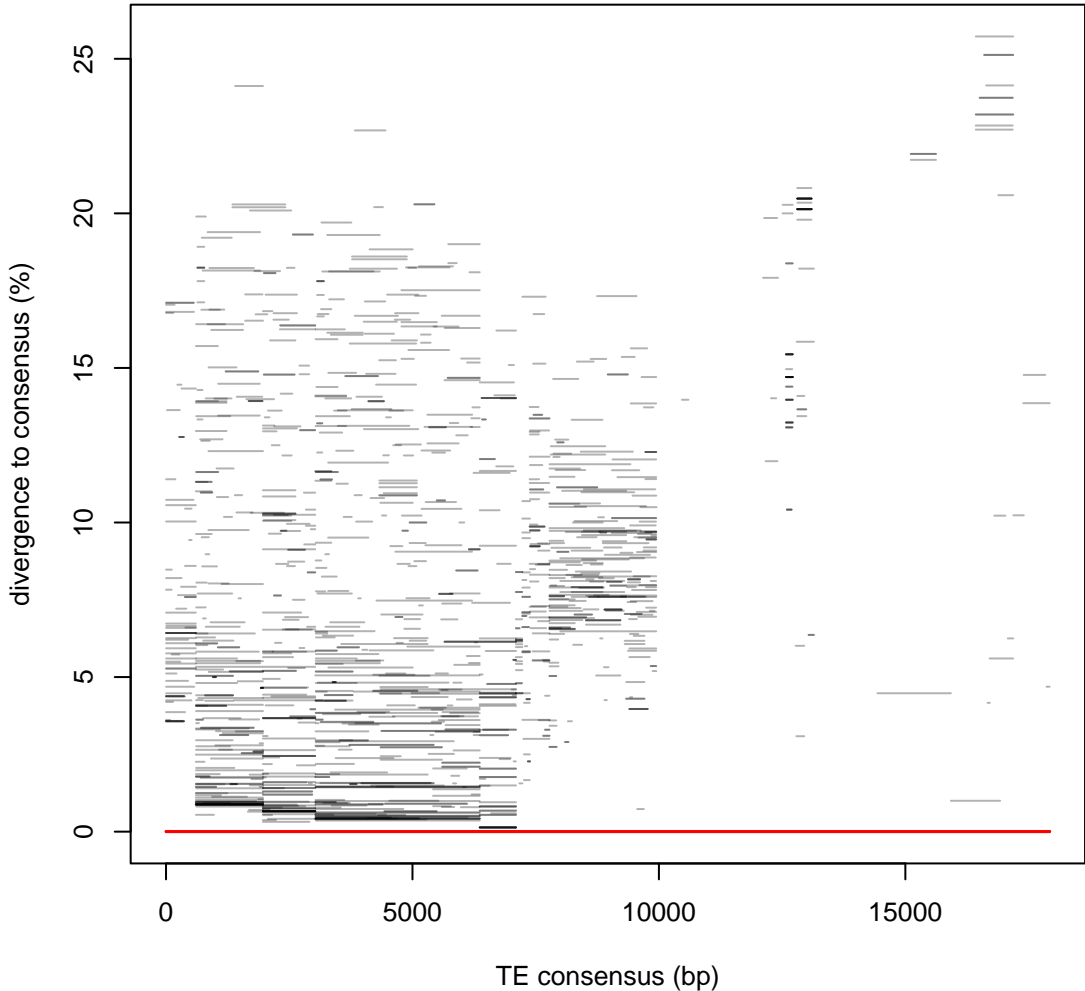
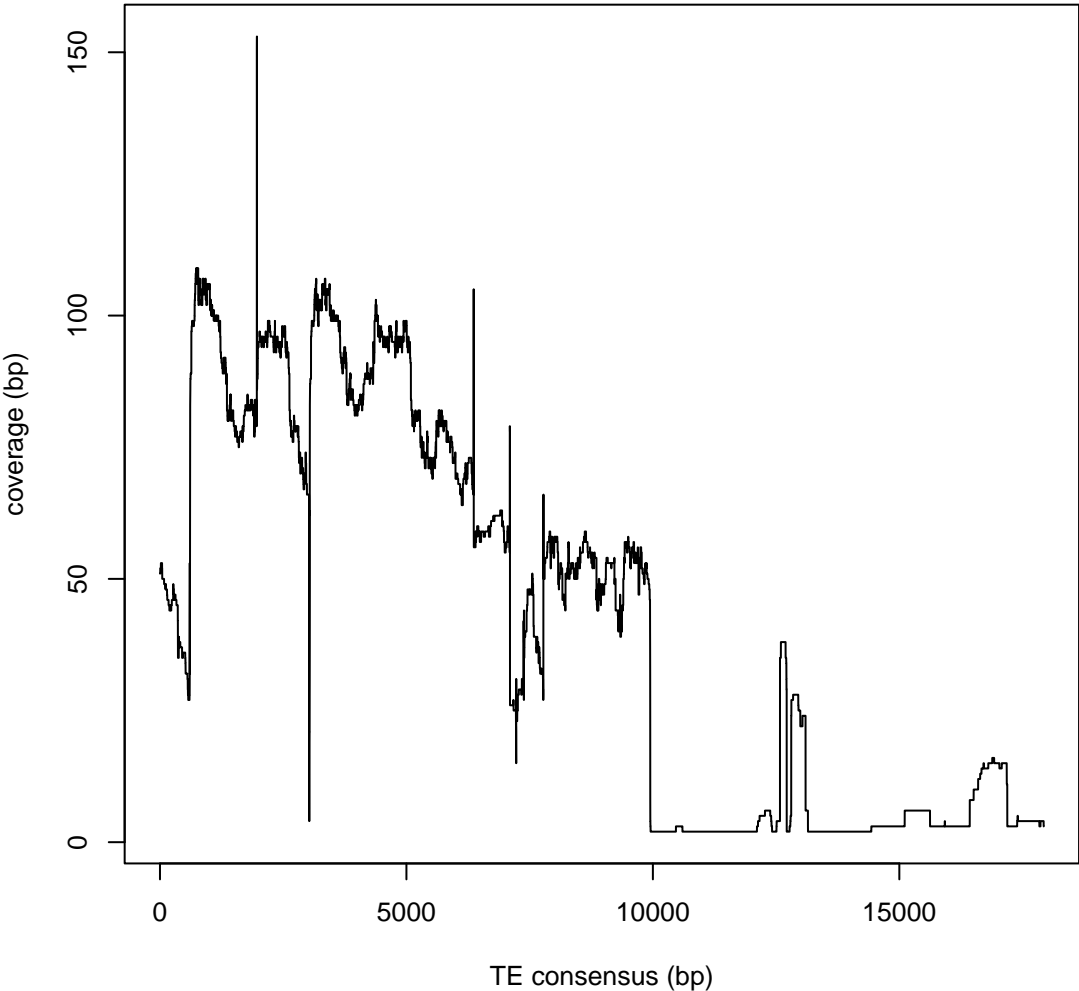


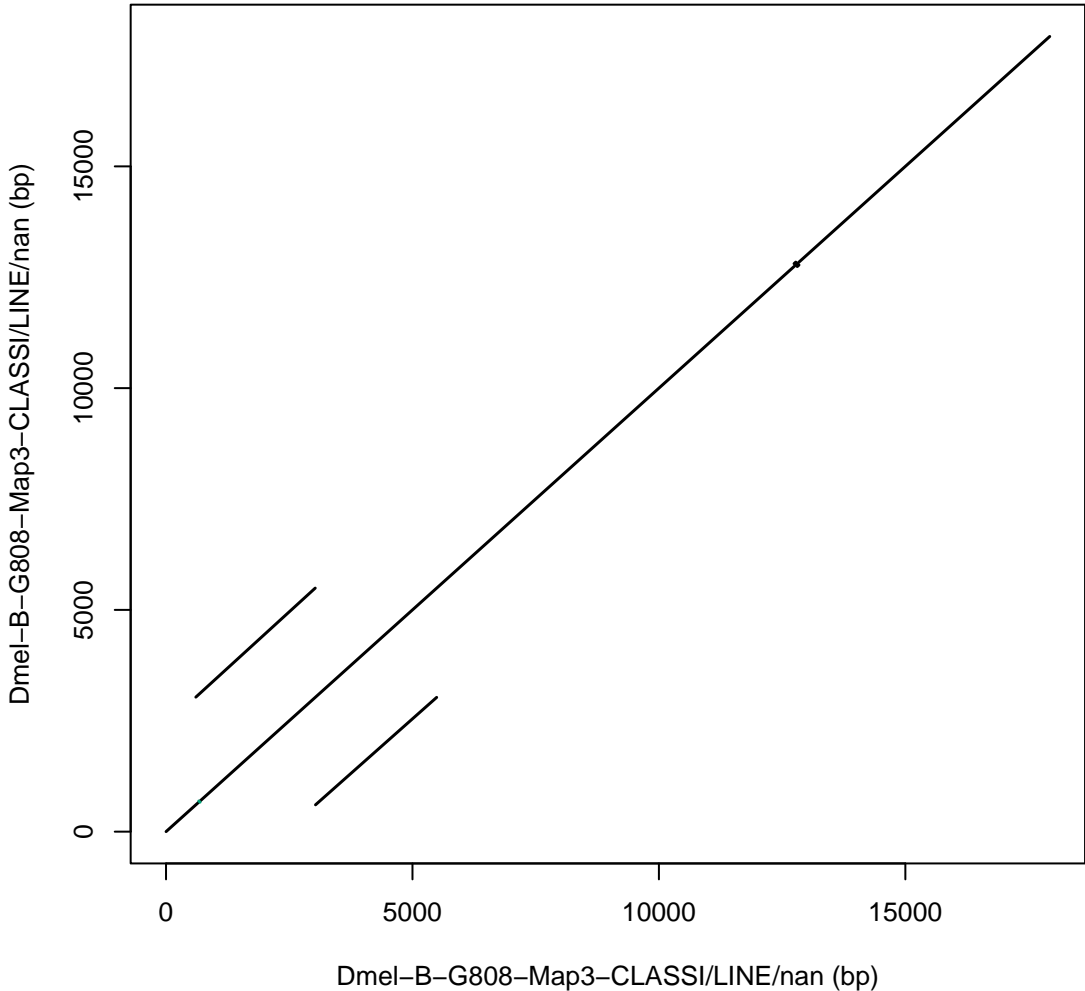
TE: Dmel-B-G808-Map3-CLASSI/LINE/nan
consensus size: 17929bp; fragments: 1274; full length: 2 (>=16136.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

