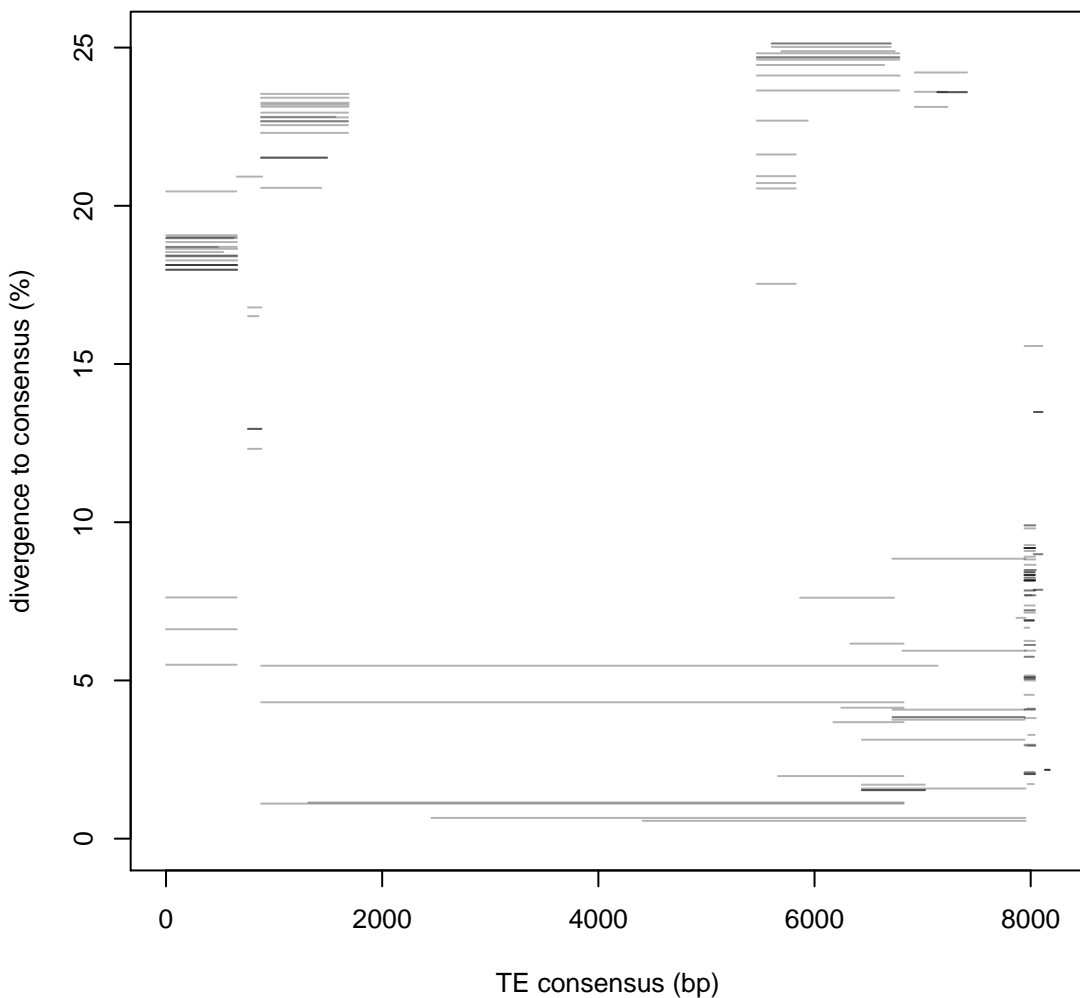
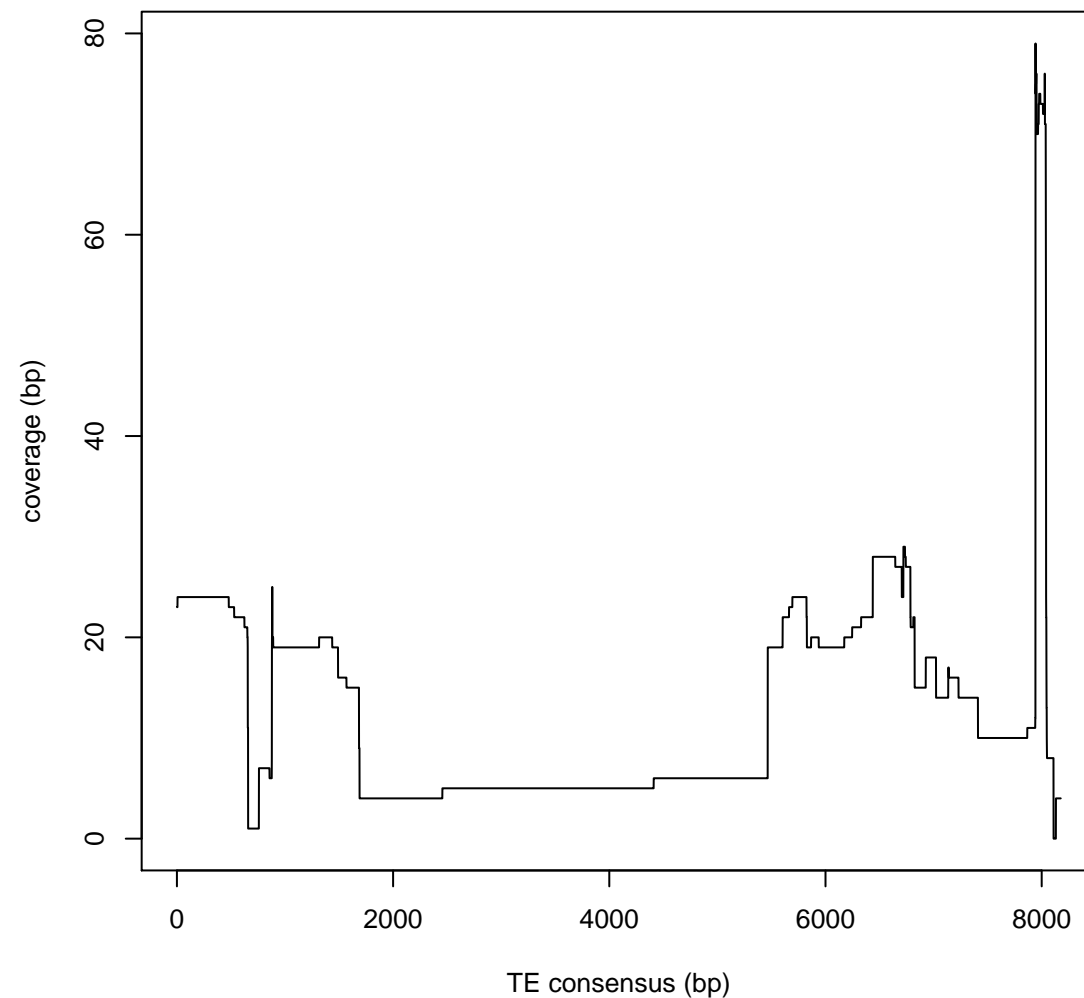


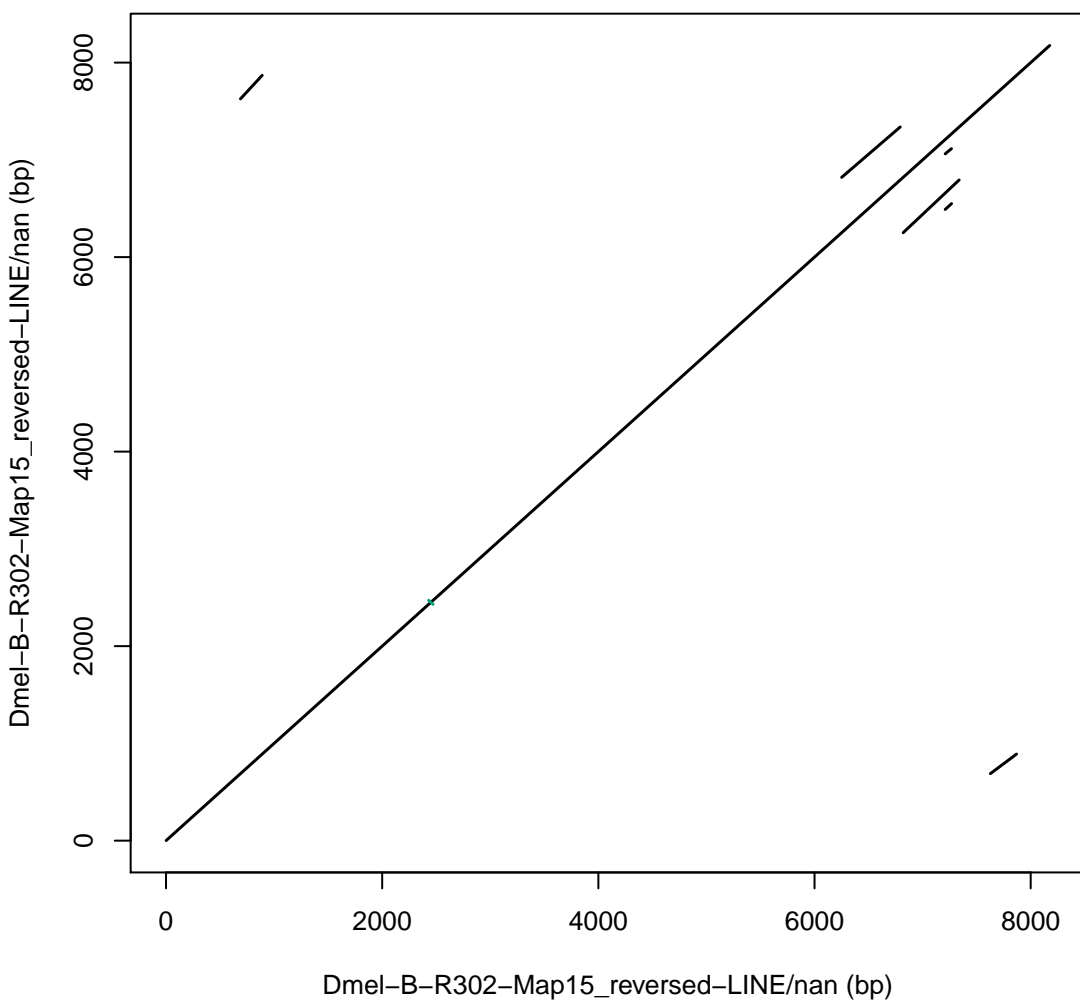
TE: Dmel-B-R302-Map15_reversed-LINE/nan
consensus size: 8176bp; fragments: 179; full length: 0 (>=7358.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

