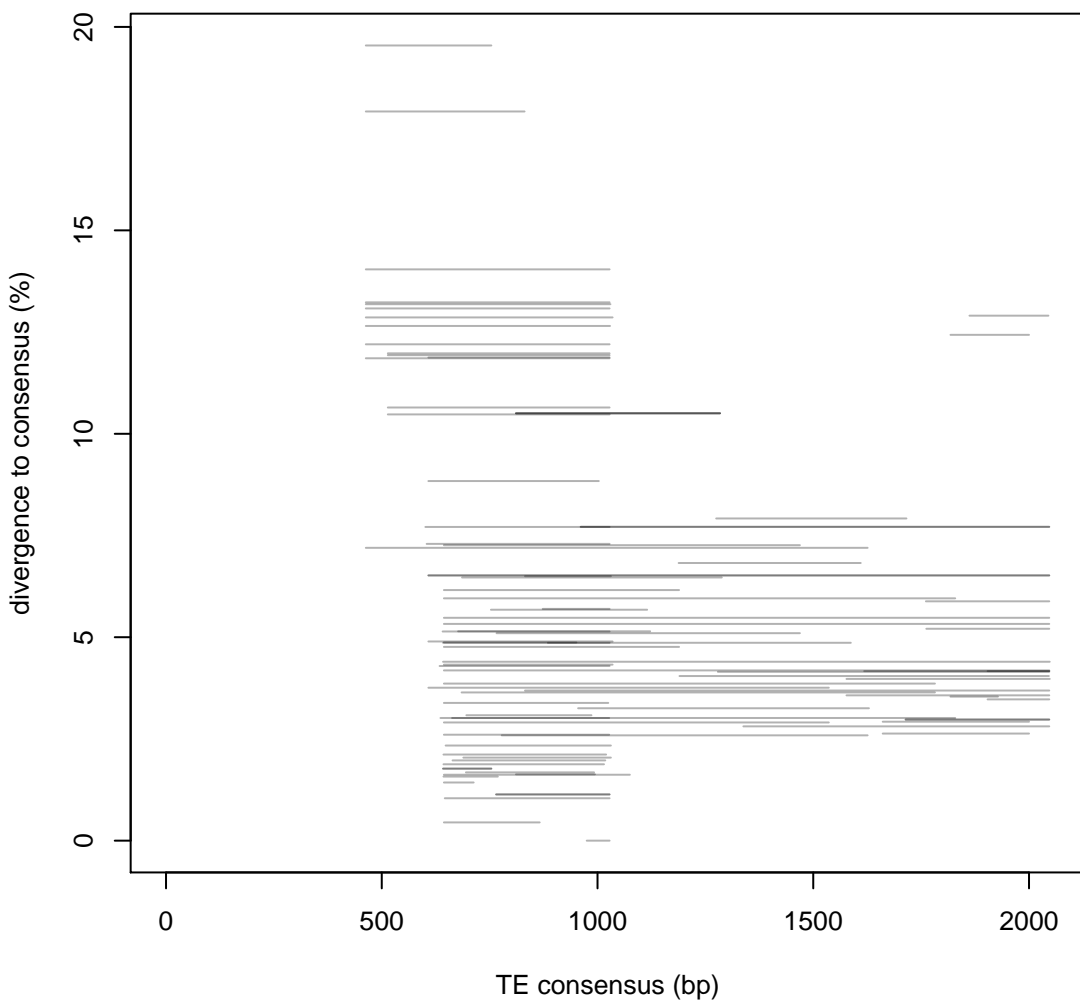
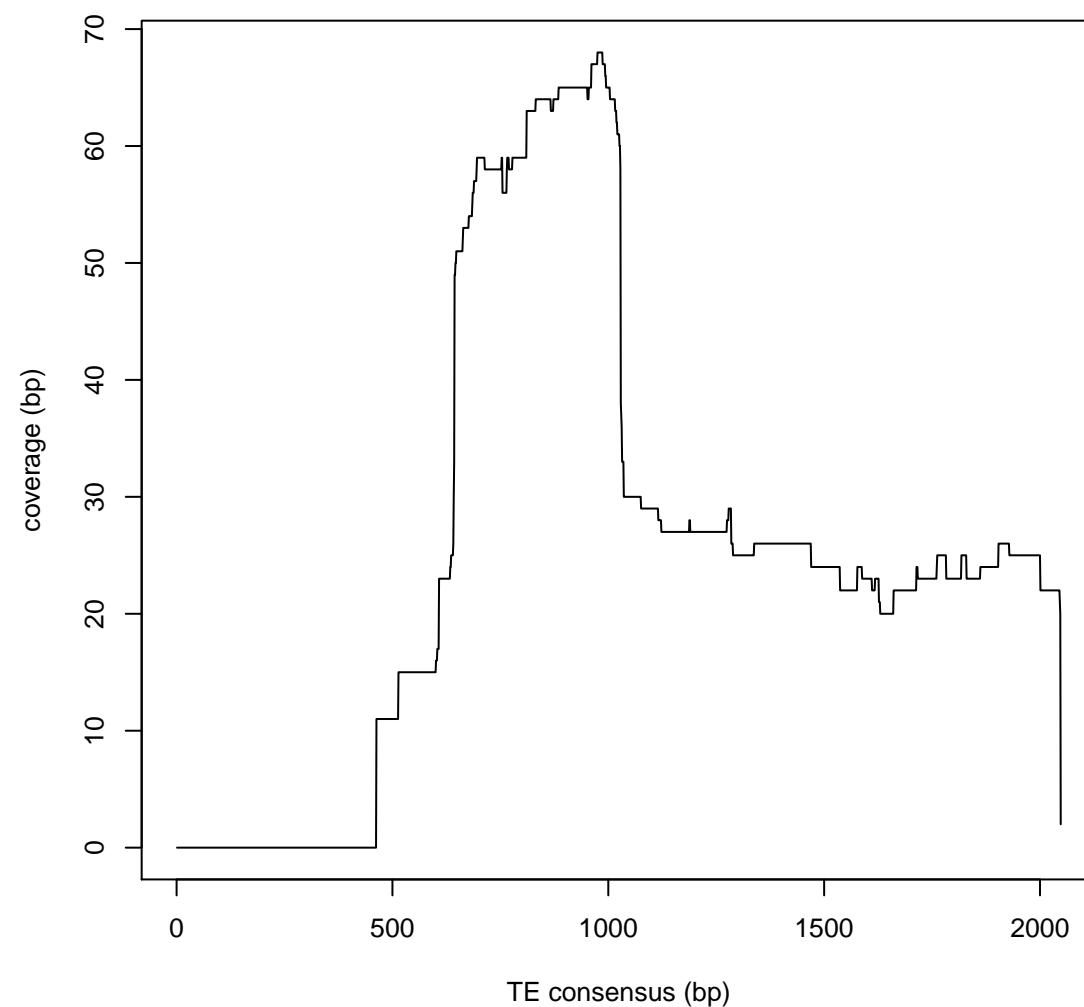


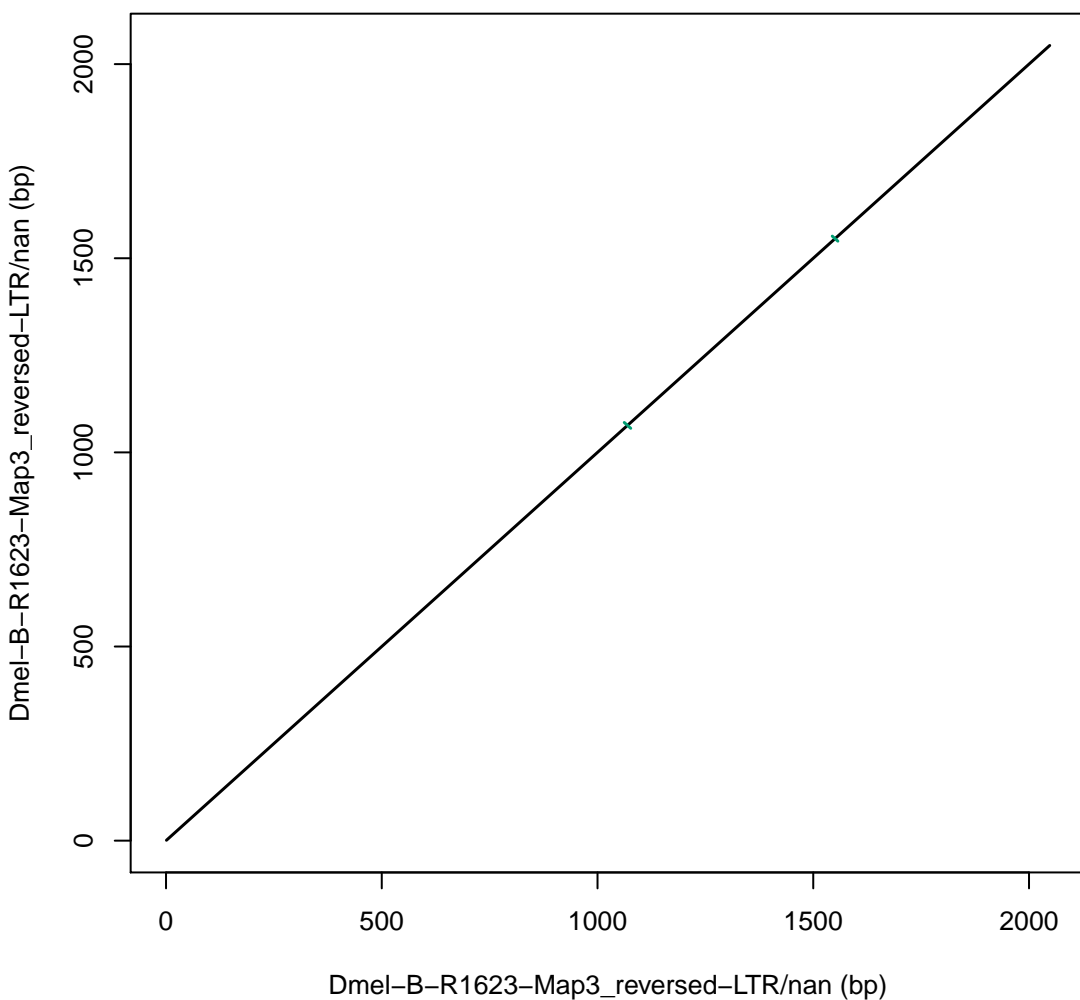
TE: Dmel-B-R1623-Map3\_reversed-LTR/nan  
consensus size: 2048bp; fragments: 95; full length: 0 (>=1843.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

