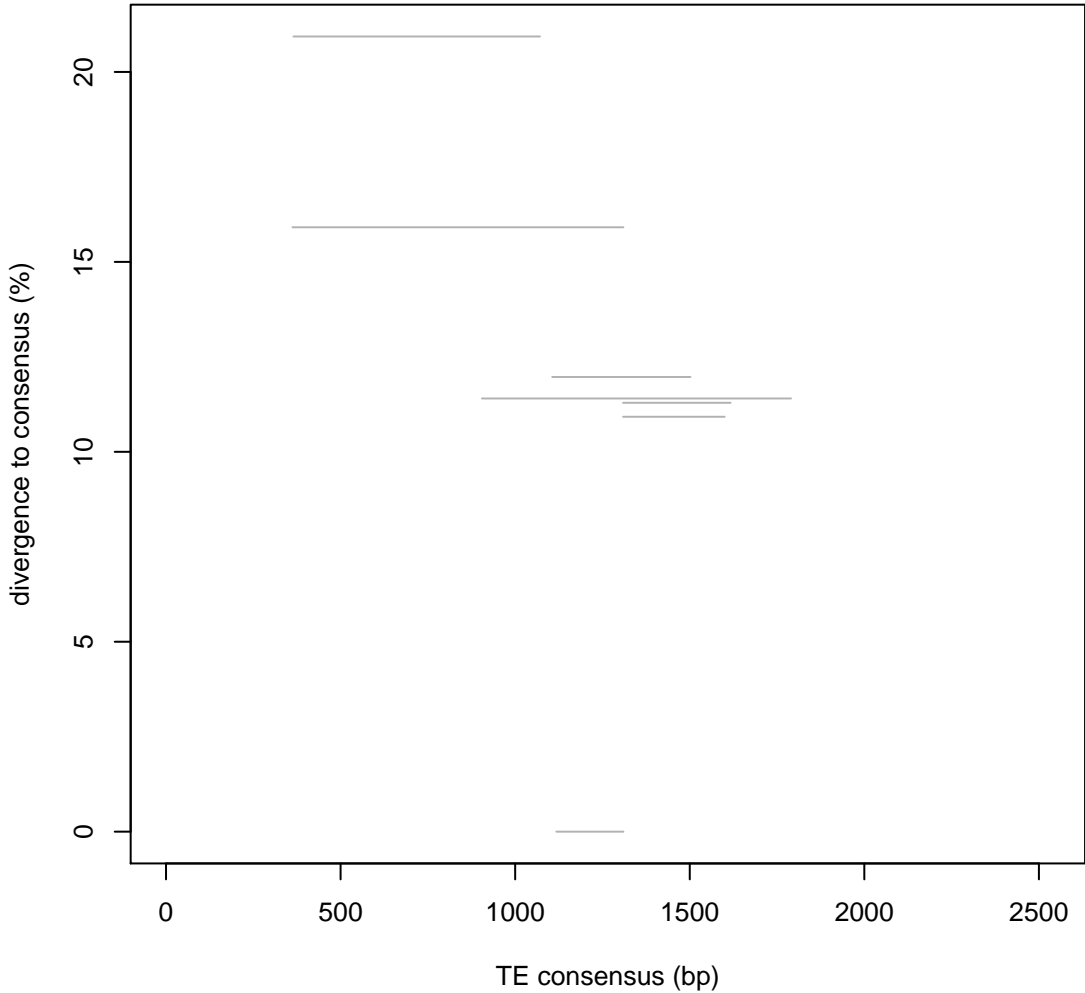
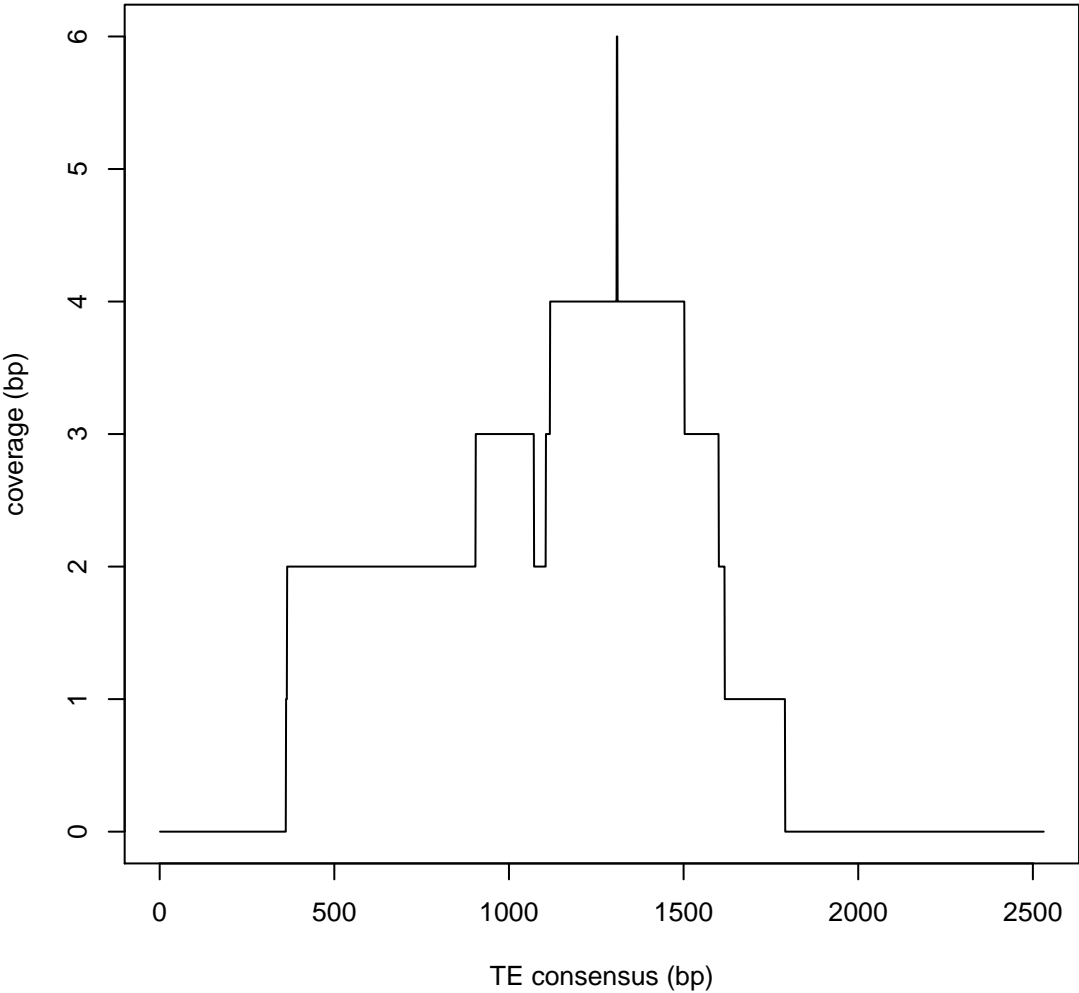


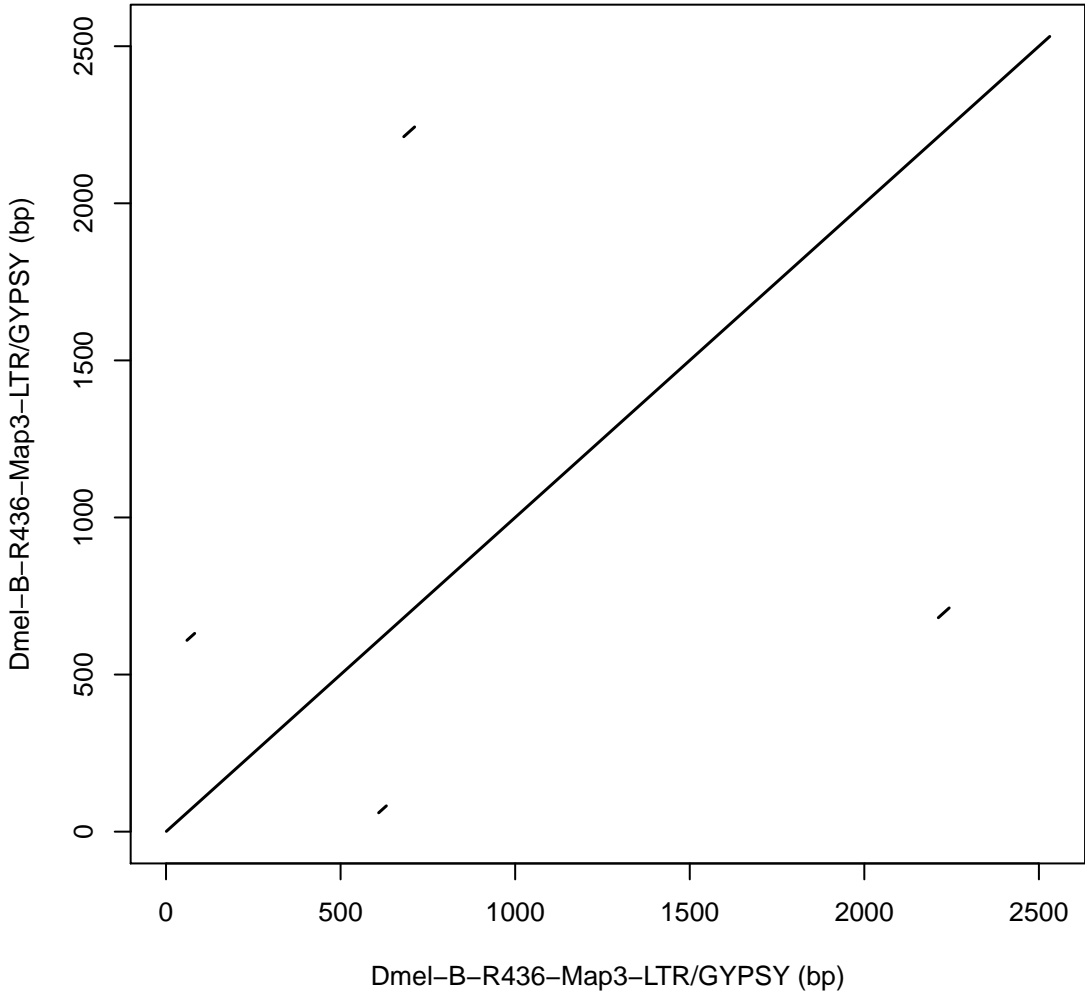
TE: Dmel-B-R436-Map3-LTR/GYPSY  
consensus size: 2531bp; fragments: 7; full length: 0 (>=2277.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

