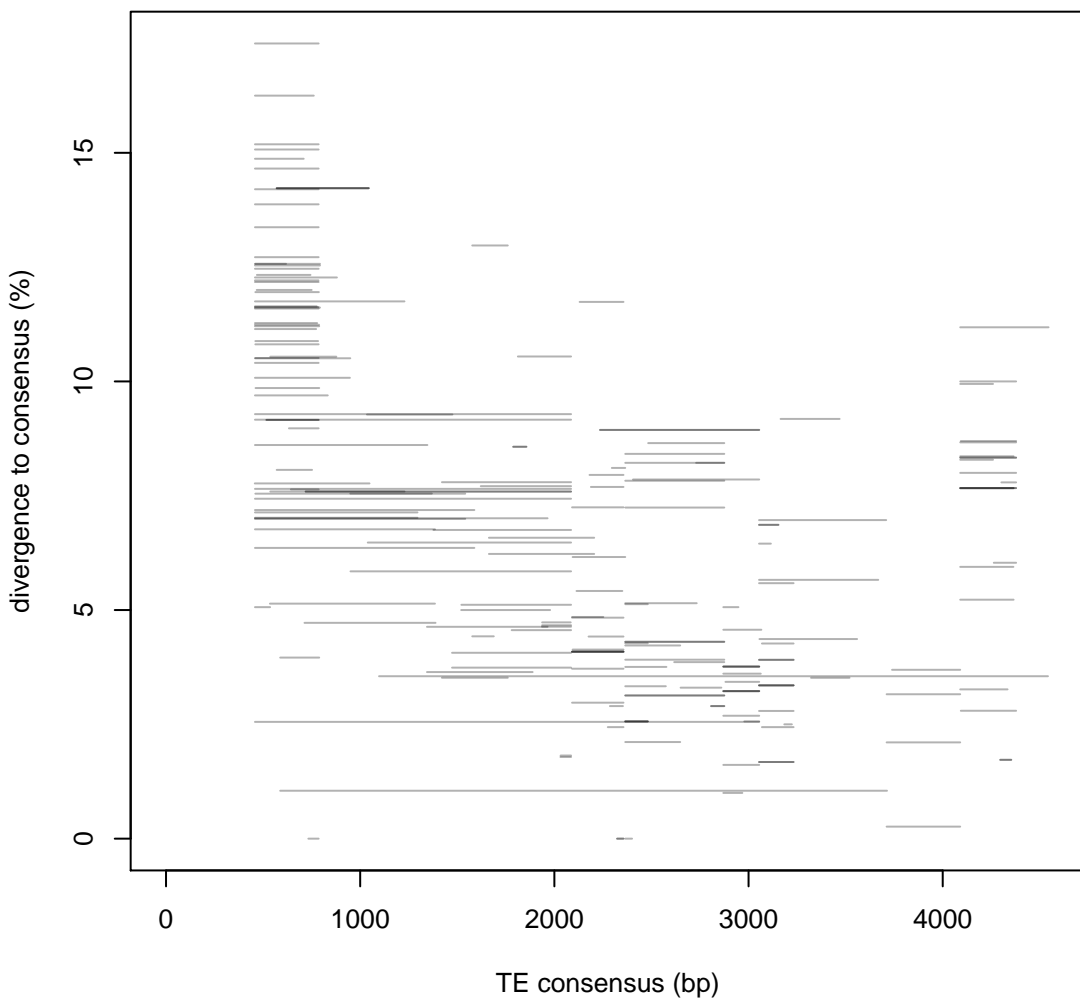
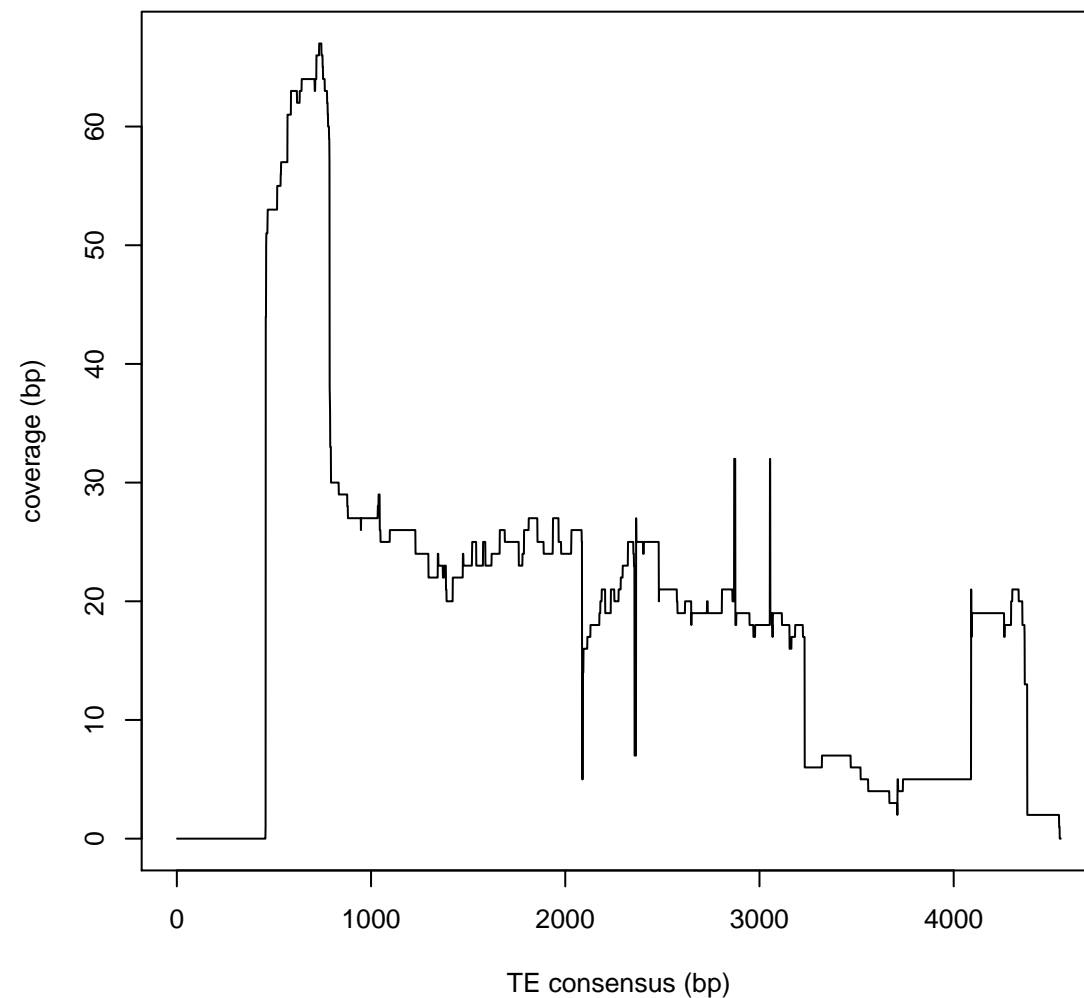


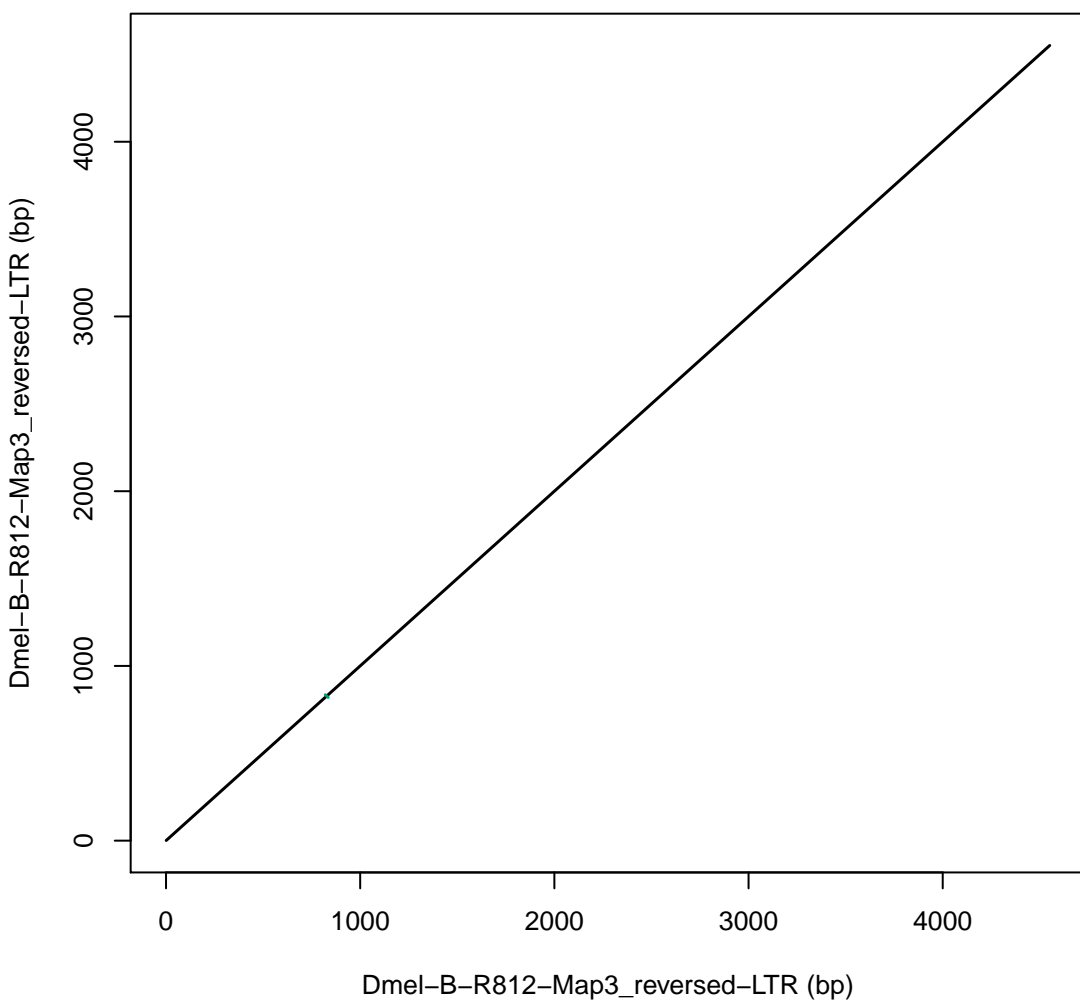
TE: Dmel-B-R812-Map3_reversed-LTR
consensus size: 4551bp; fragments: 208; full length: 0 (>=4095.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

