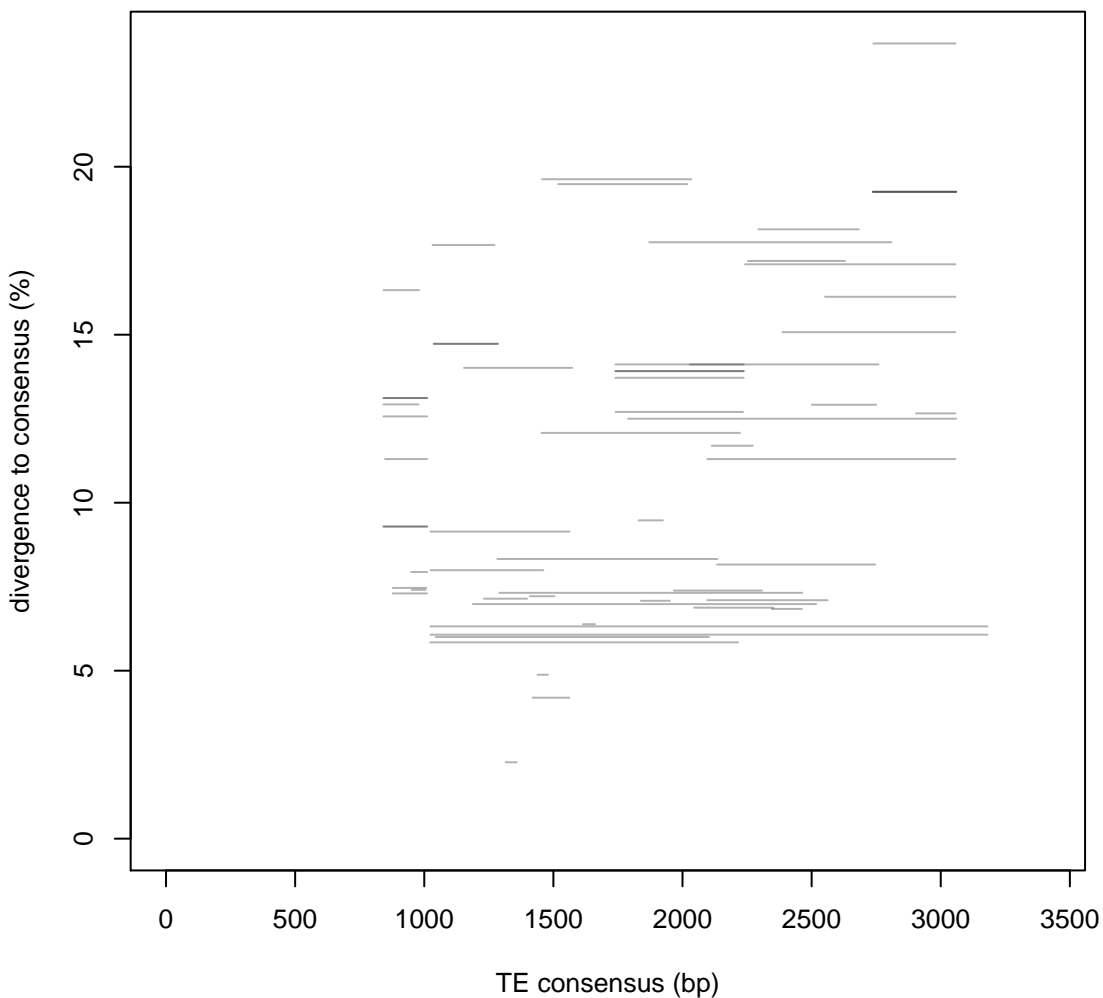
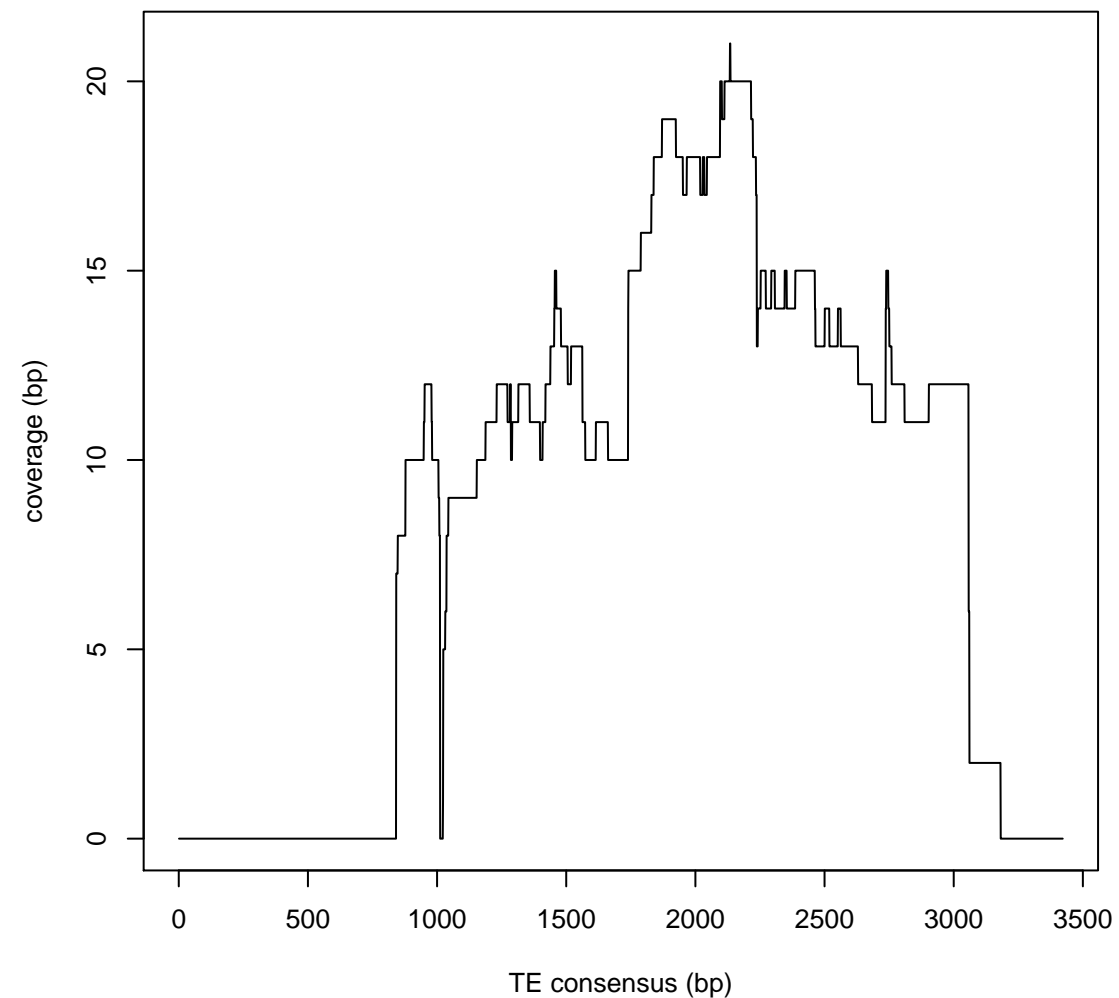


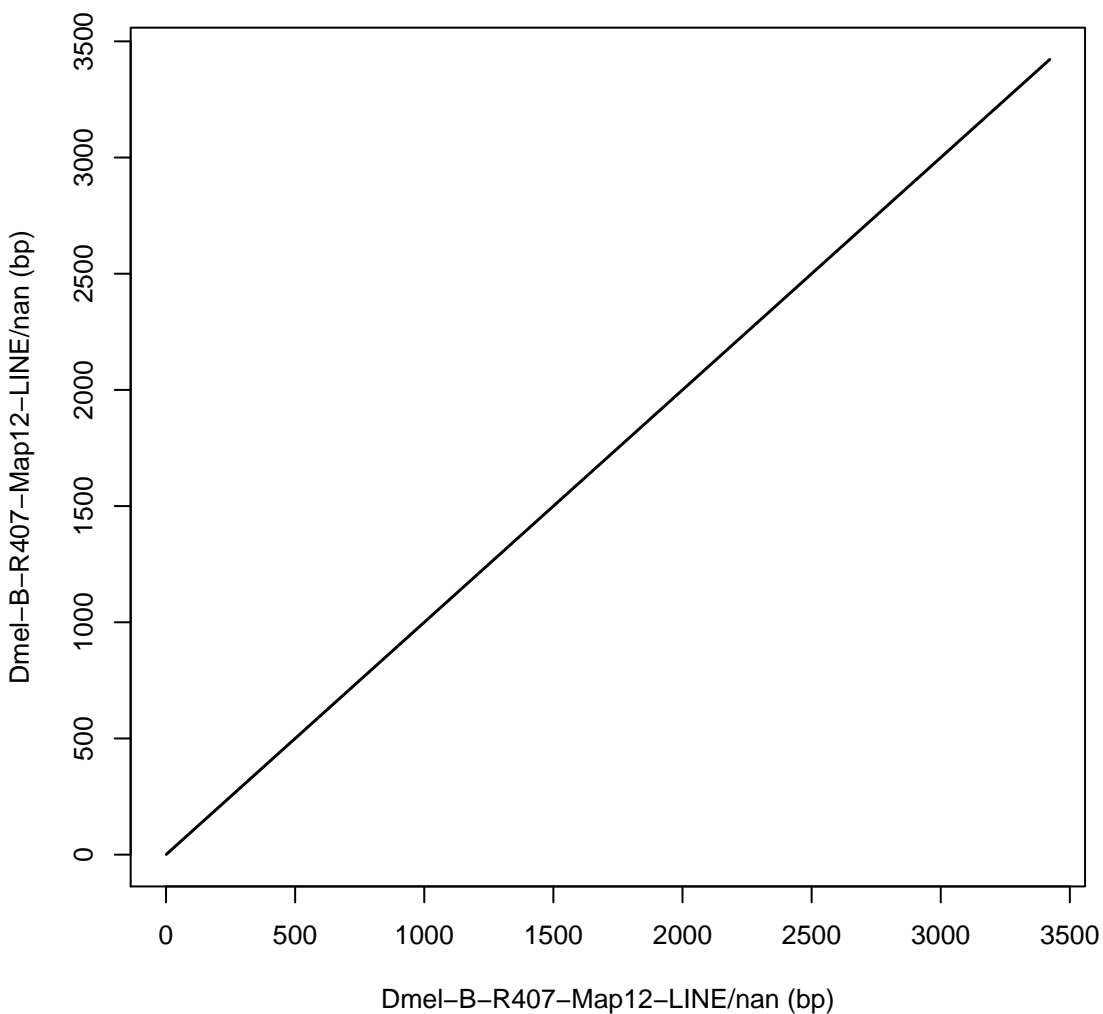
TE: Dmel-B-R407-Map12-LINE/nan  
consensus size: 3422bp; fragments: 62; full length: 0 ( $\geq 3079.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

