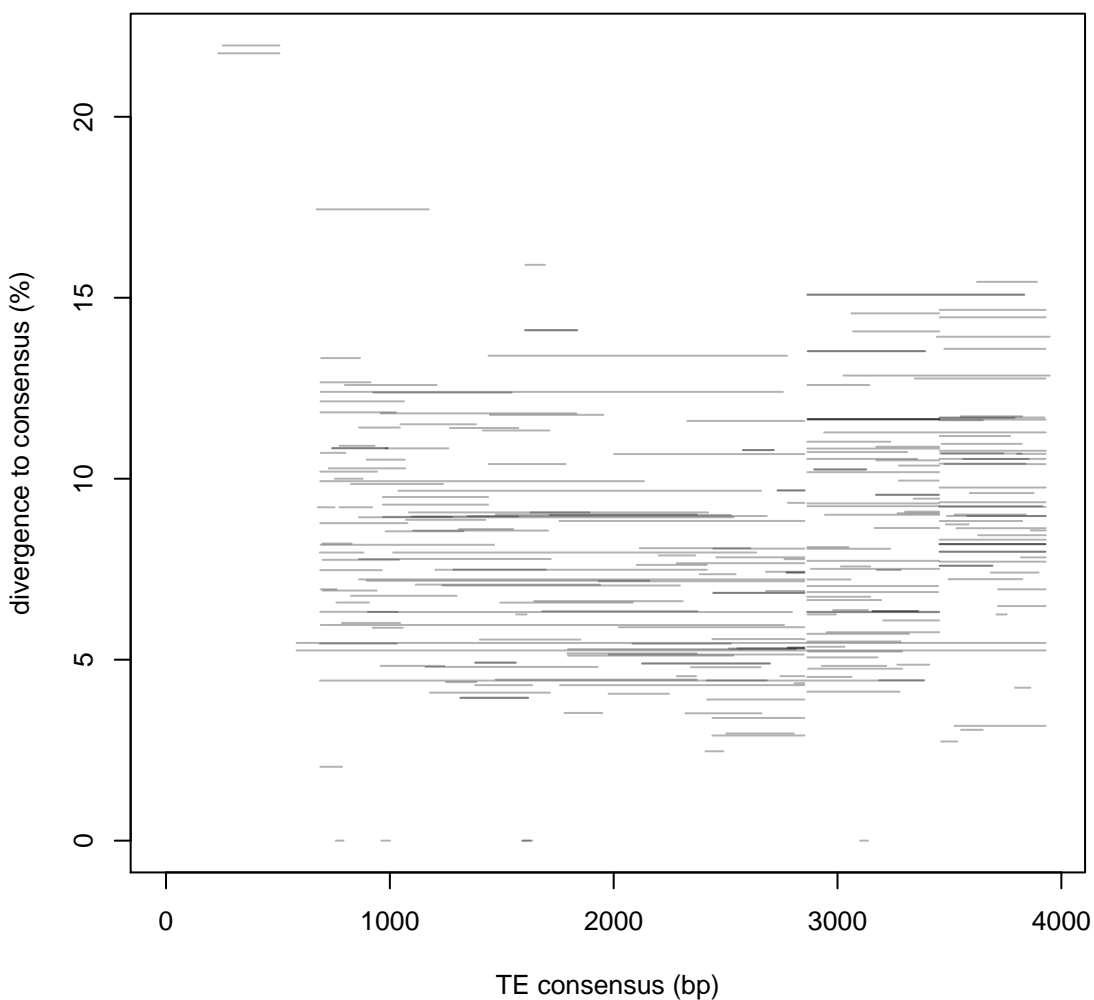
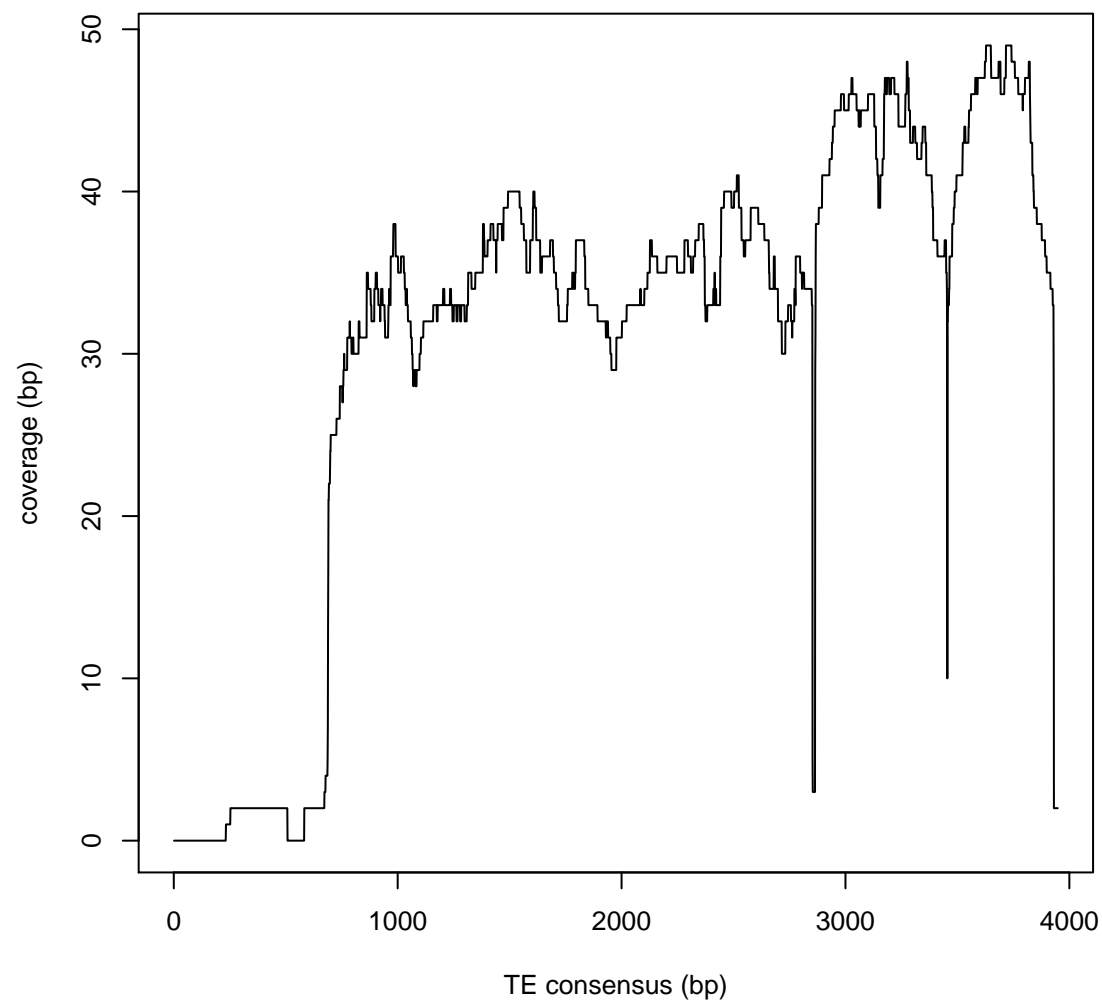


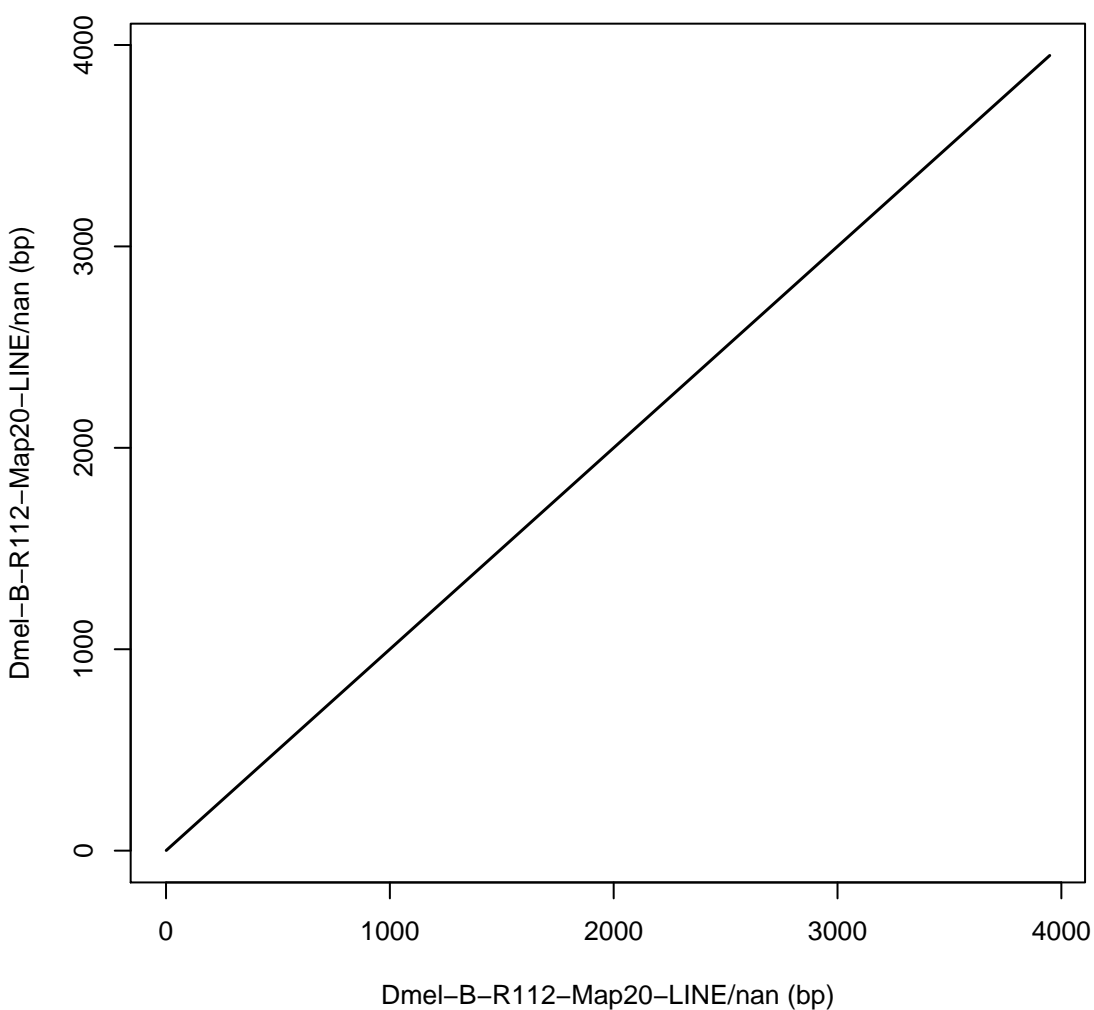
TE: Dmel-B-R112-Map20-LINE/nan  
consensus size: 3948bp; fragments: 262; full length: 0 ( $\geq 3553.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

