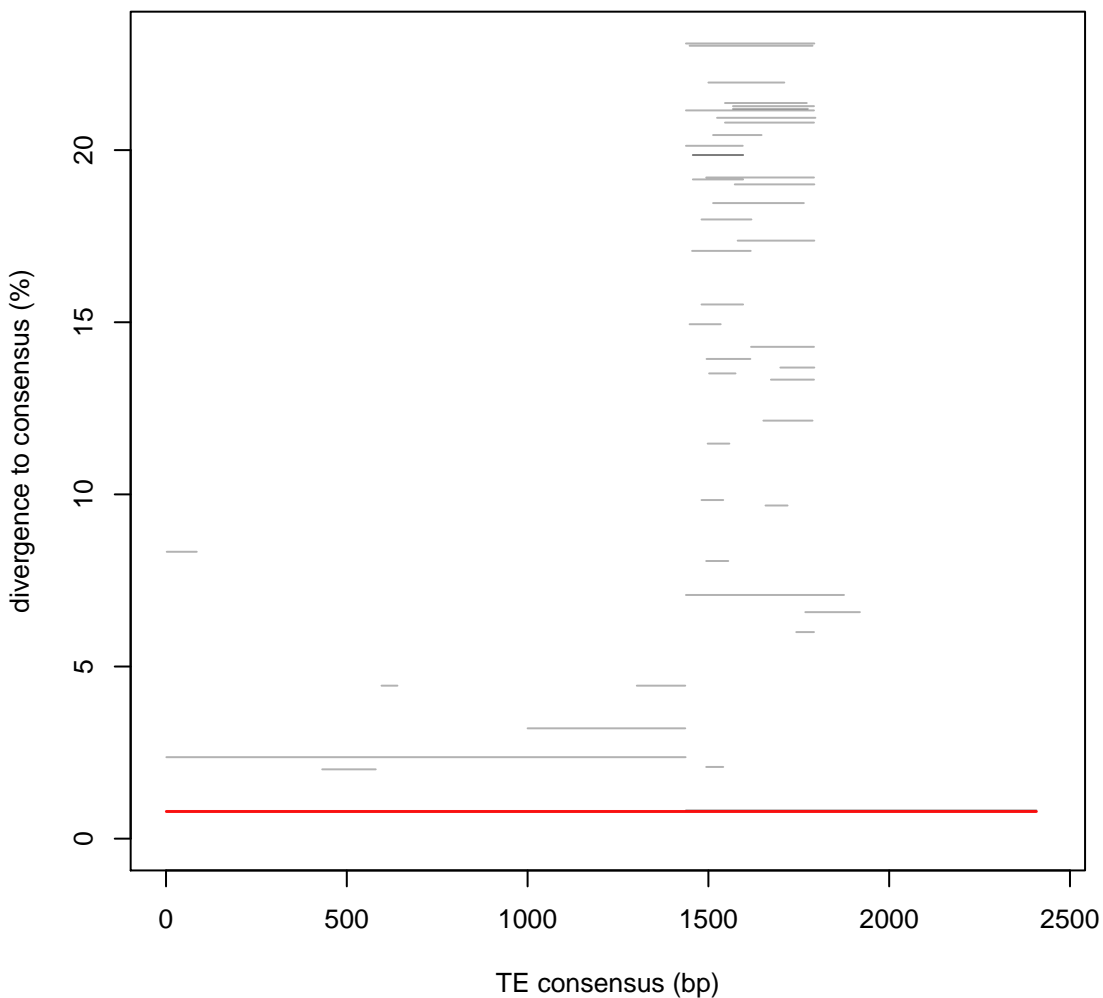
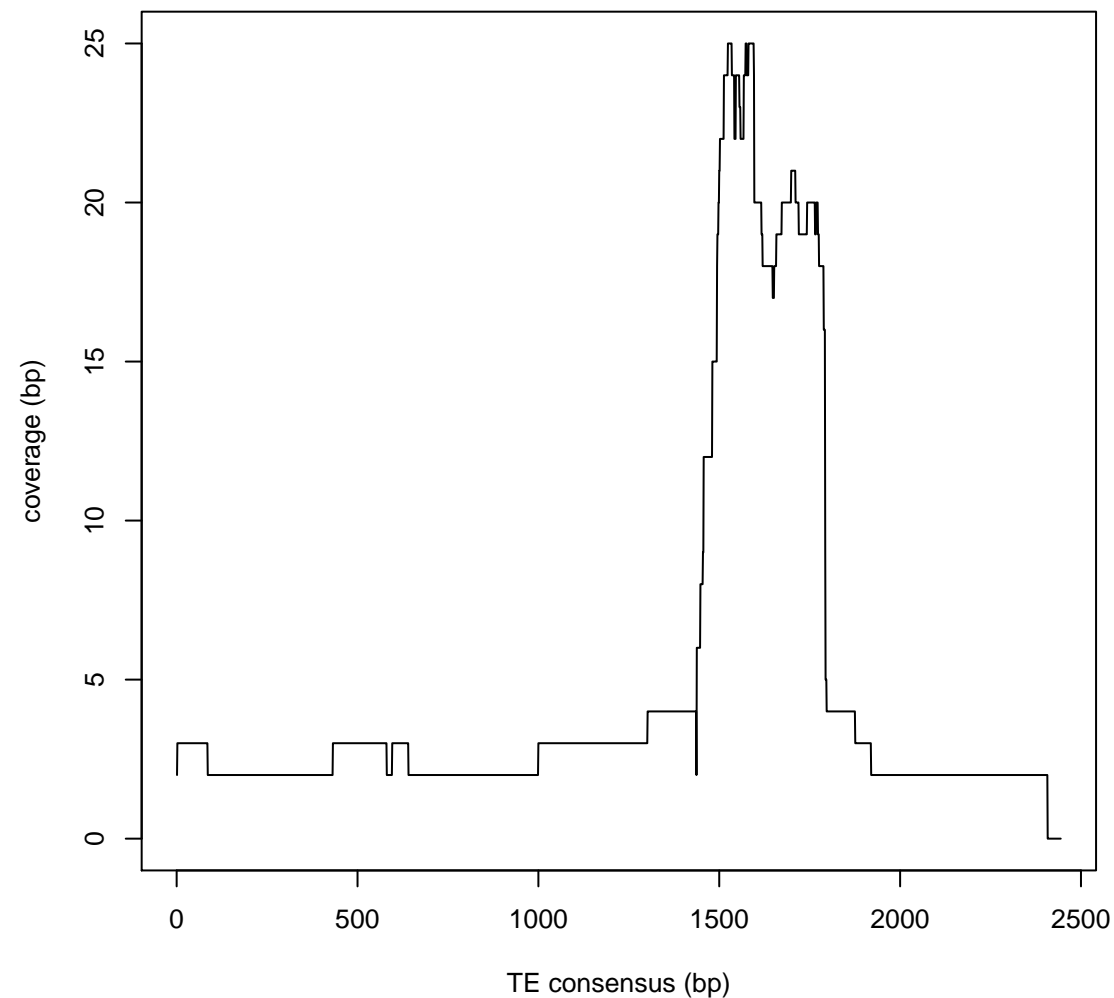


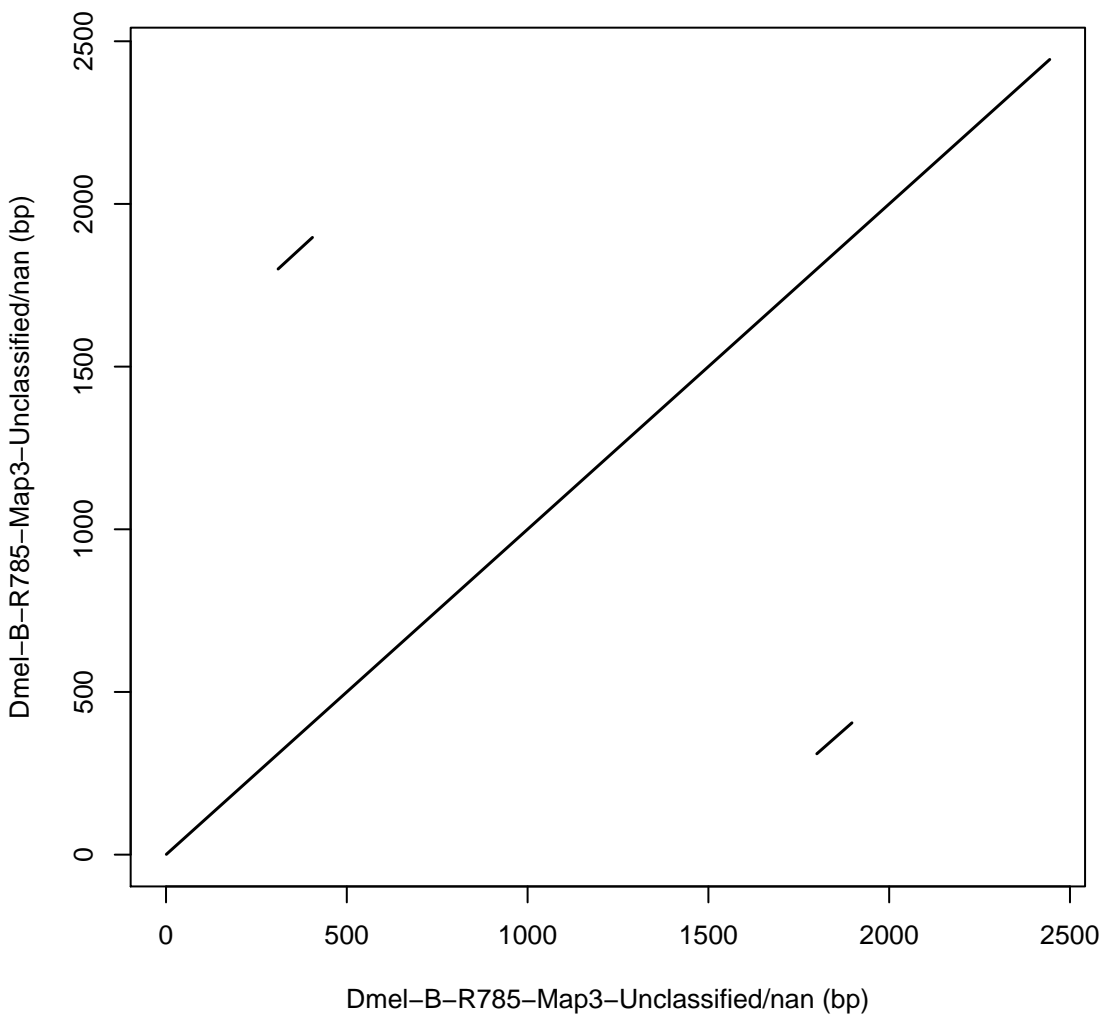
TE: Dmel-B-R785-Map3-Unclassified/nan
consensus size: 2444bp; fragments: 44; full length: 1 ($\geq 2199.6\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

