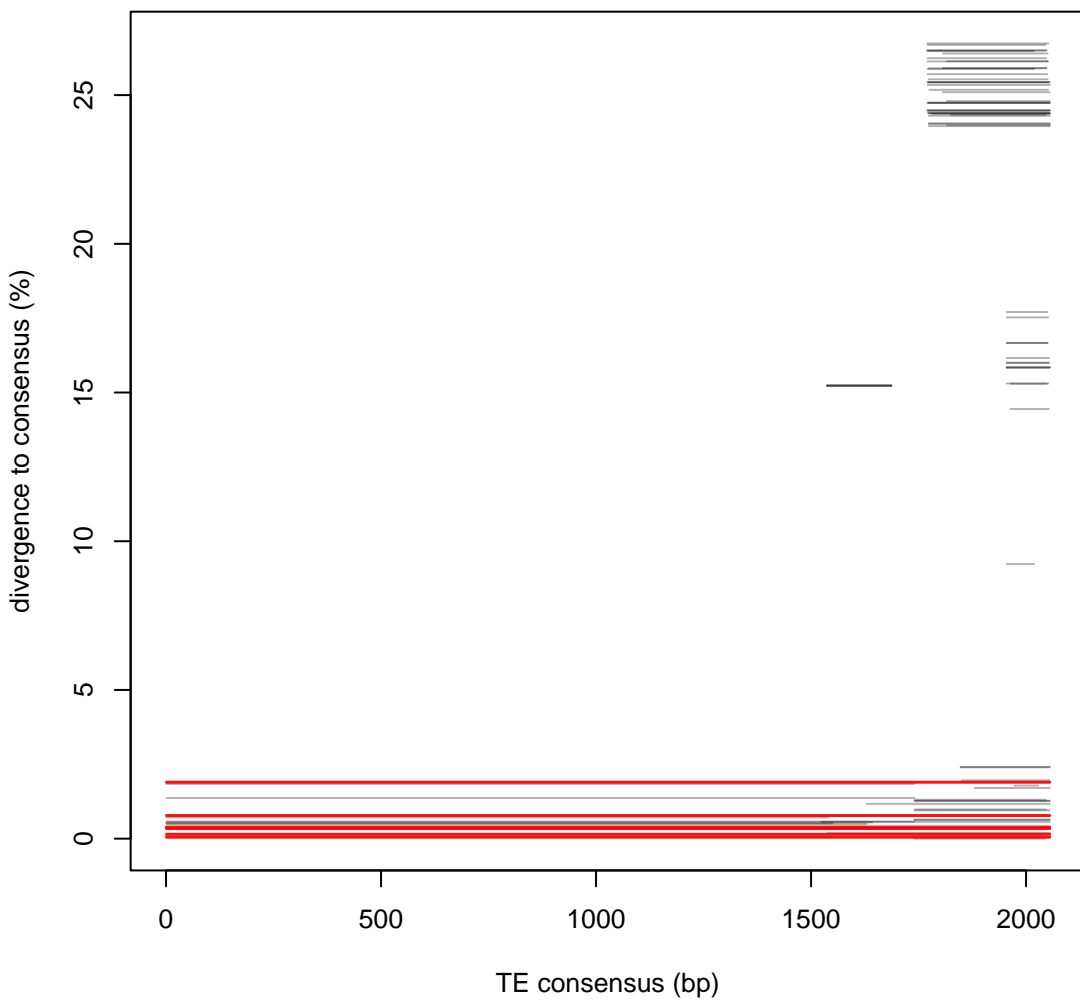
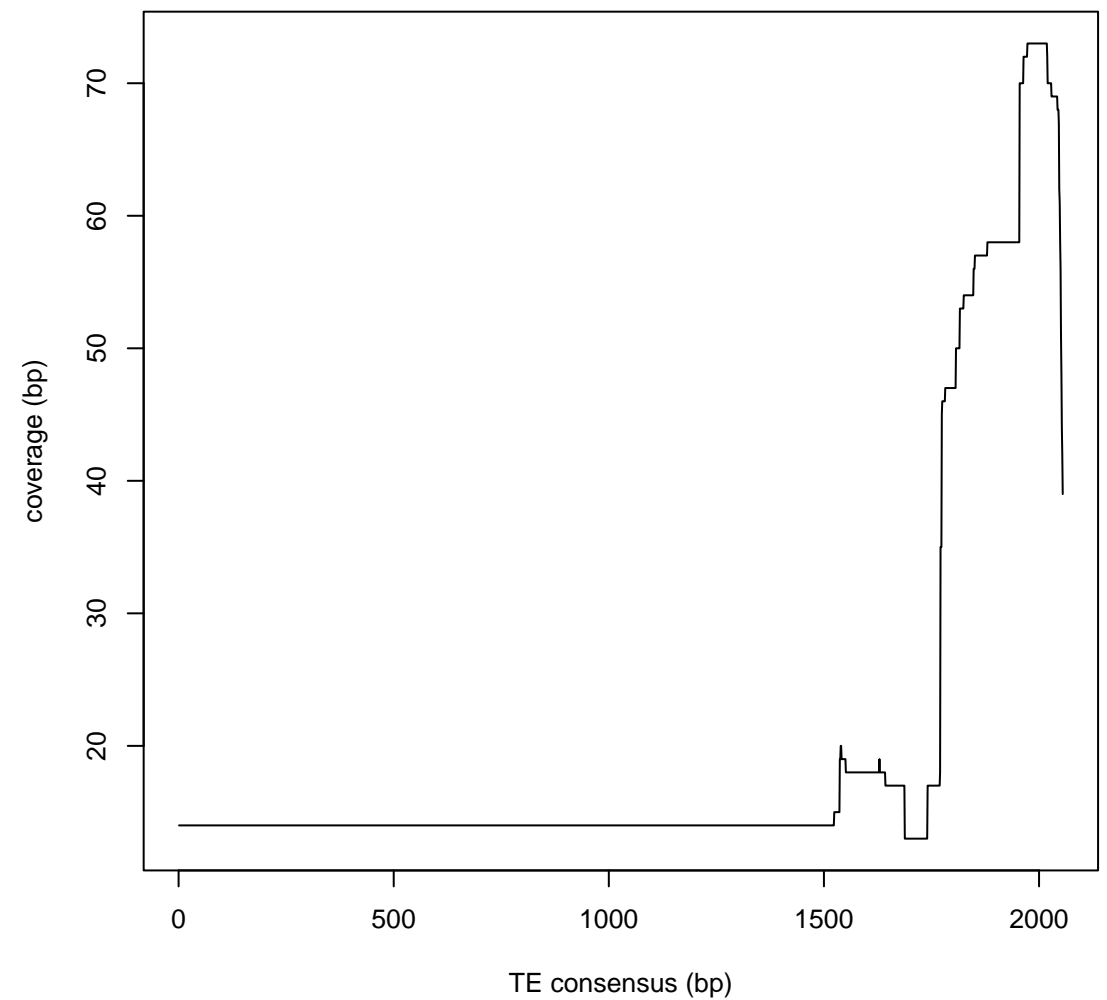


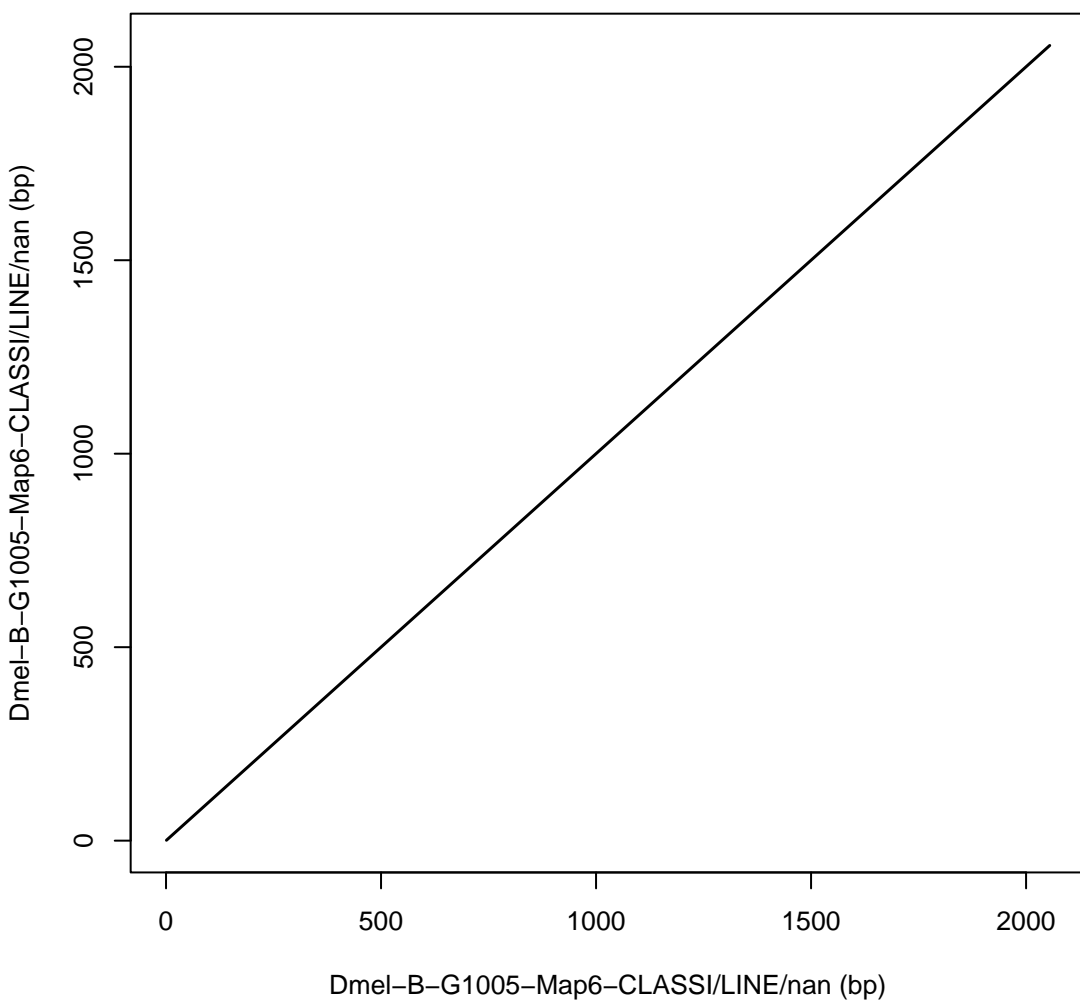
TE: Dmel-B-G1005-Map6-CLASSI/LINE/nan
consensus size: 2055bp; fragments: 85; full length: 6 ($\geq 1849.5\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

