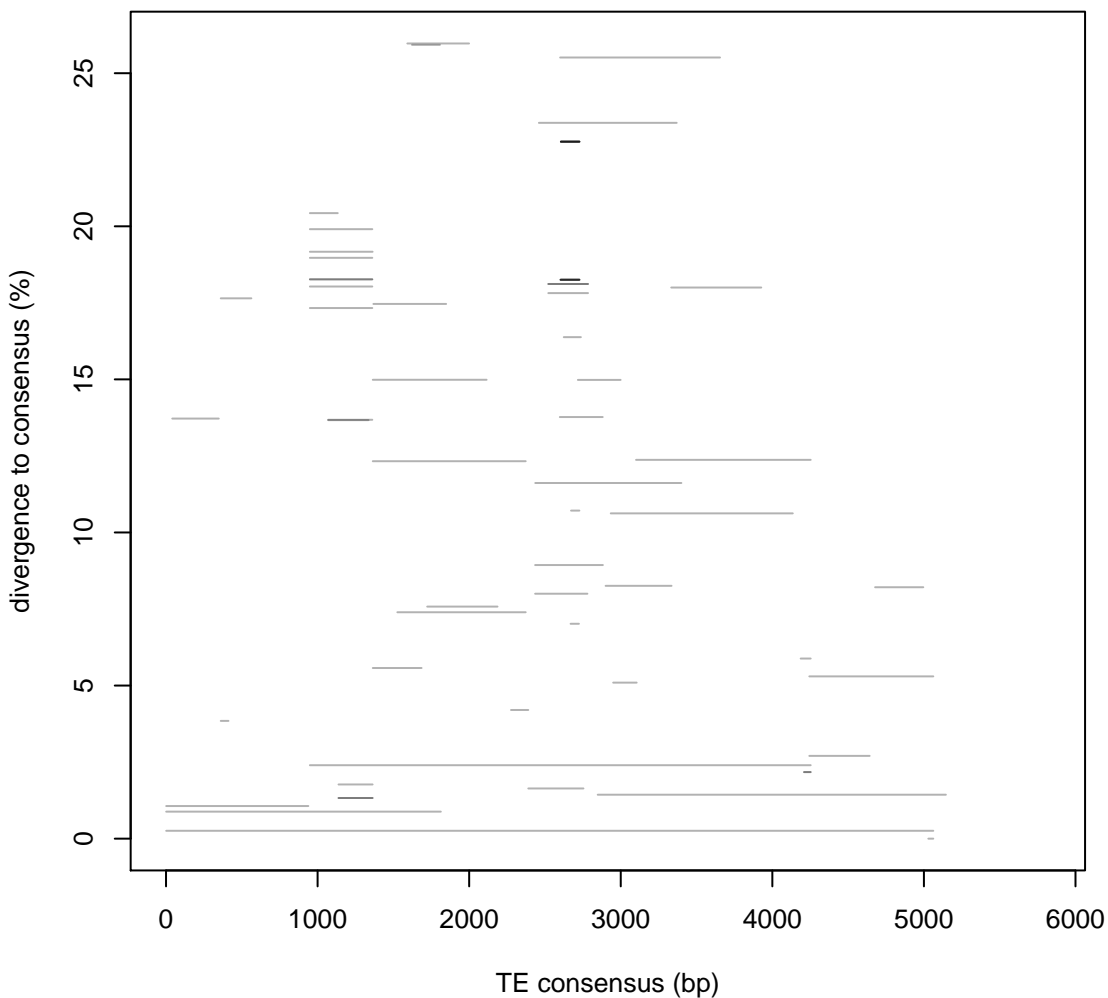
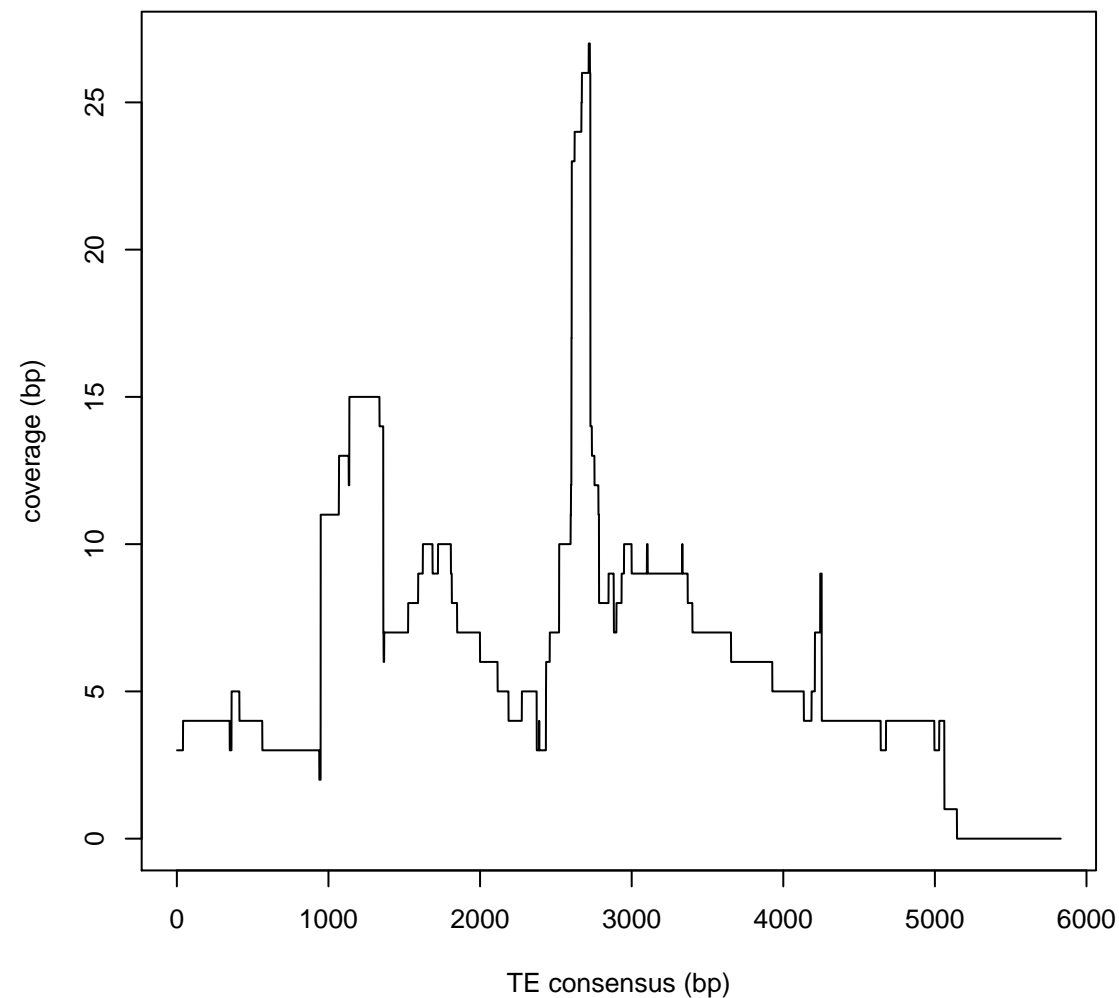


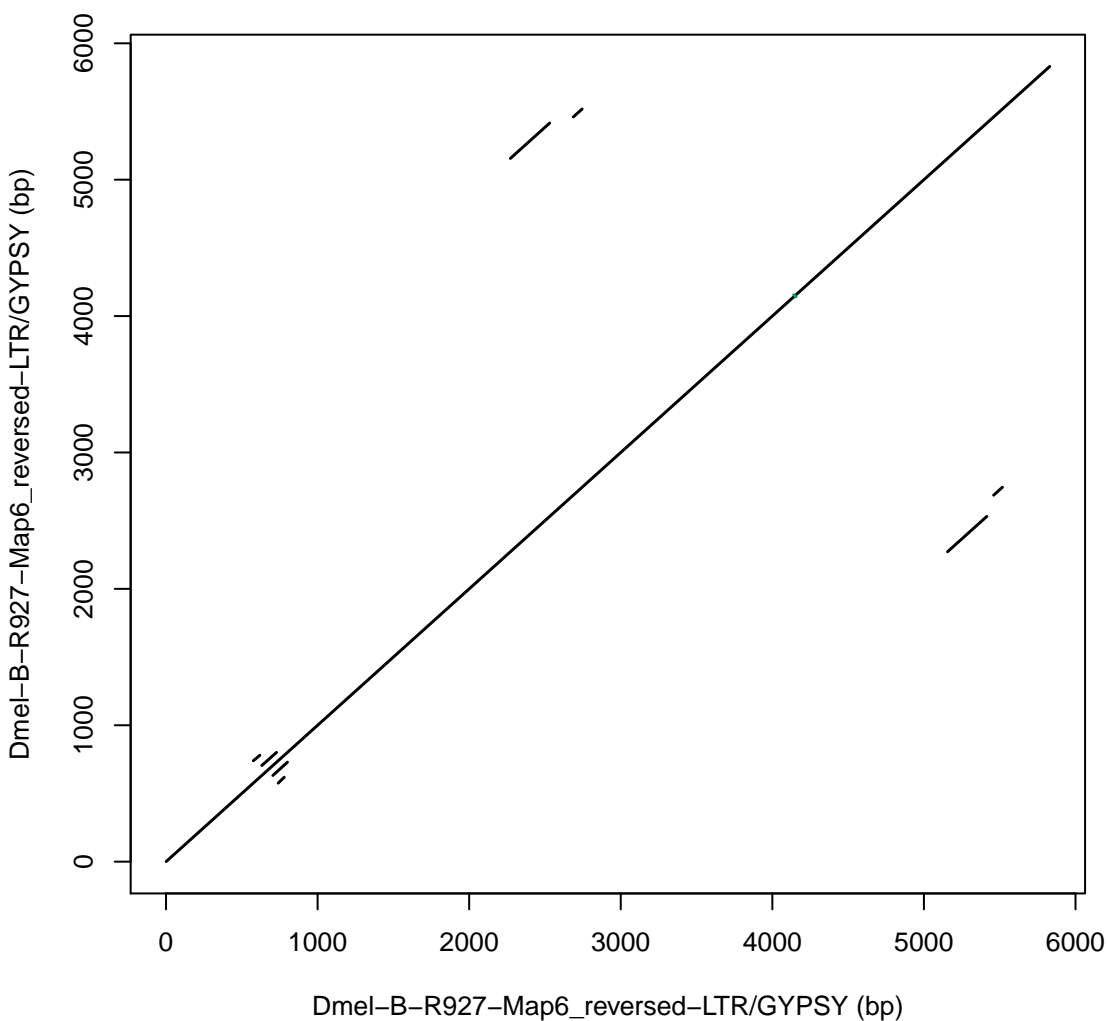
TE: Dmel-B-R927-Map6_reversed-LTR/GYPY
consensus size: 5830bp; fragments: 67; full length: 0 (>=5247bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

