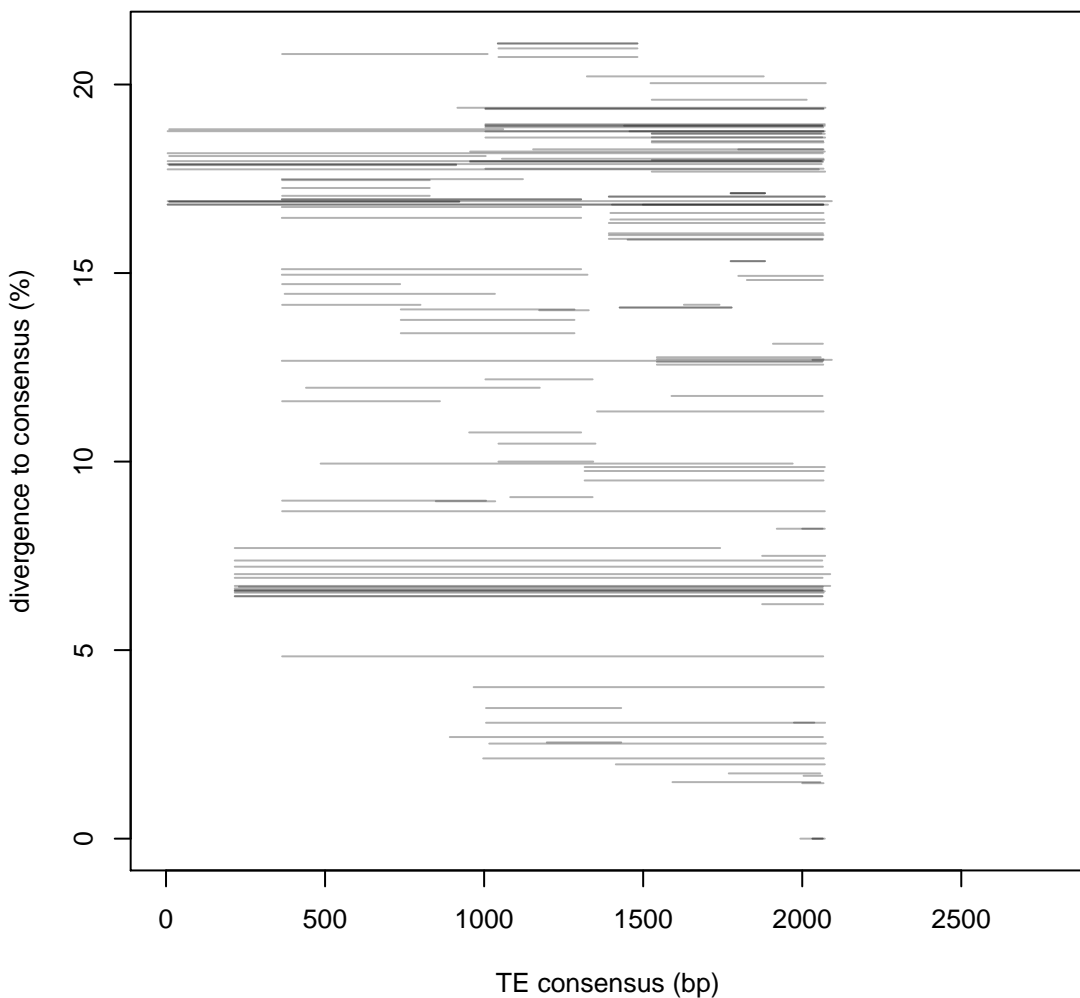
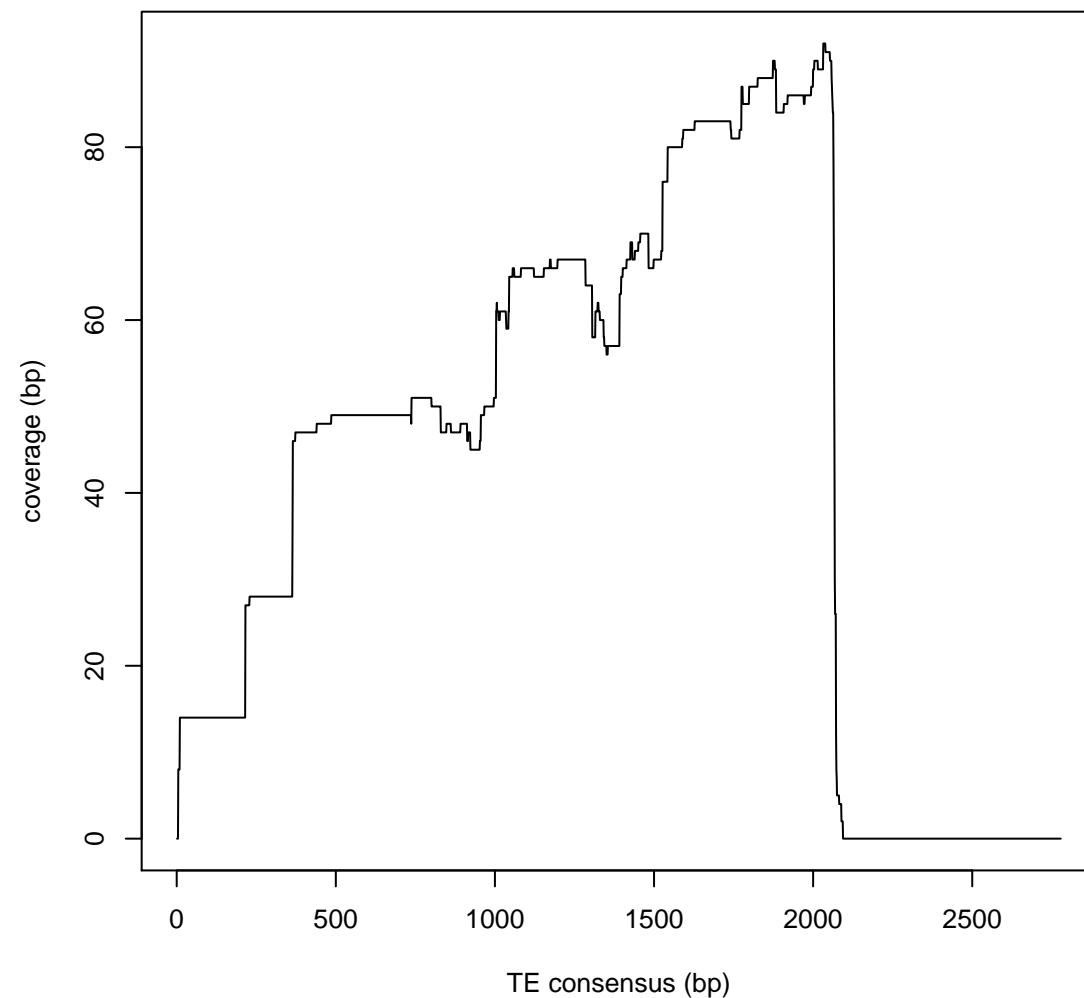


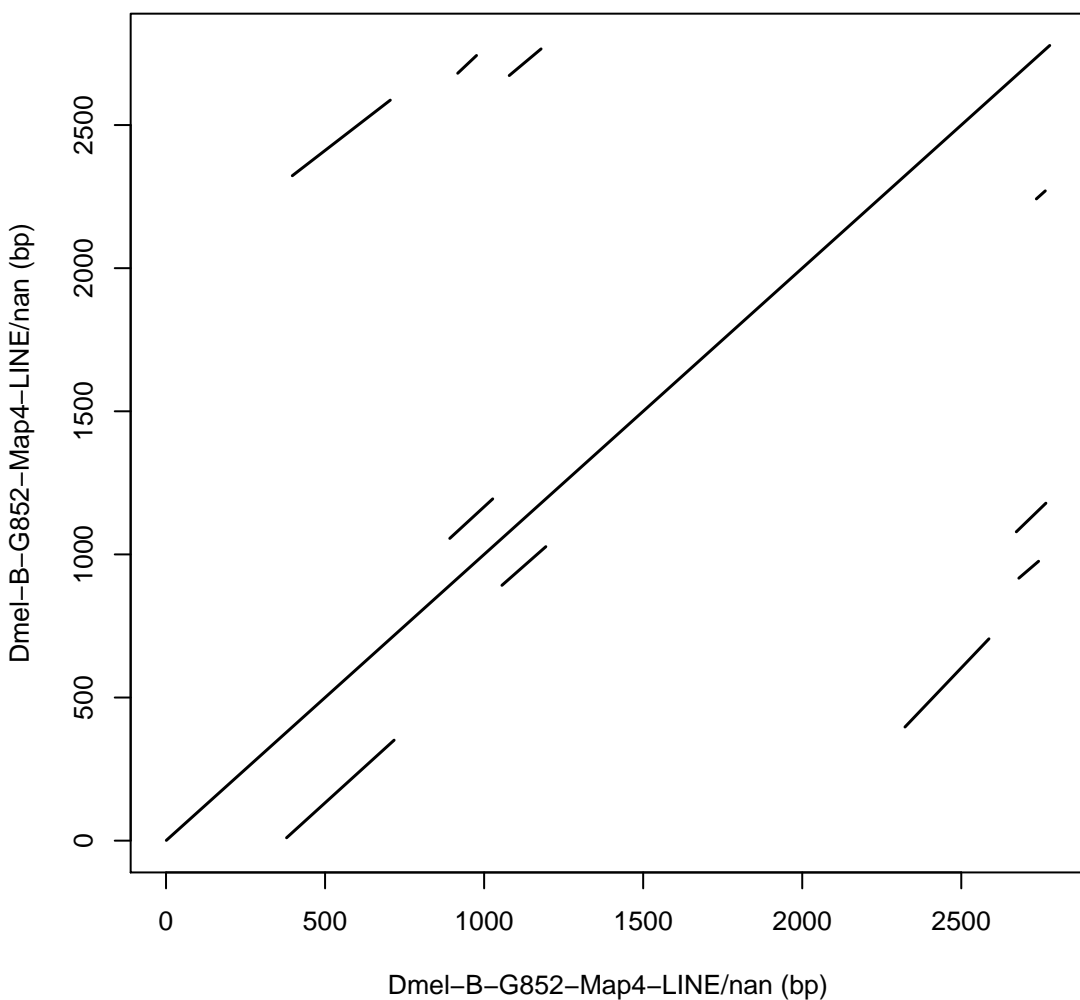
TE: Dmel-B-G852-Map4-LINE/nan  
consensus size: 2778bp; fragments: 143; full length: 0 (>=2500.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

