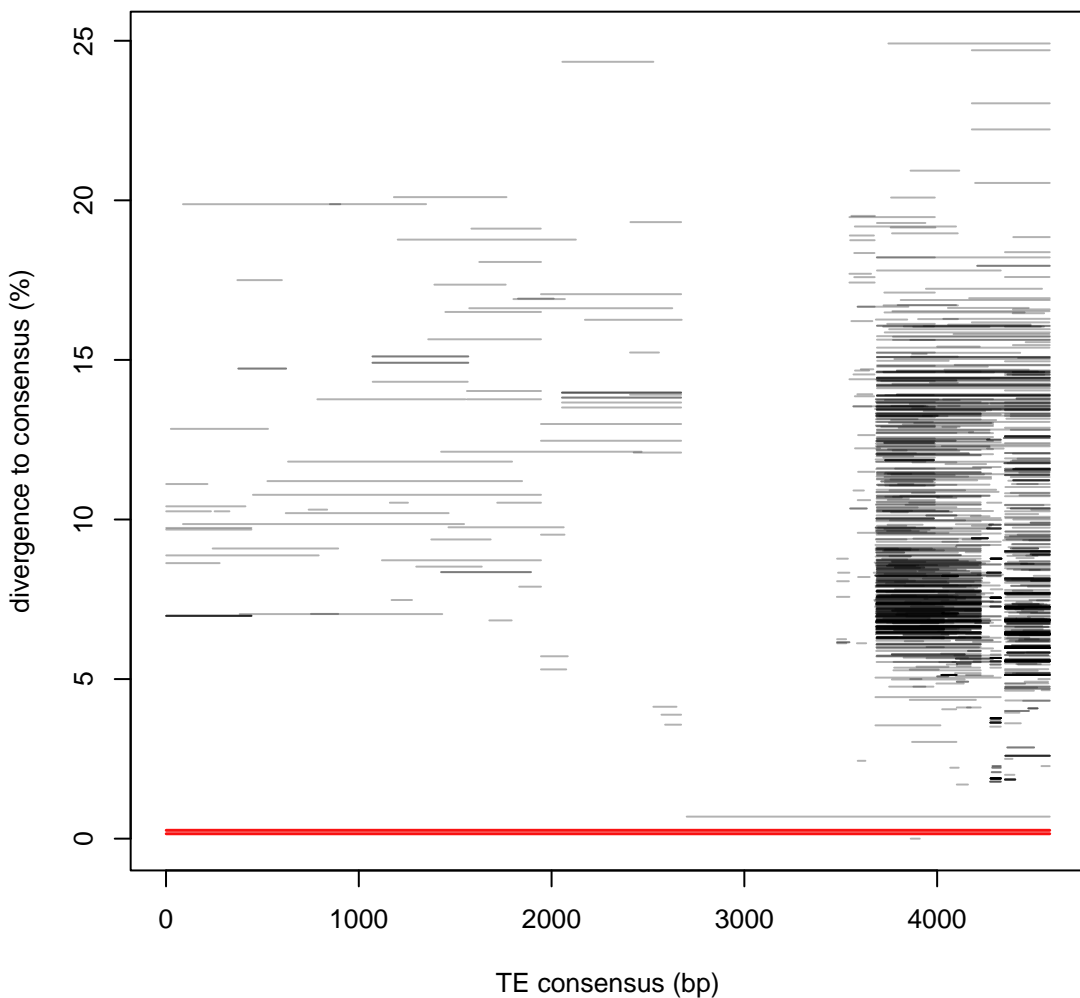
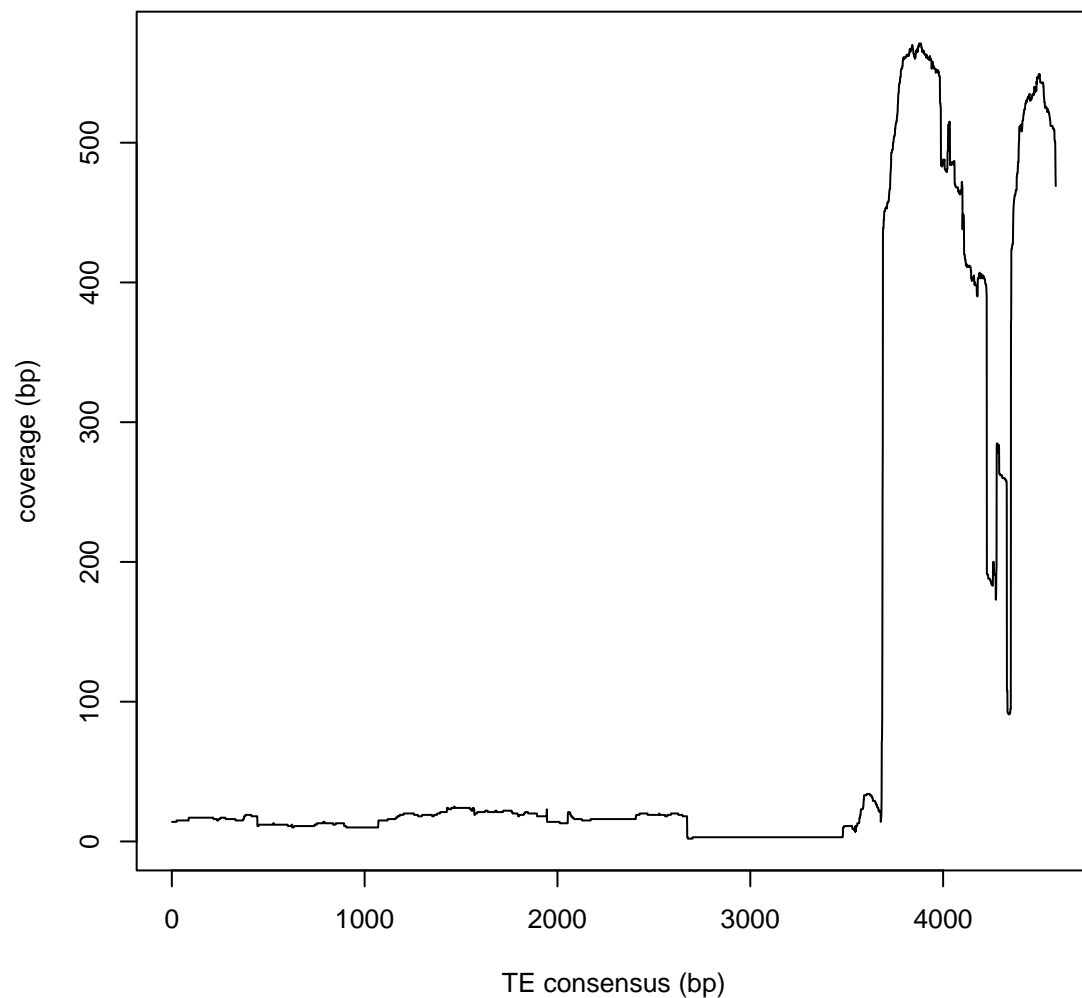


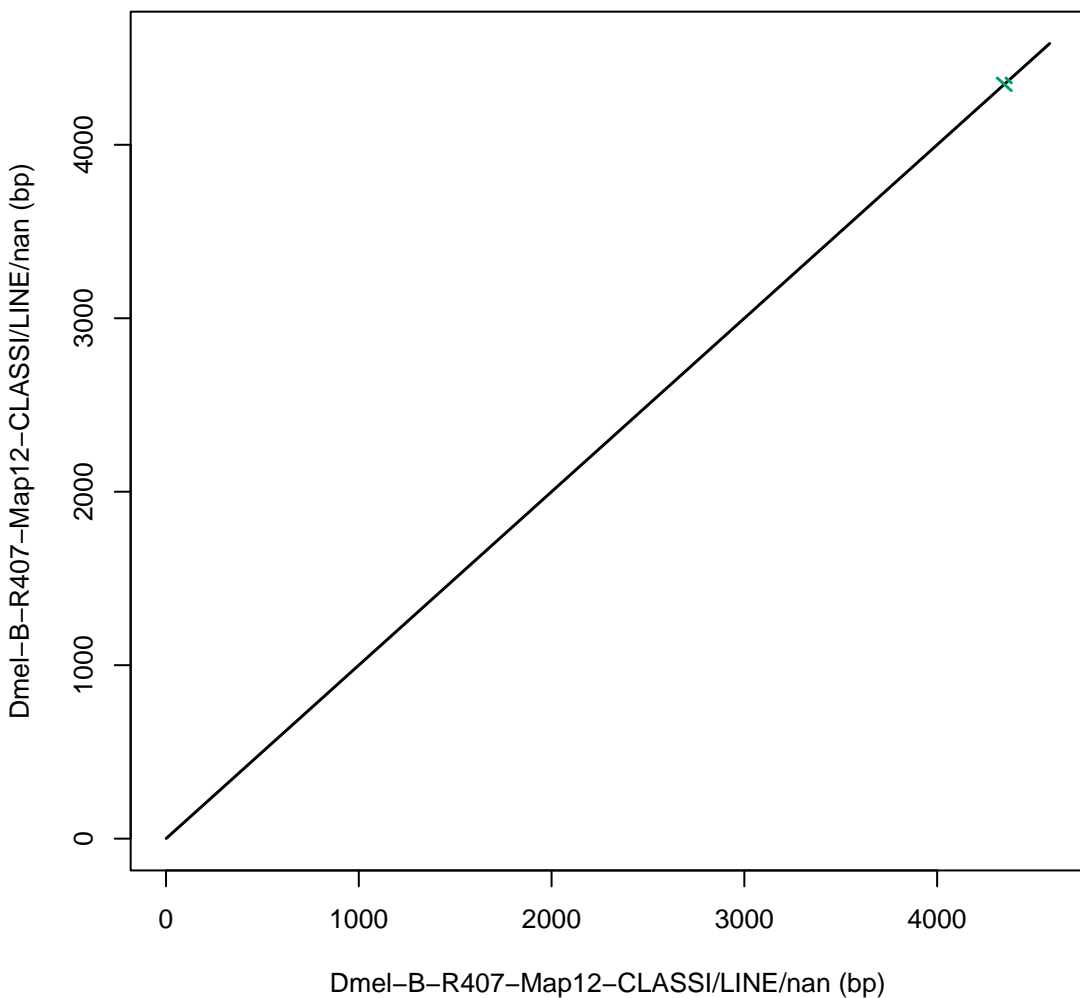
TE: Dmel-B-R407-Map12-CLASSI/LINE/nan
consensus size: 4584bp; fragments: 1487; full length: 2 (≥ 4125.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

