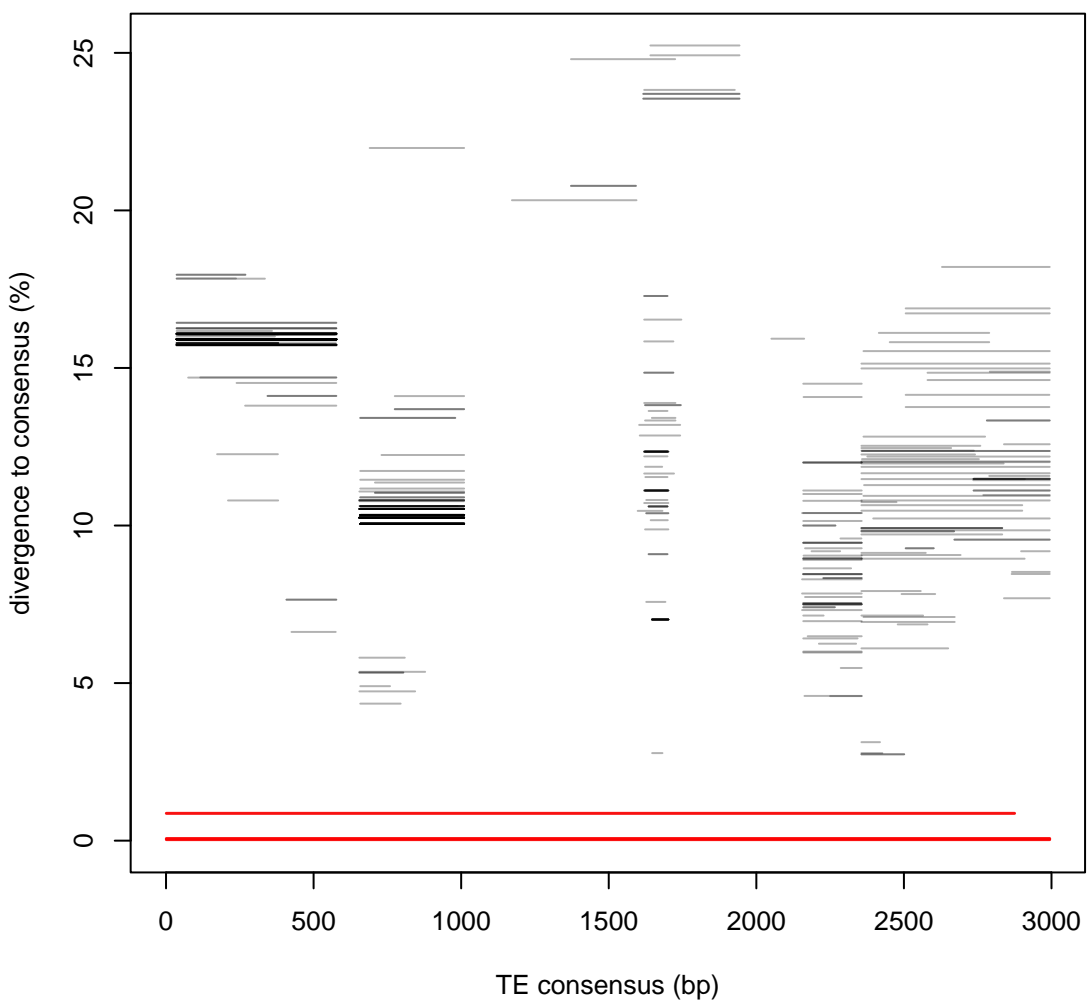
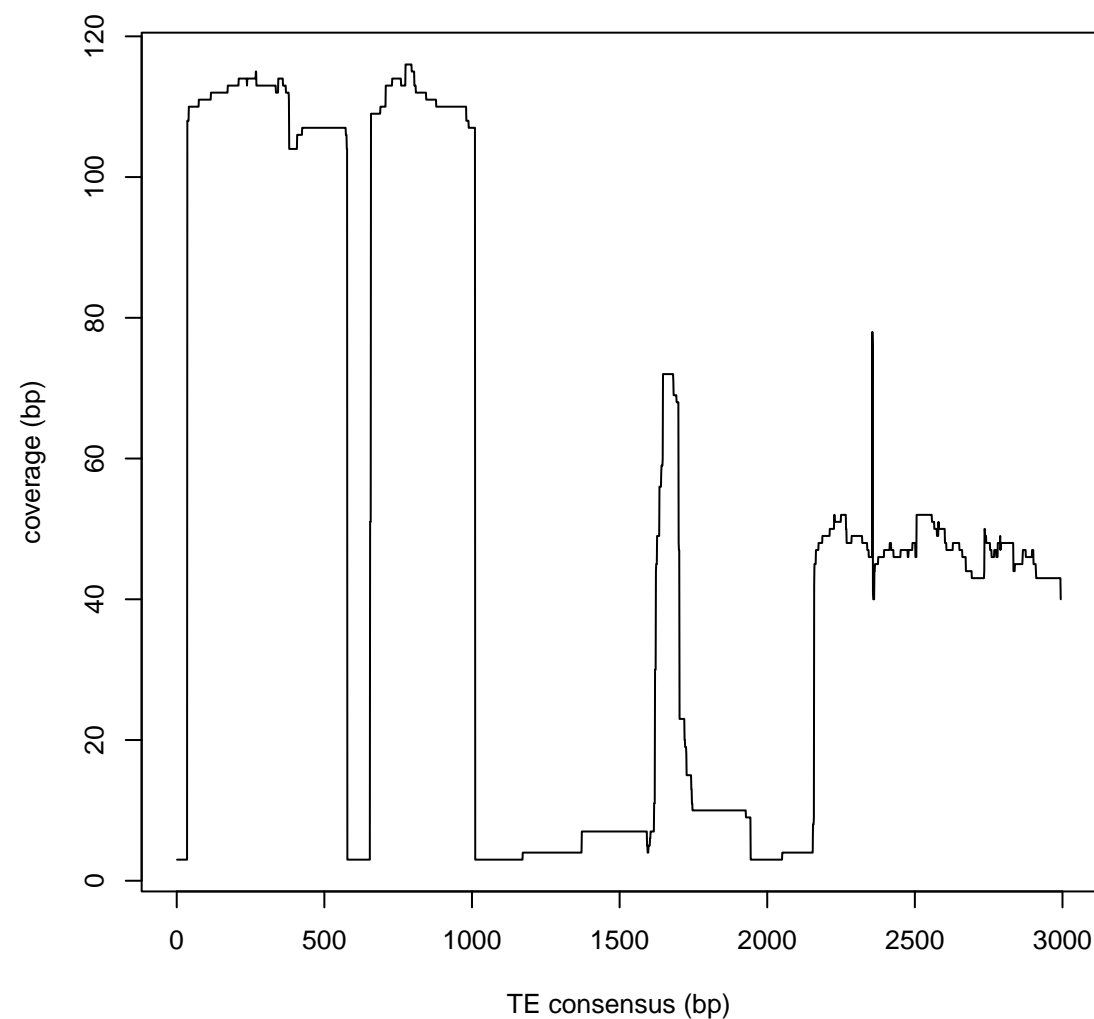


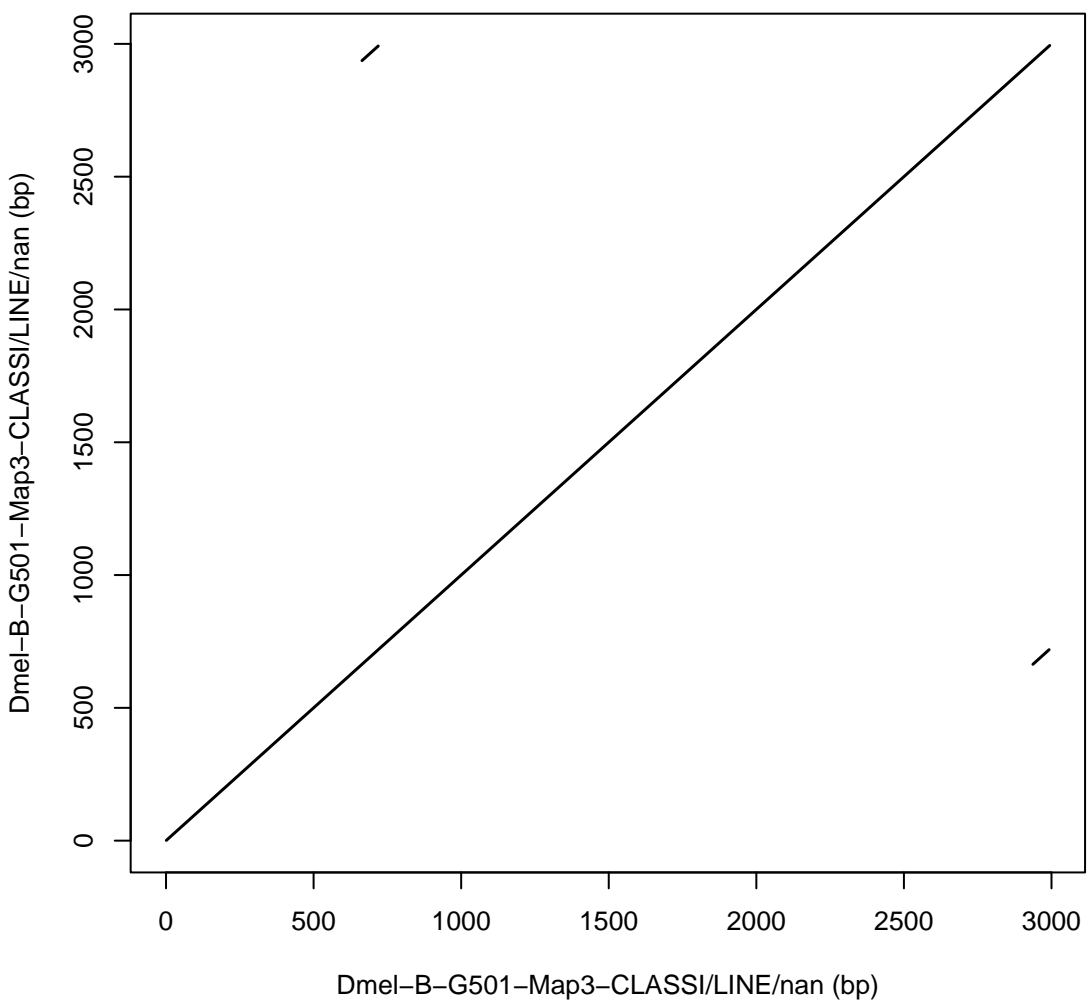
TE: Dmel-B-G501-Map3-CLASSI/LINE/nan
consensus size: 2994bp; fragments: 436; full length: 3 (>=2694.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

