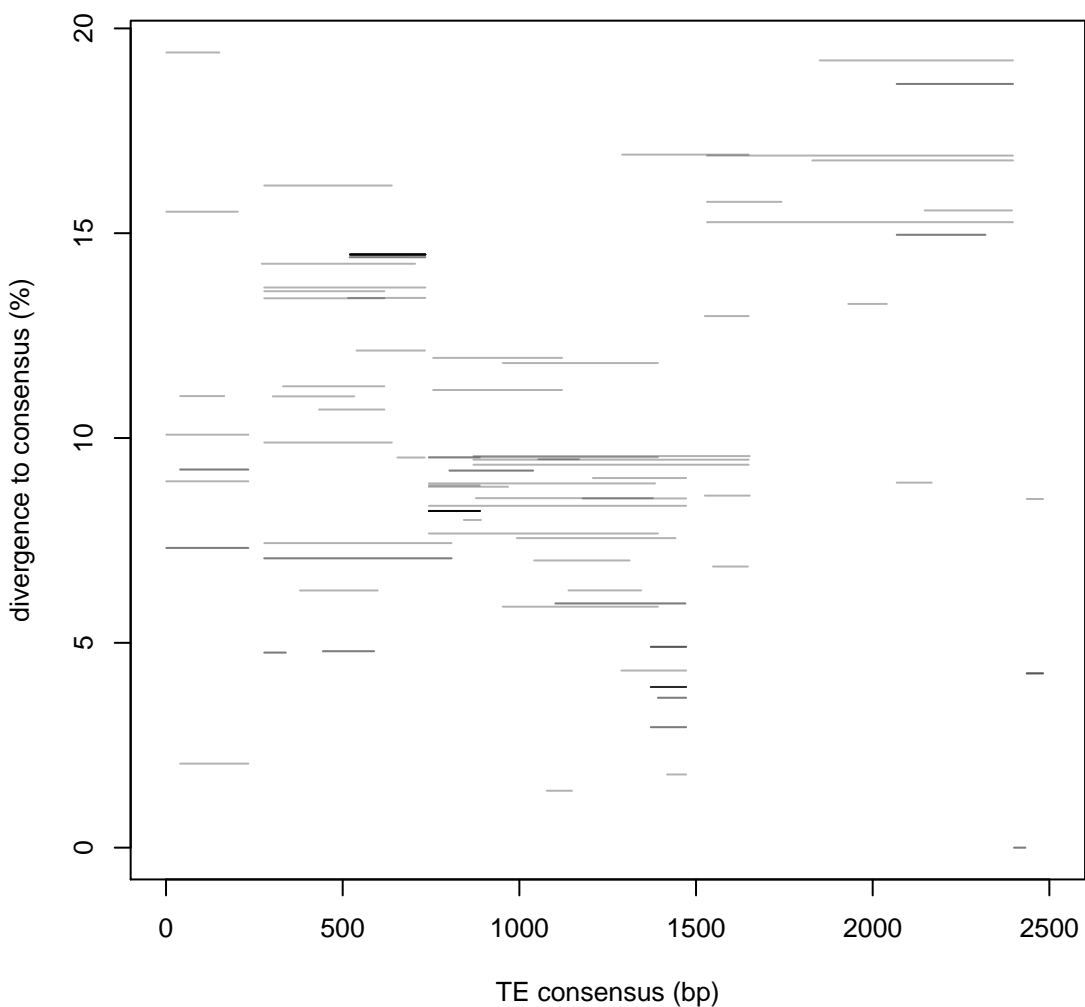
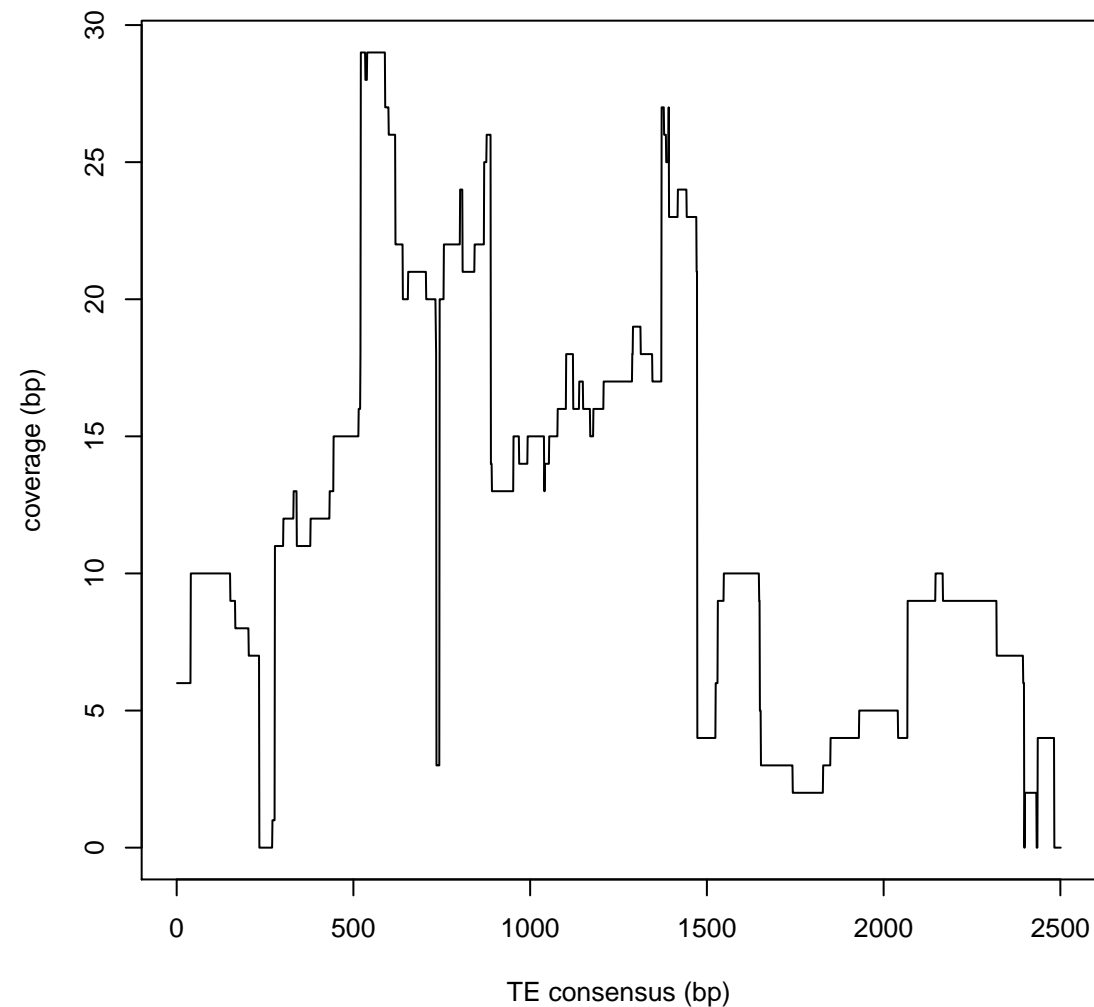


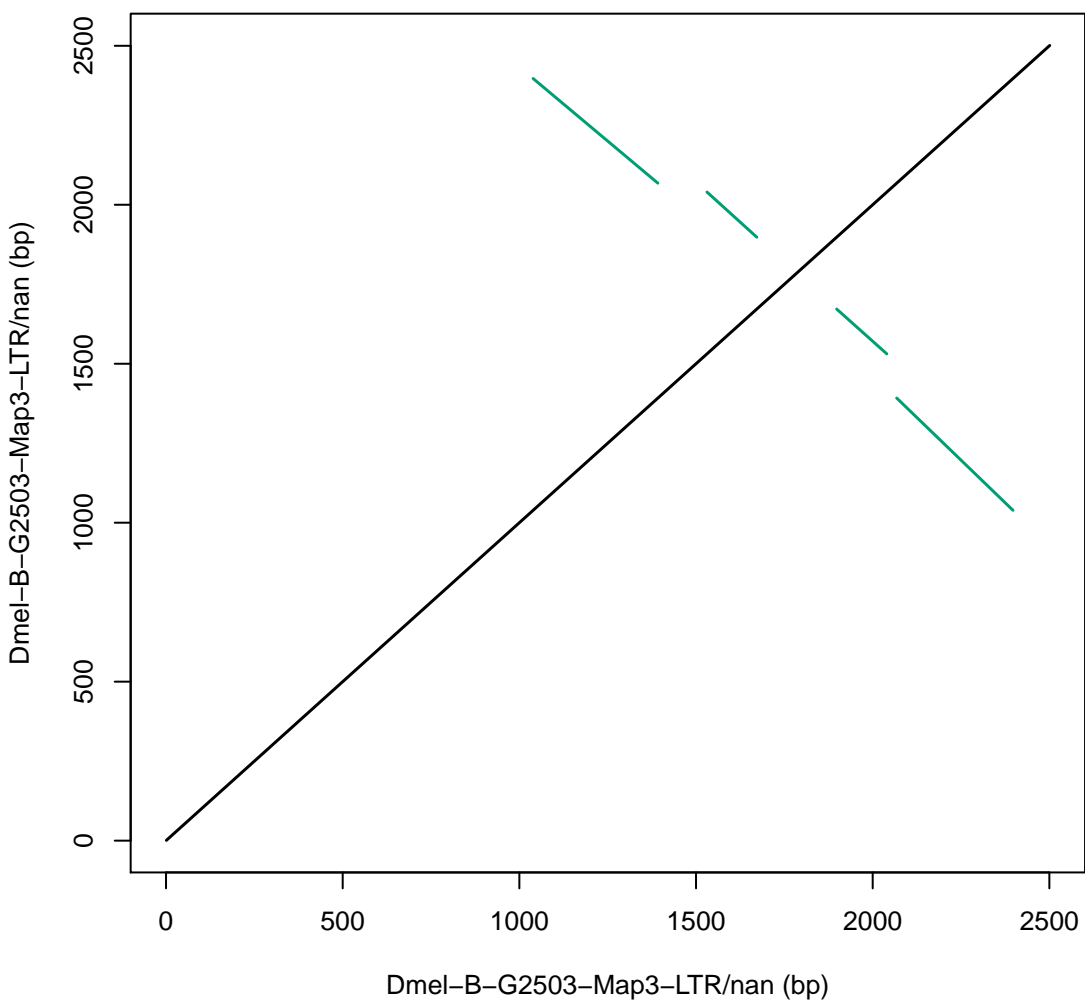
TE: Dmel-B-G2503-Map3-LTR/nan  
consensus size: 2501bp; fragments: 116; full length: 0 (>=2250.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

