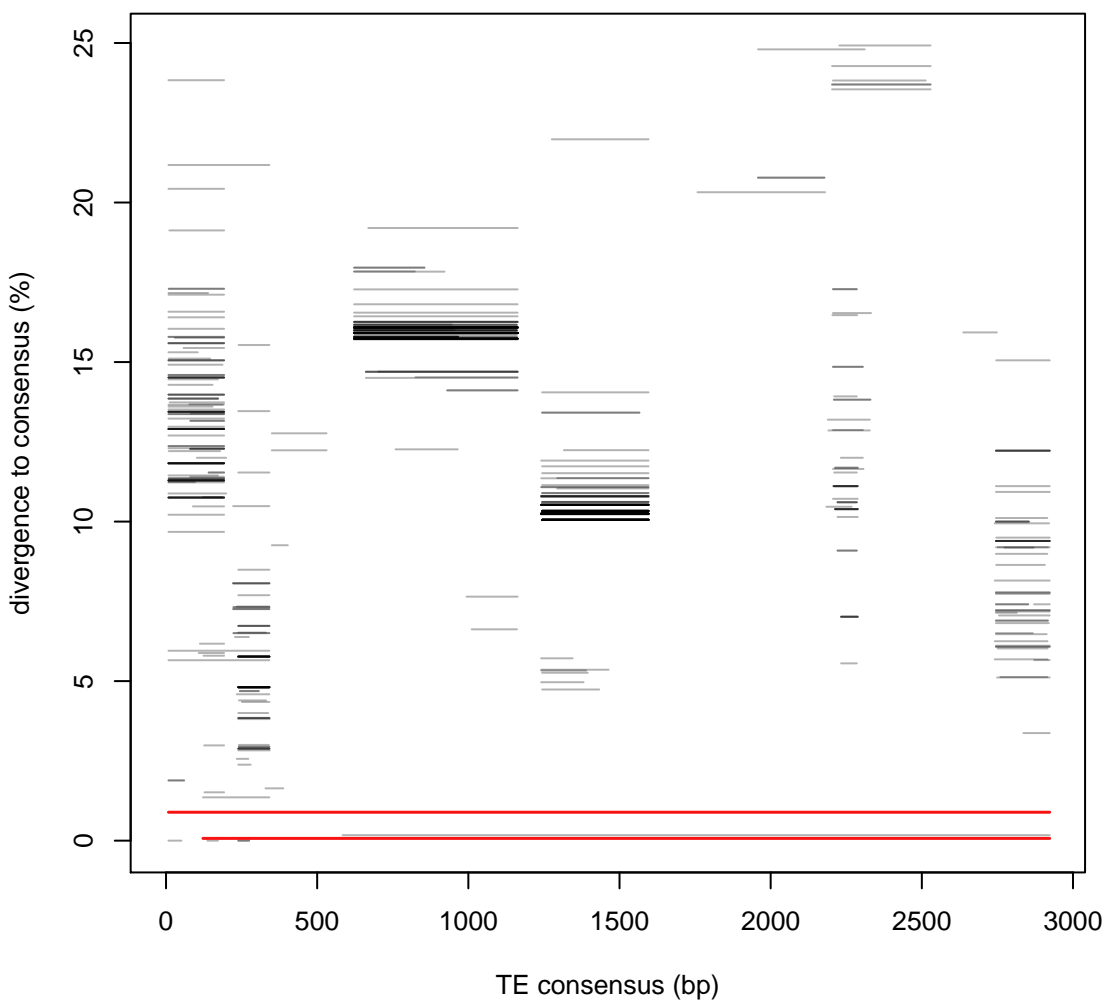
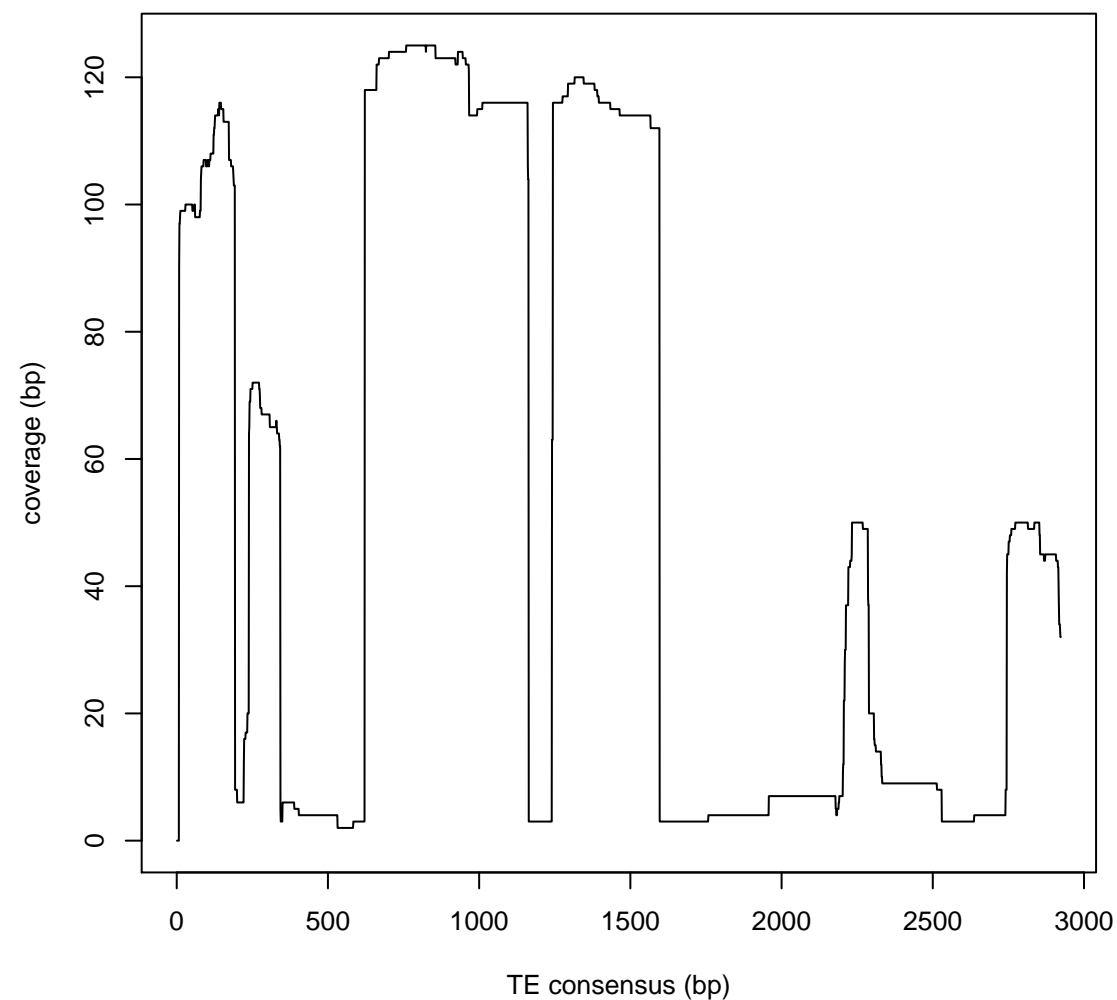


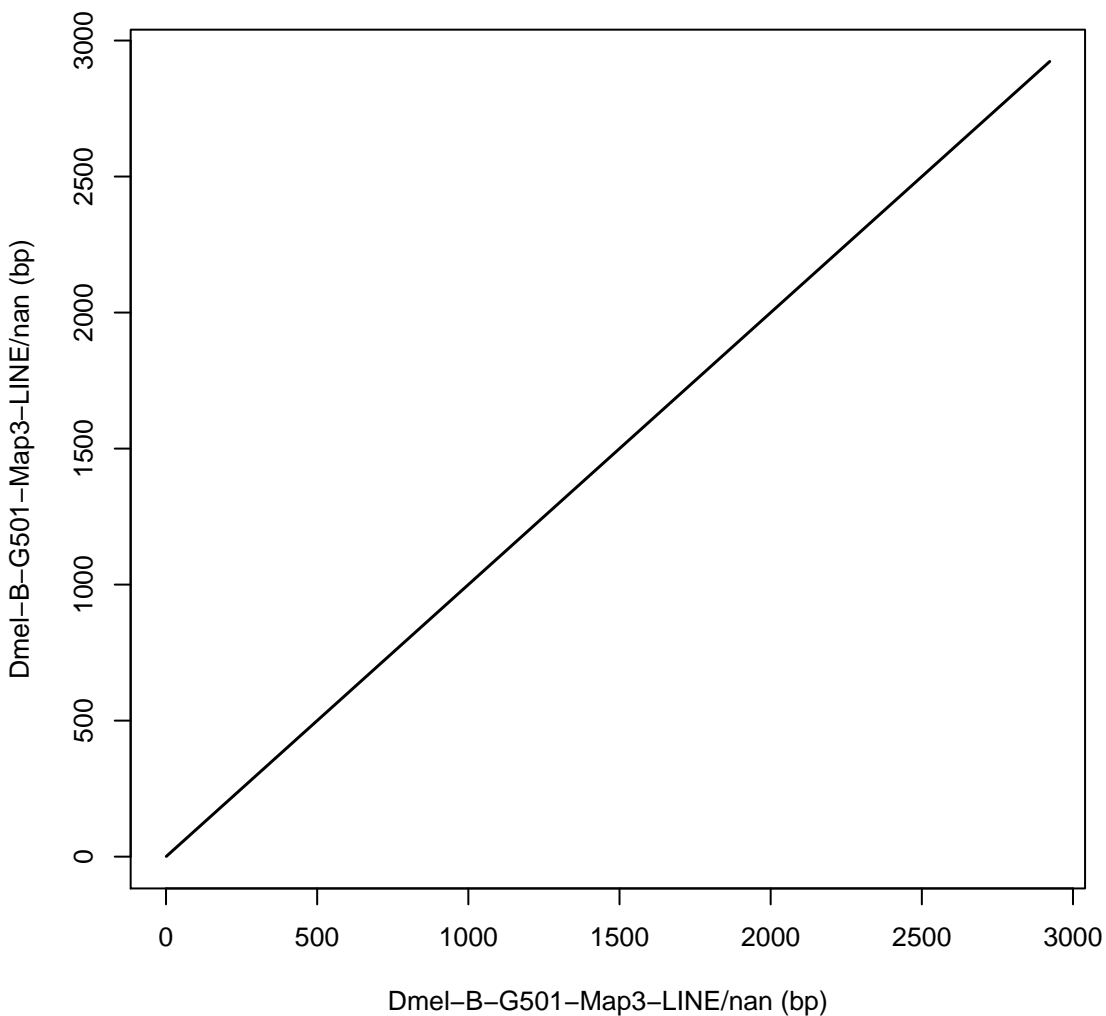
TE: Dmel-B-G501-Map3-LINE/nan  
consensus size: 2923bp; fragments: 538; full length: 2 ( $\geq 2630.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

