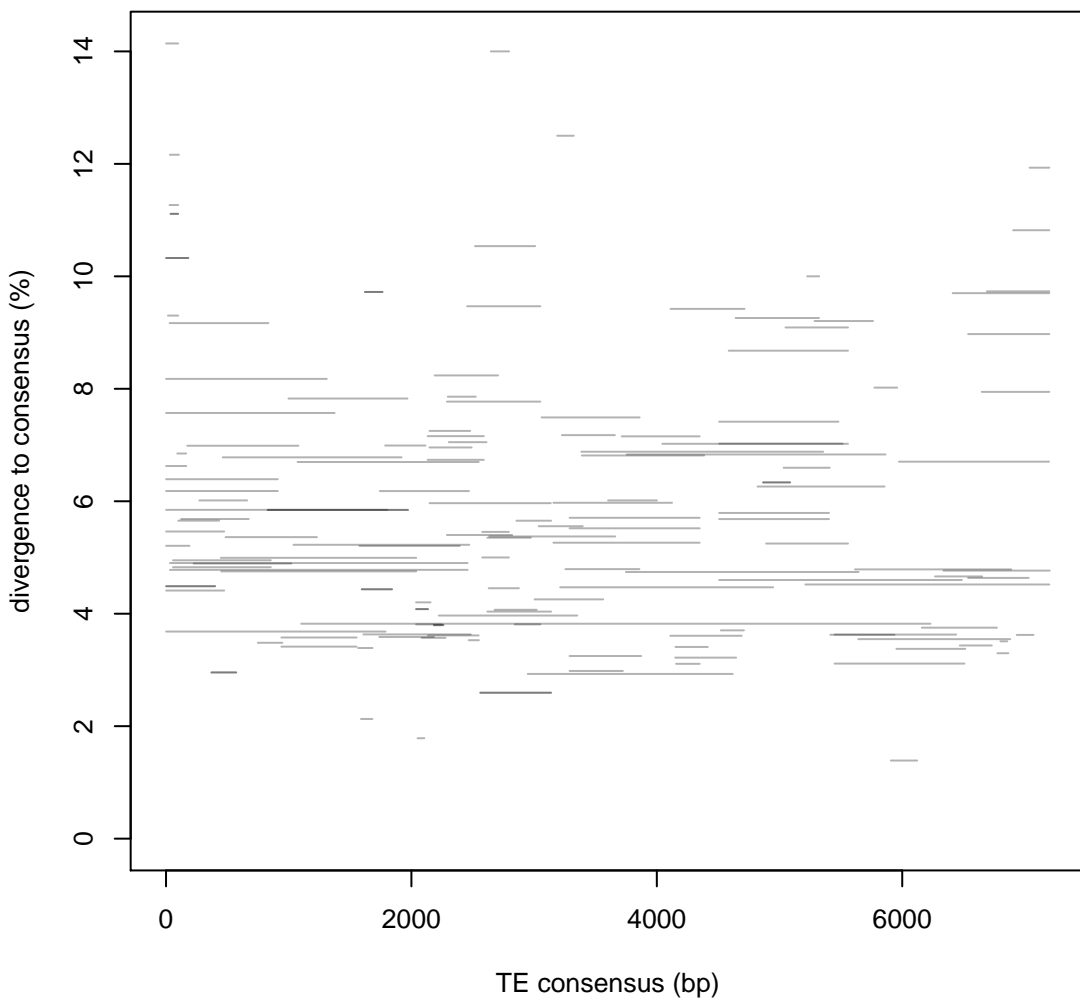
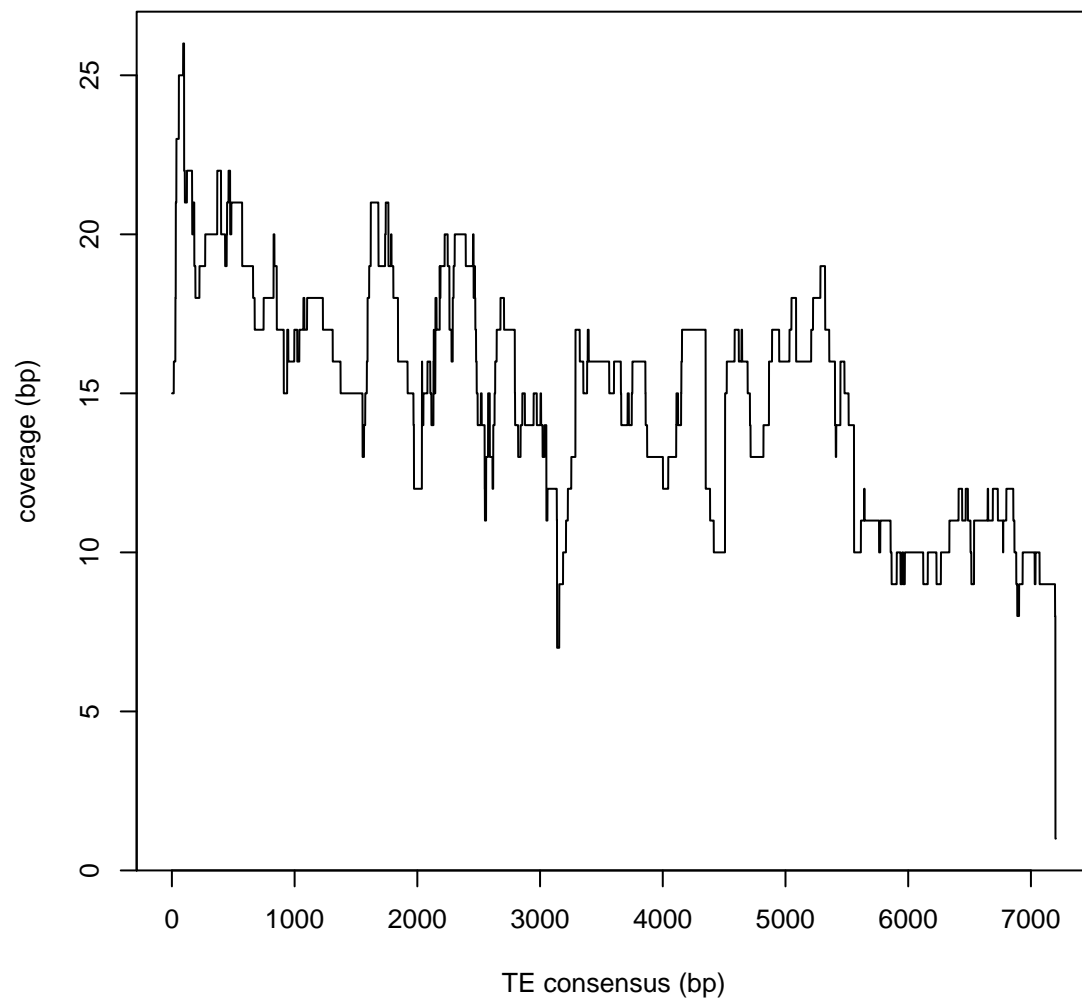


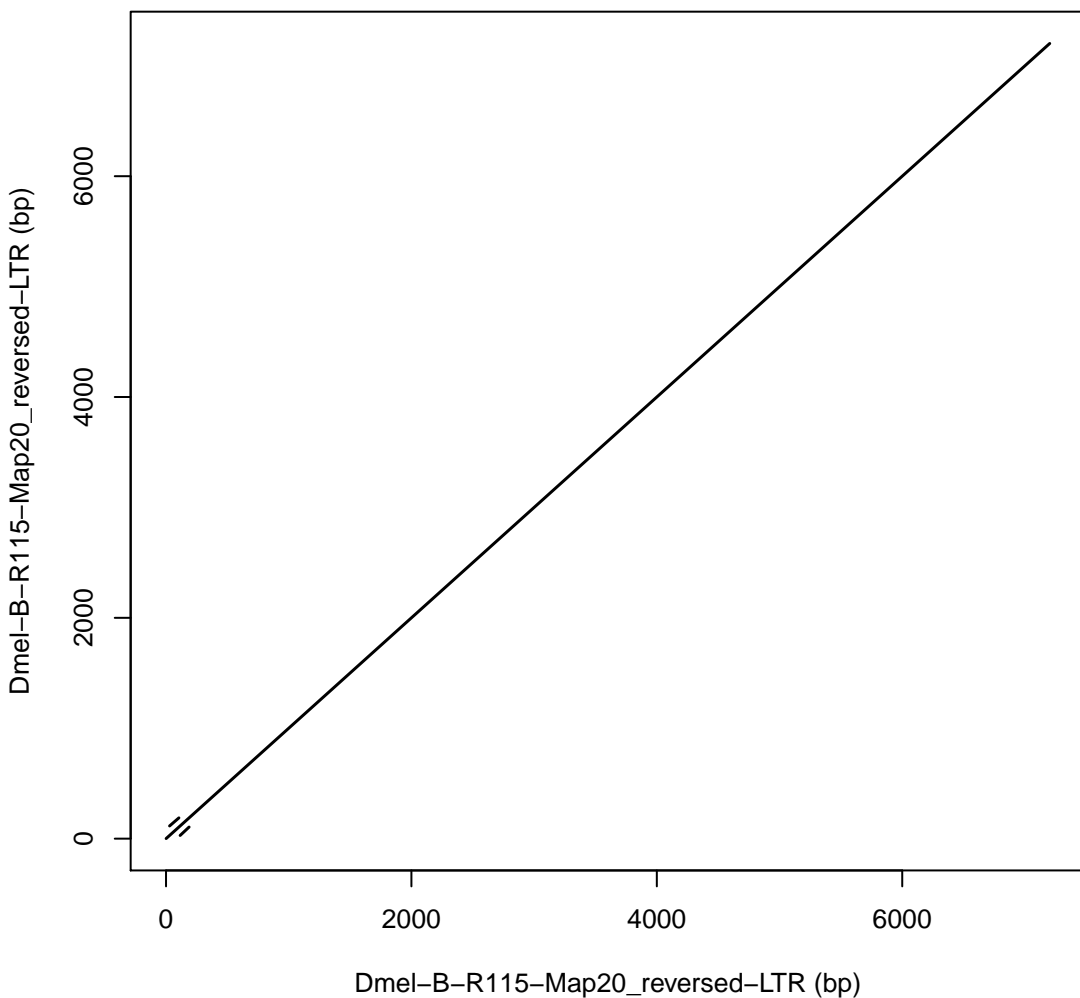
TE: Dmel-B-R115-Map20_reversed-LTR
consensus size: 7202bp; fragments: 158; full length: 0 (>=6481.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

