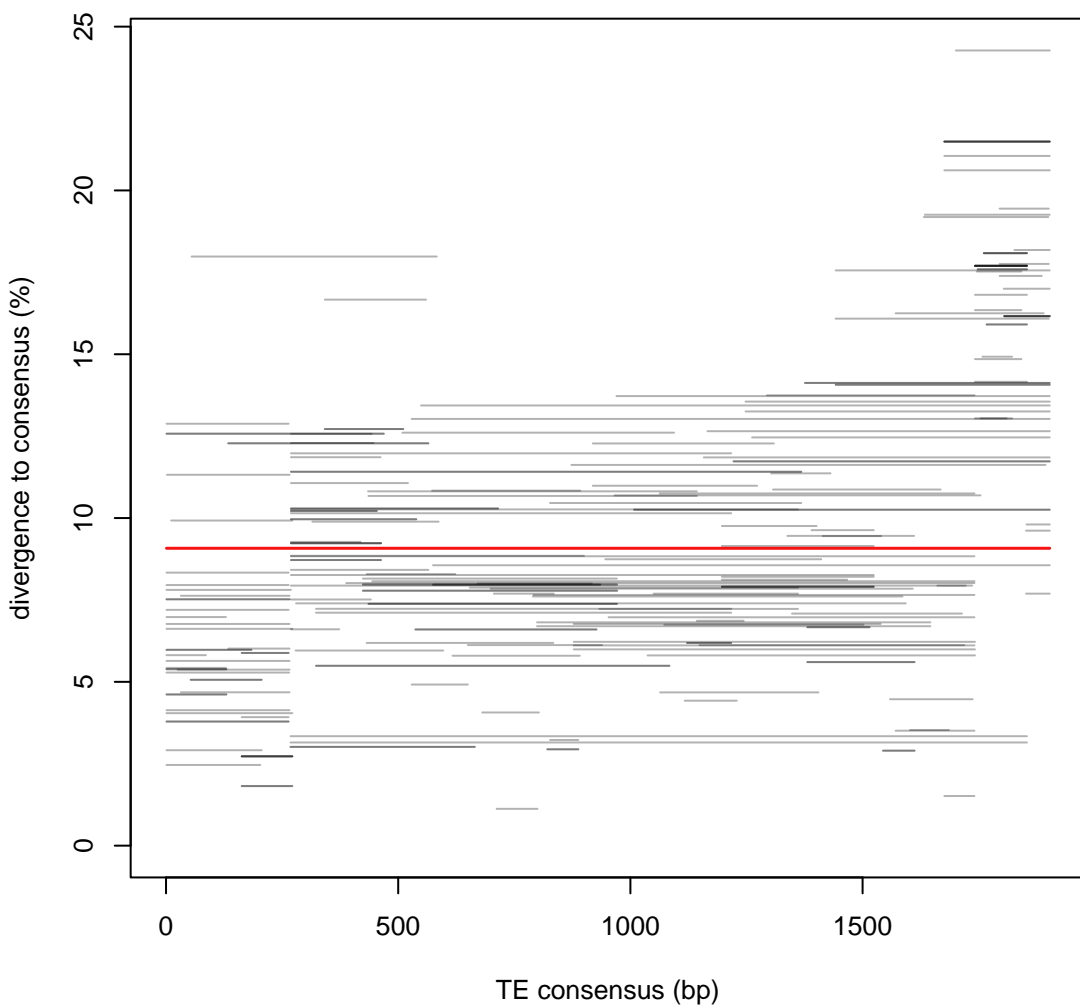
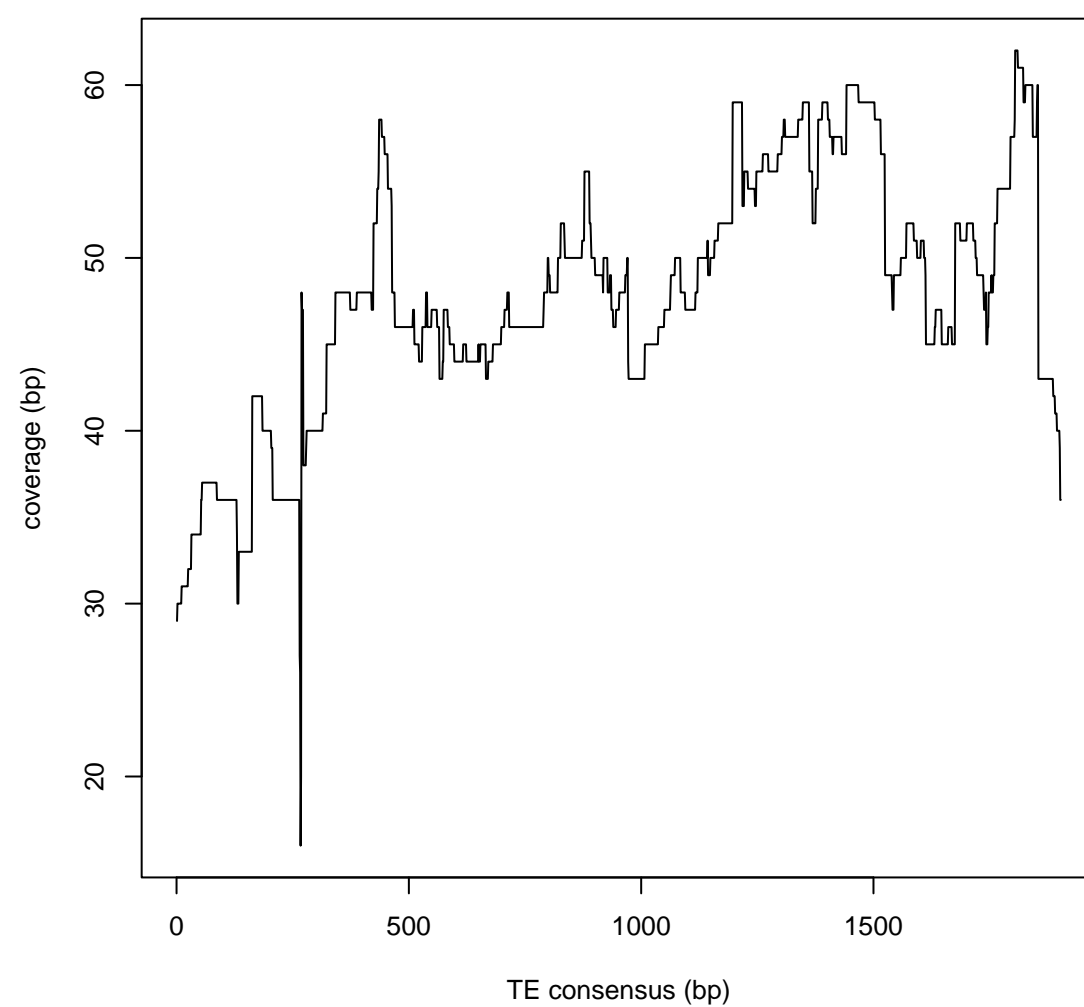


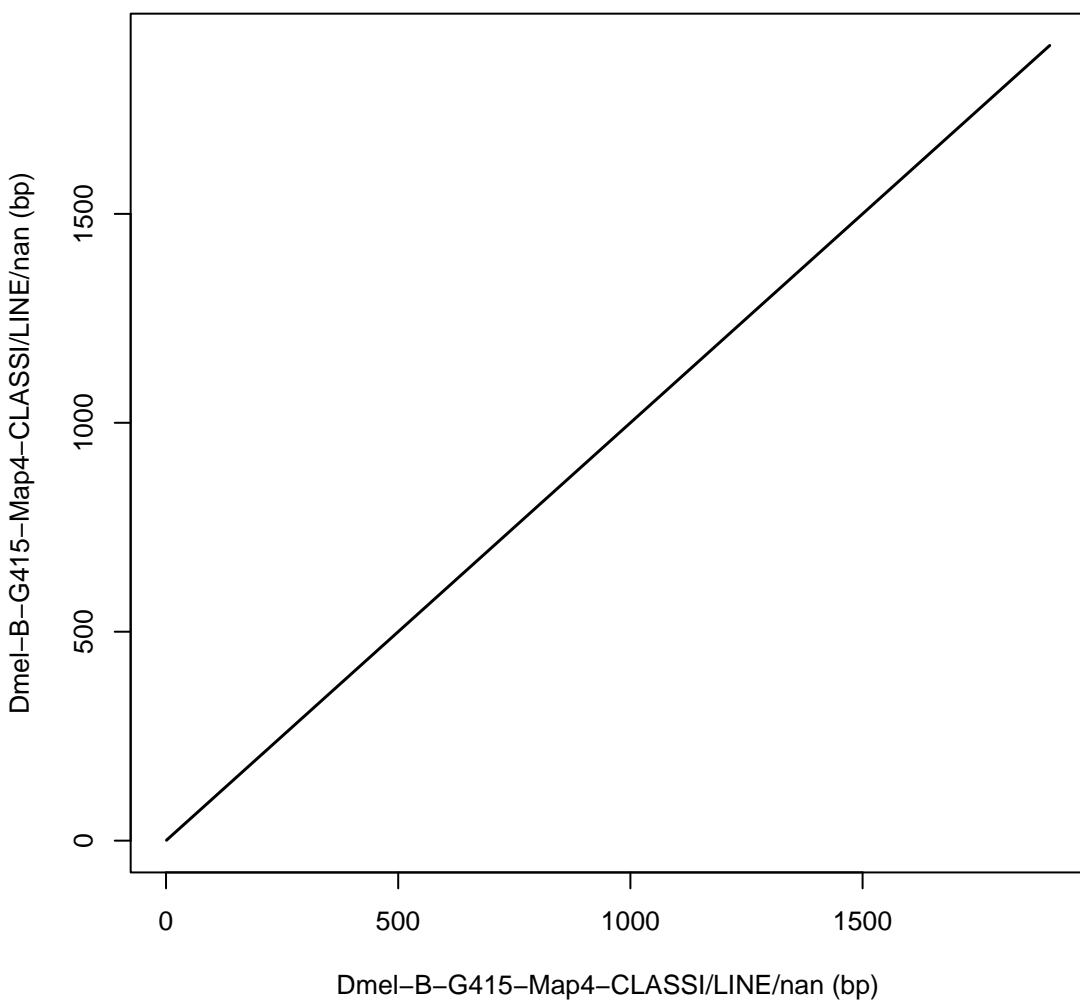
TE: Dmel-B-G415-Map4-CLASSI/LINE/nan
consensus size: 1903bp; fragments: 235; full length: 1 (>=1712.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

