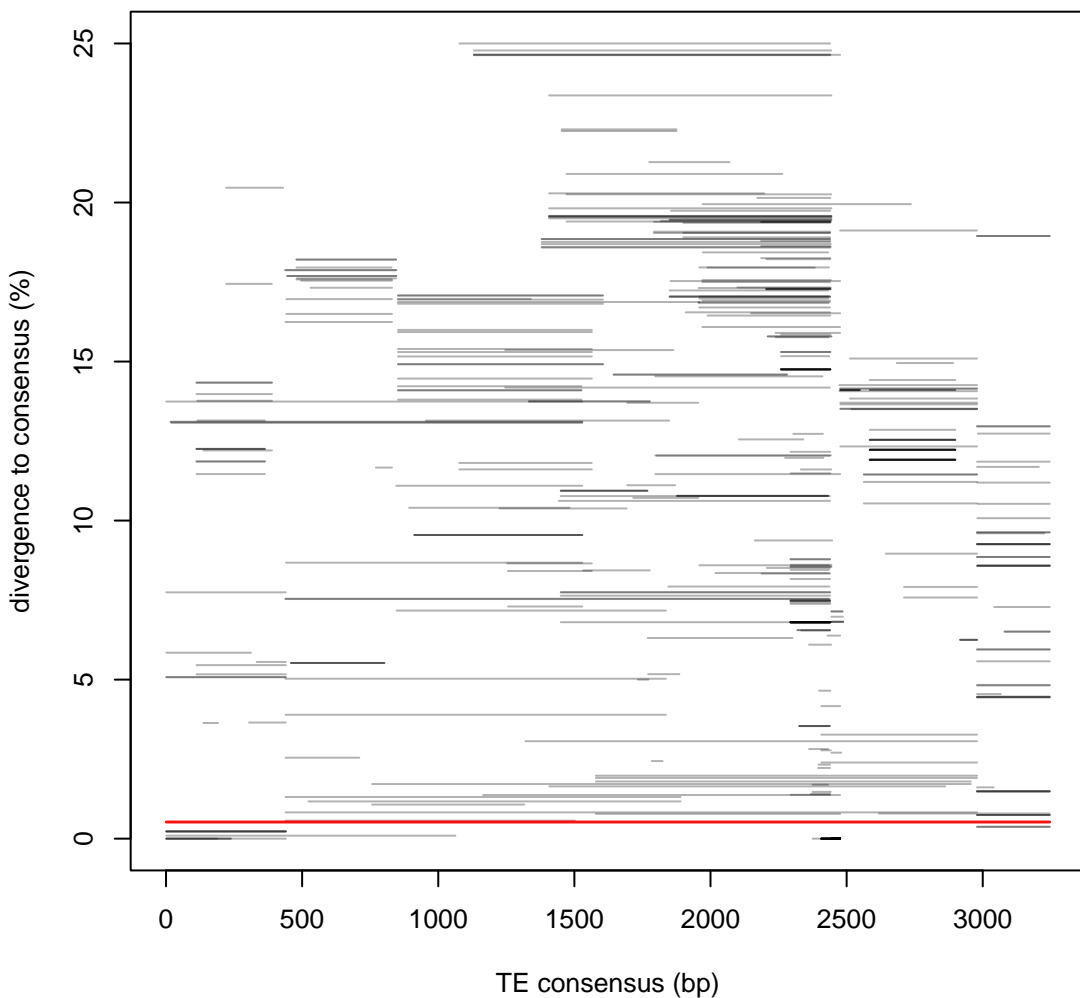
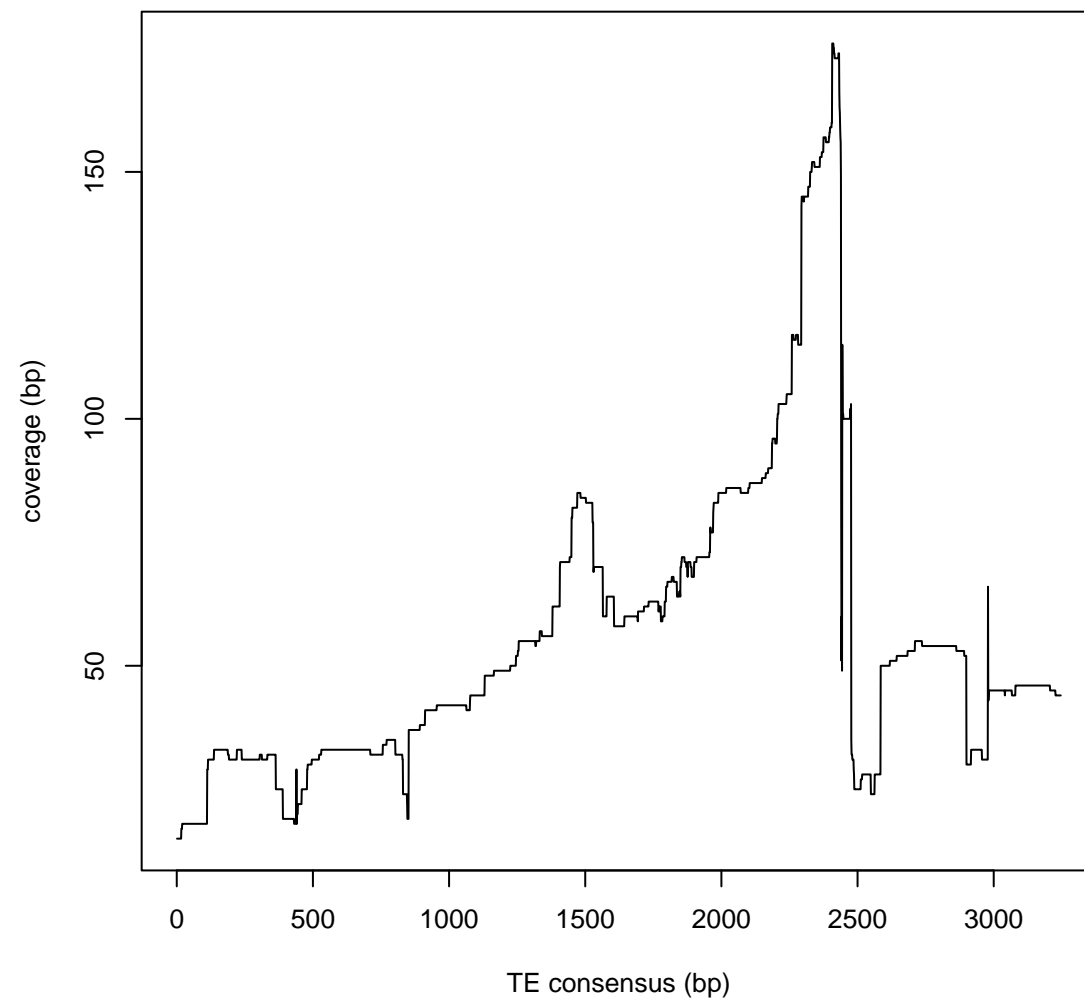


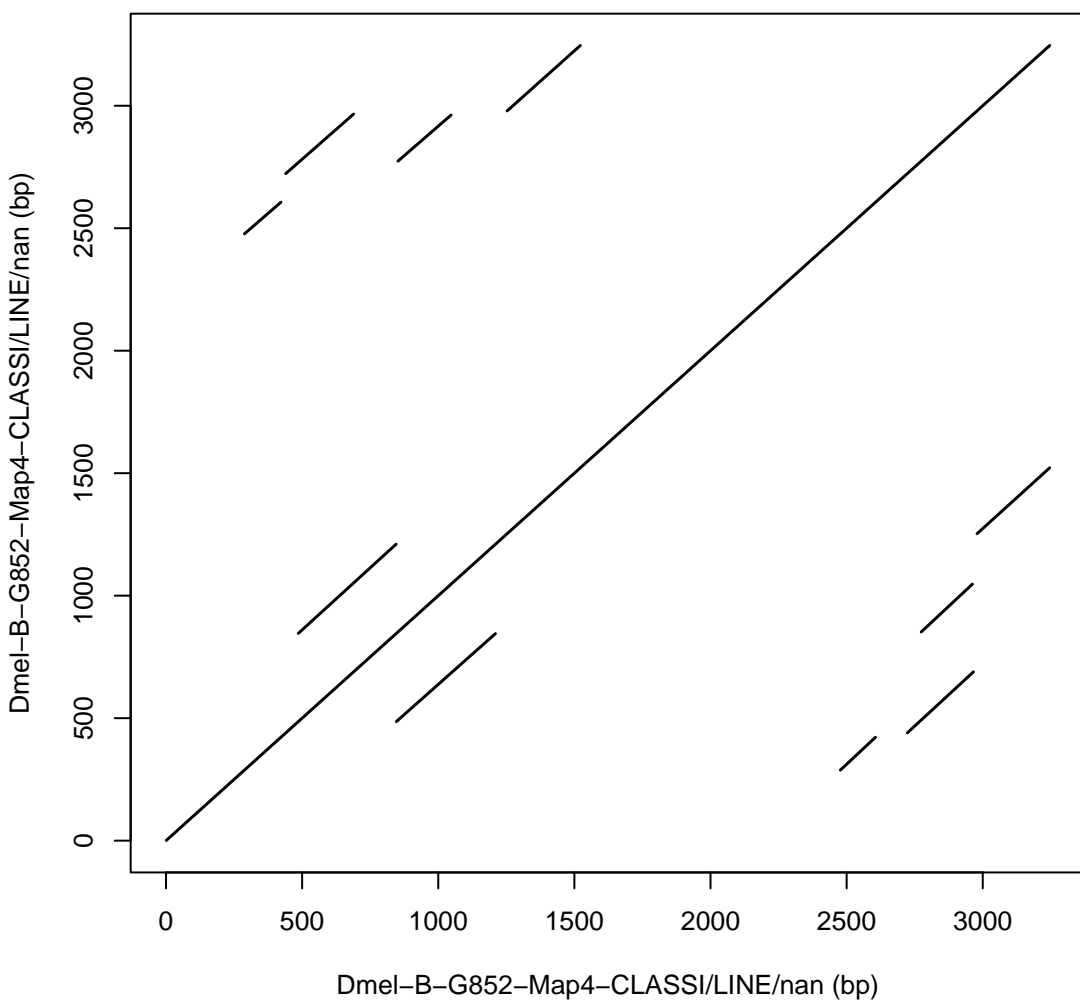
TE: Dmel-B-G852-Map4-CLASSI/LINE/nan
consensus size: 3246bp; fragments: 469; full length: 1 (>=2921.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

