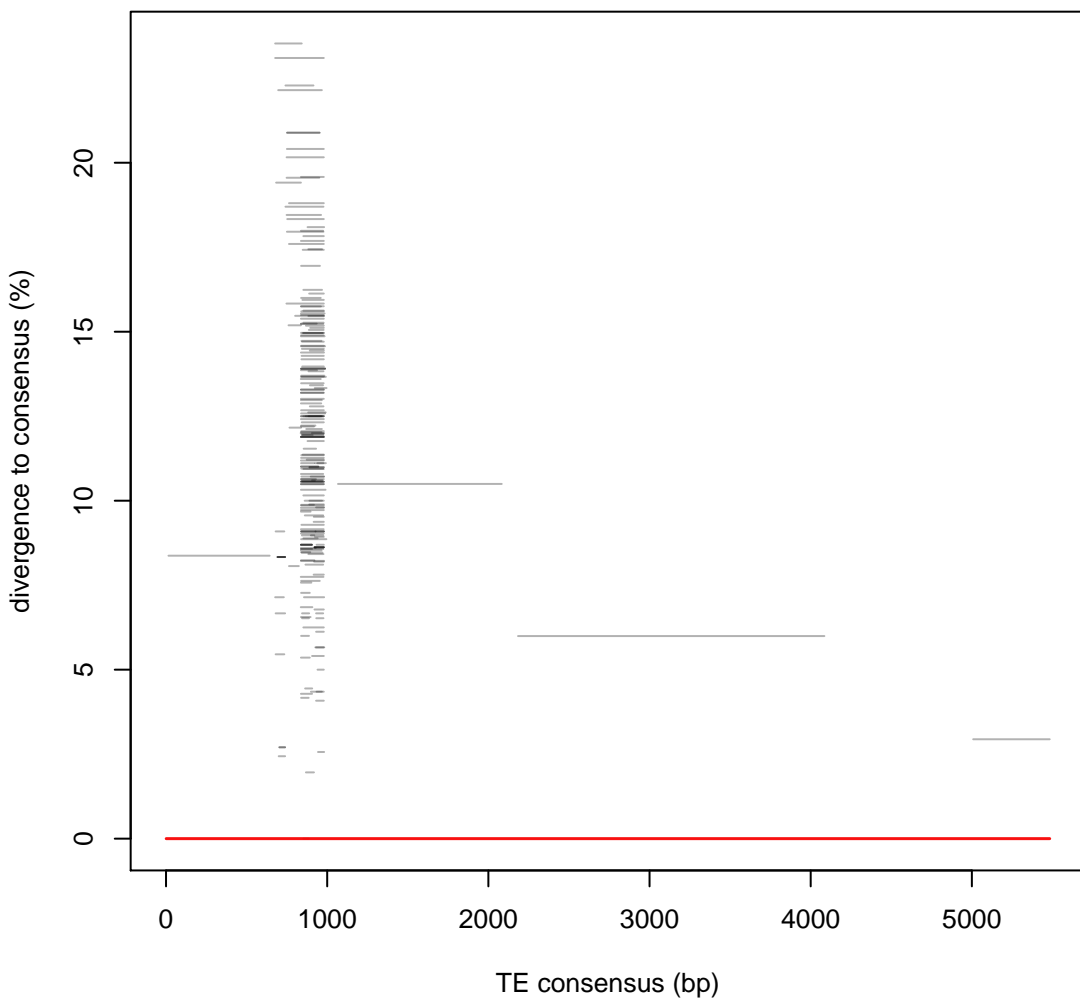
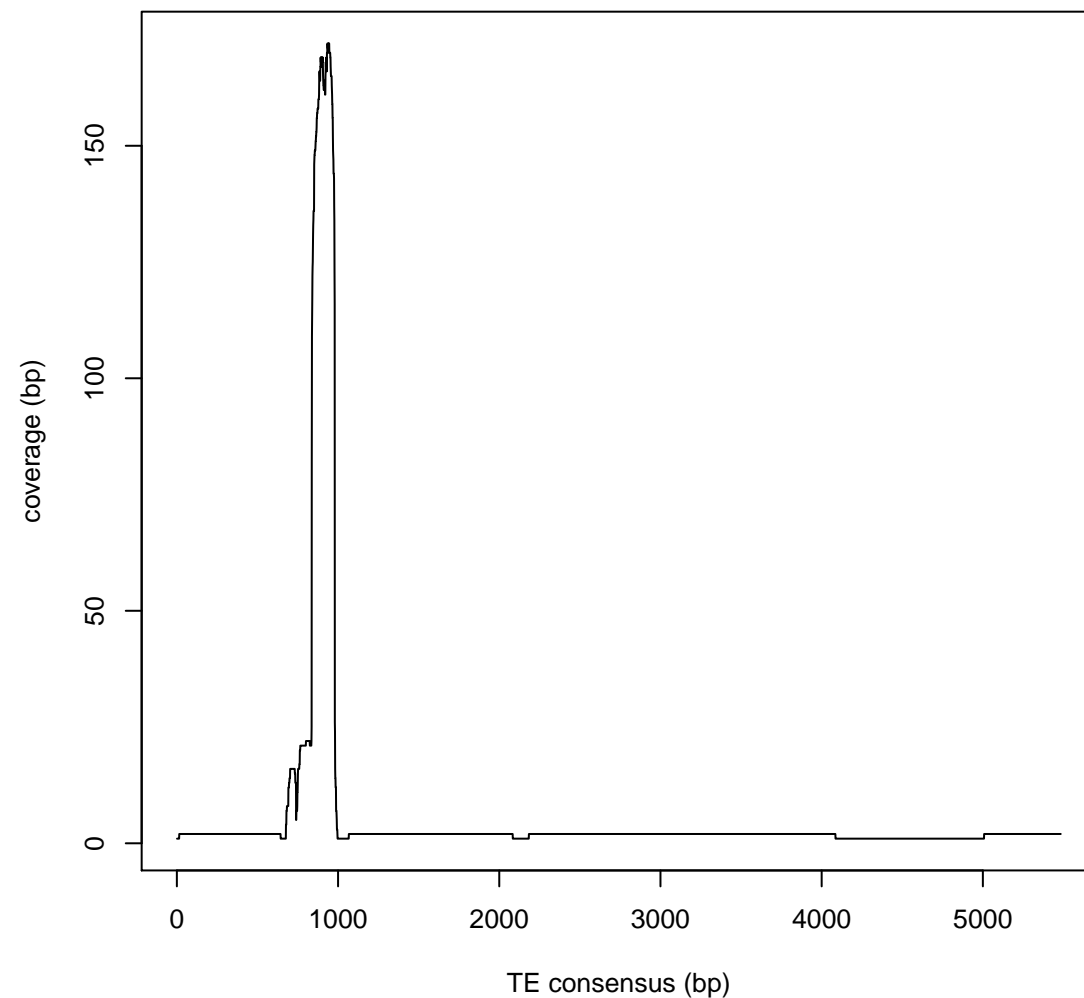


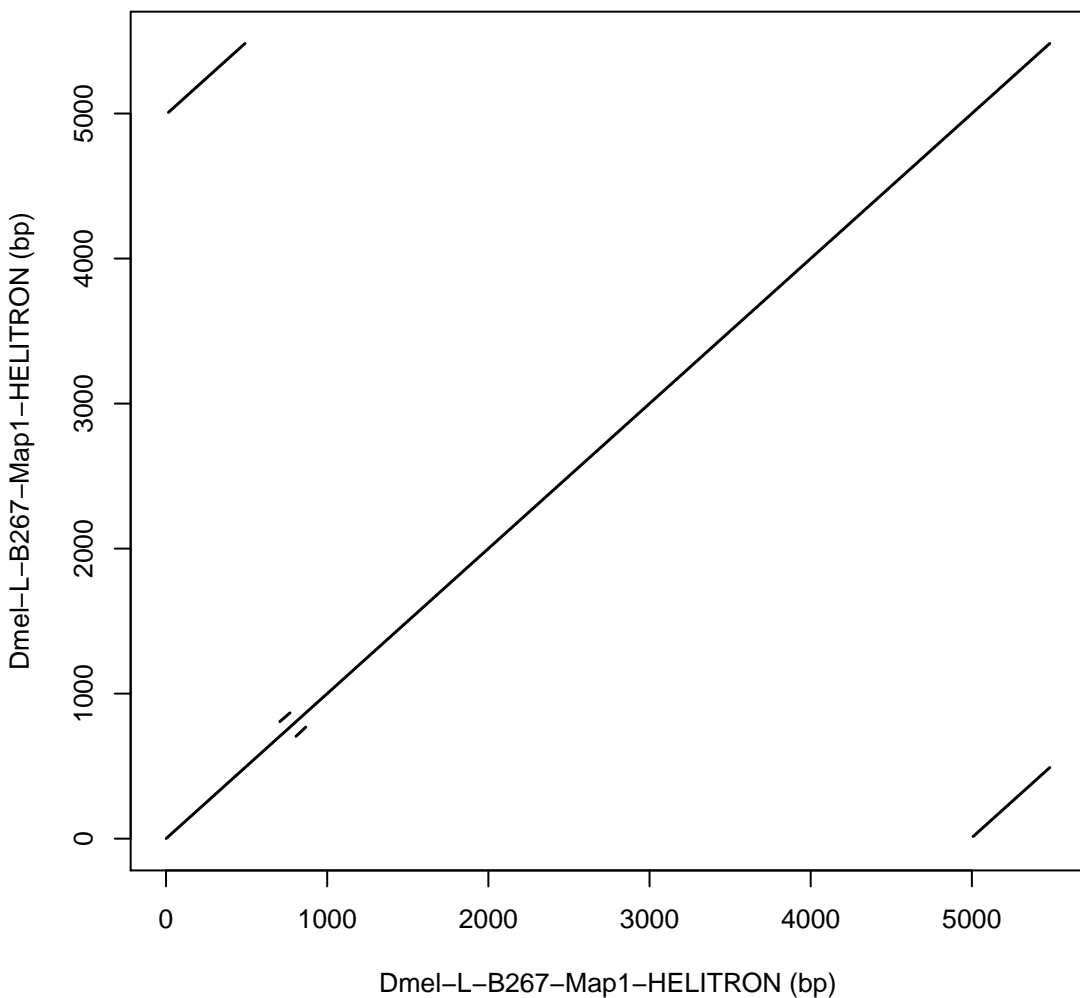
TE: Dmel-L-B267-Map1-HELITRON
consensus size: 5483bp; fragments: 228; full length: 1 (>=4934.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

