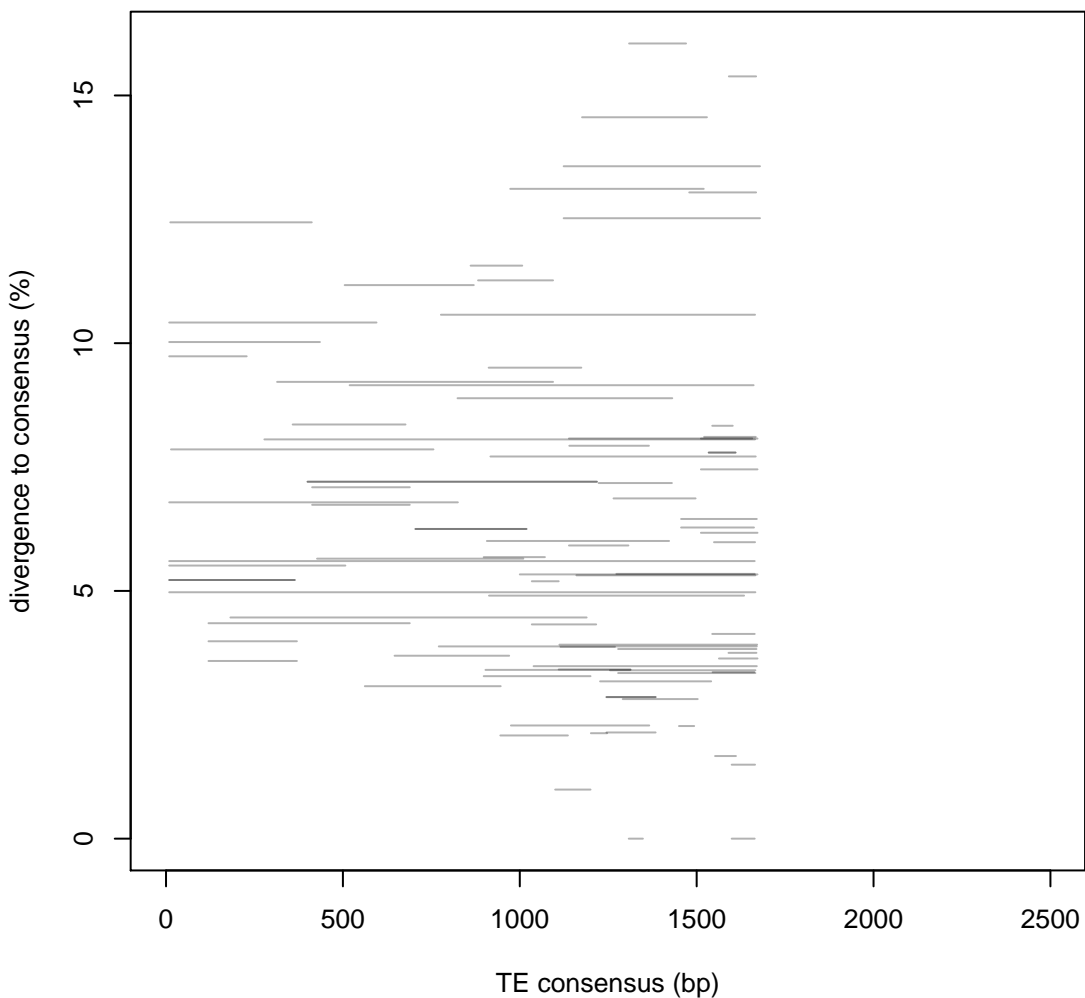
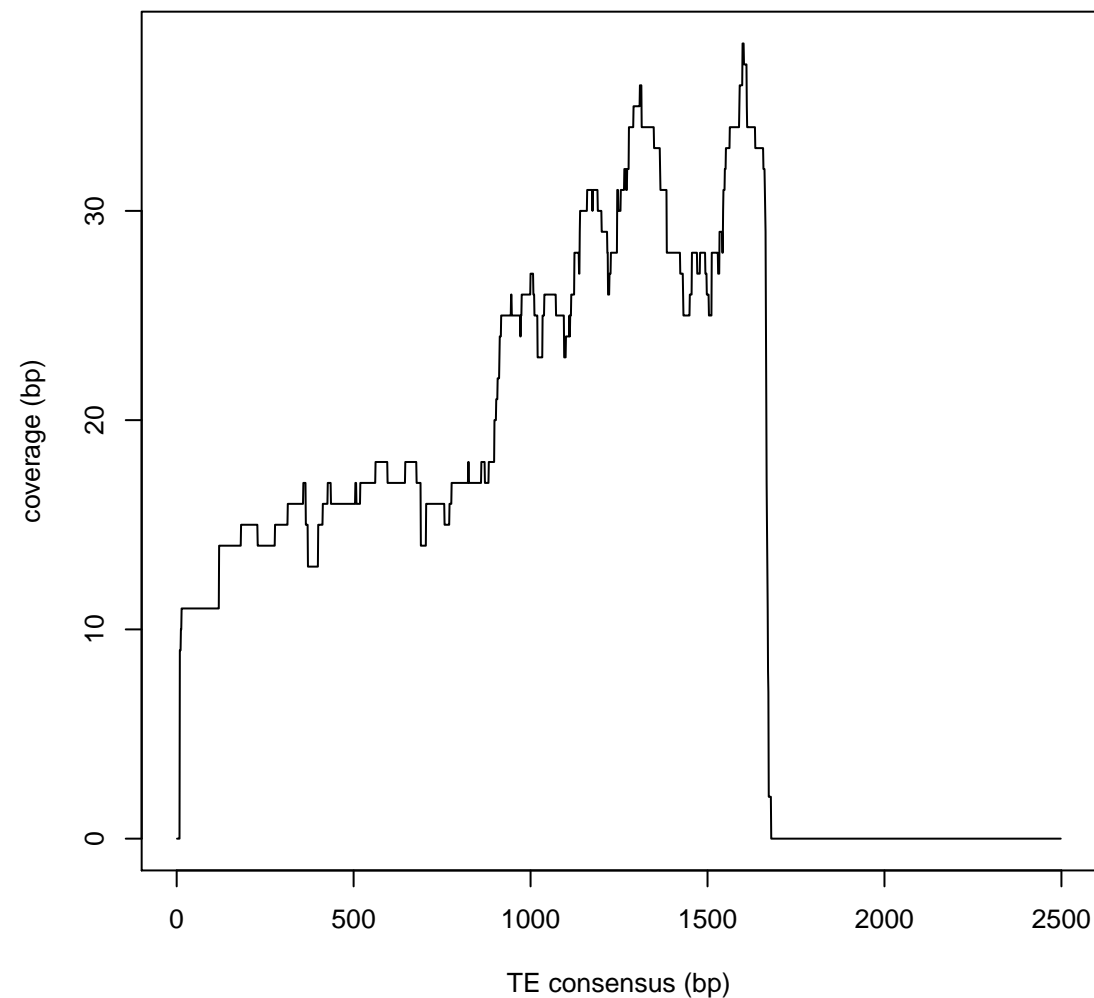


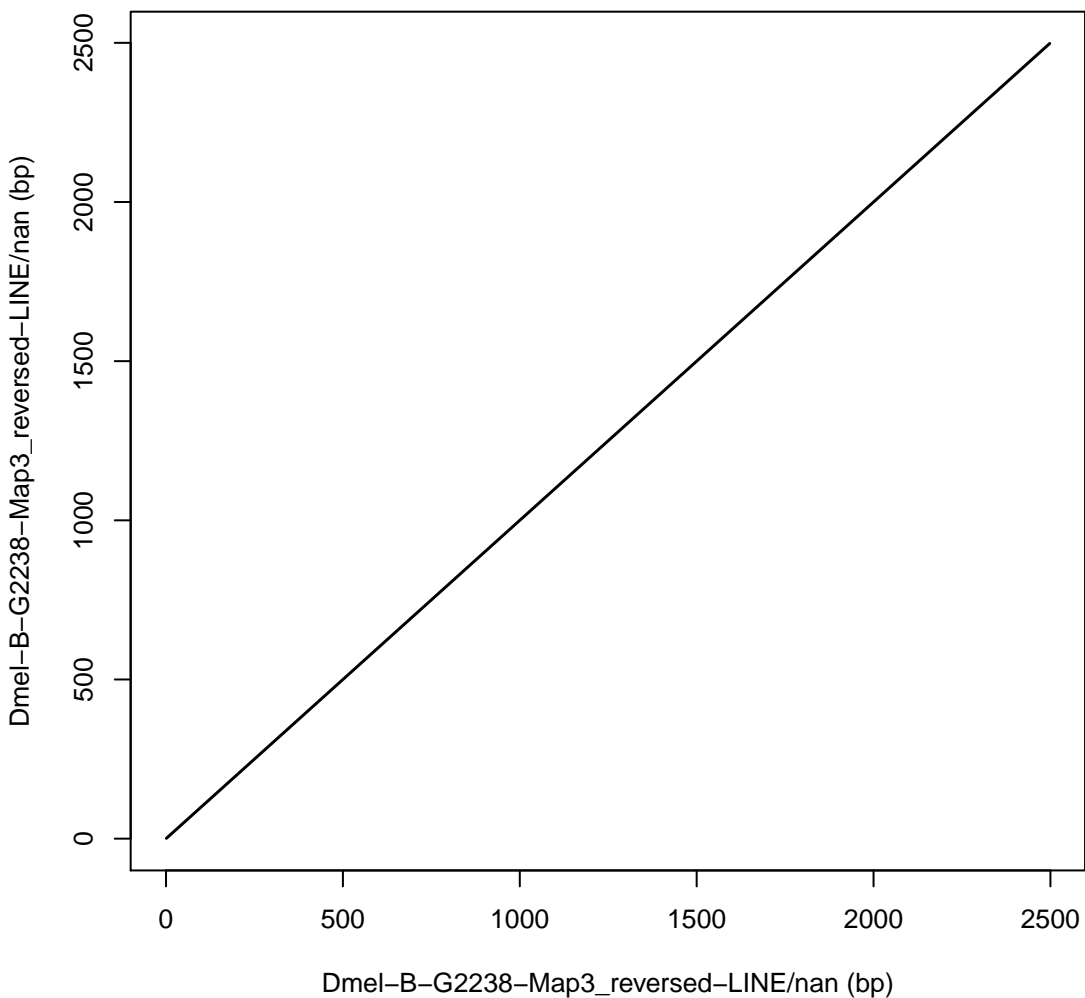
TE: Dmel-B-G2238-Map3\_reversed-LINE/nan  
consensus size: 2498bp; fragments: 93; full length: 0 ( $\geq 2248.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

