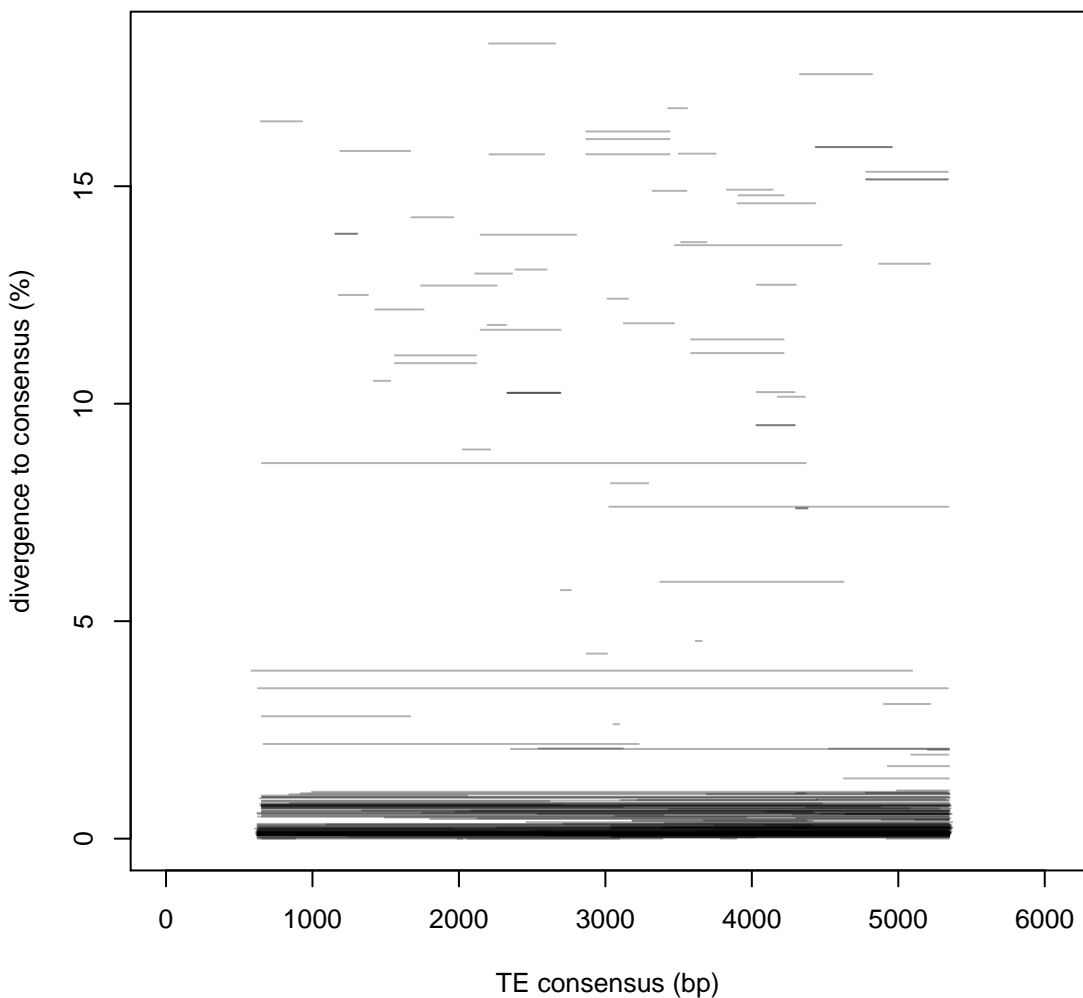
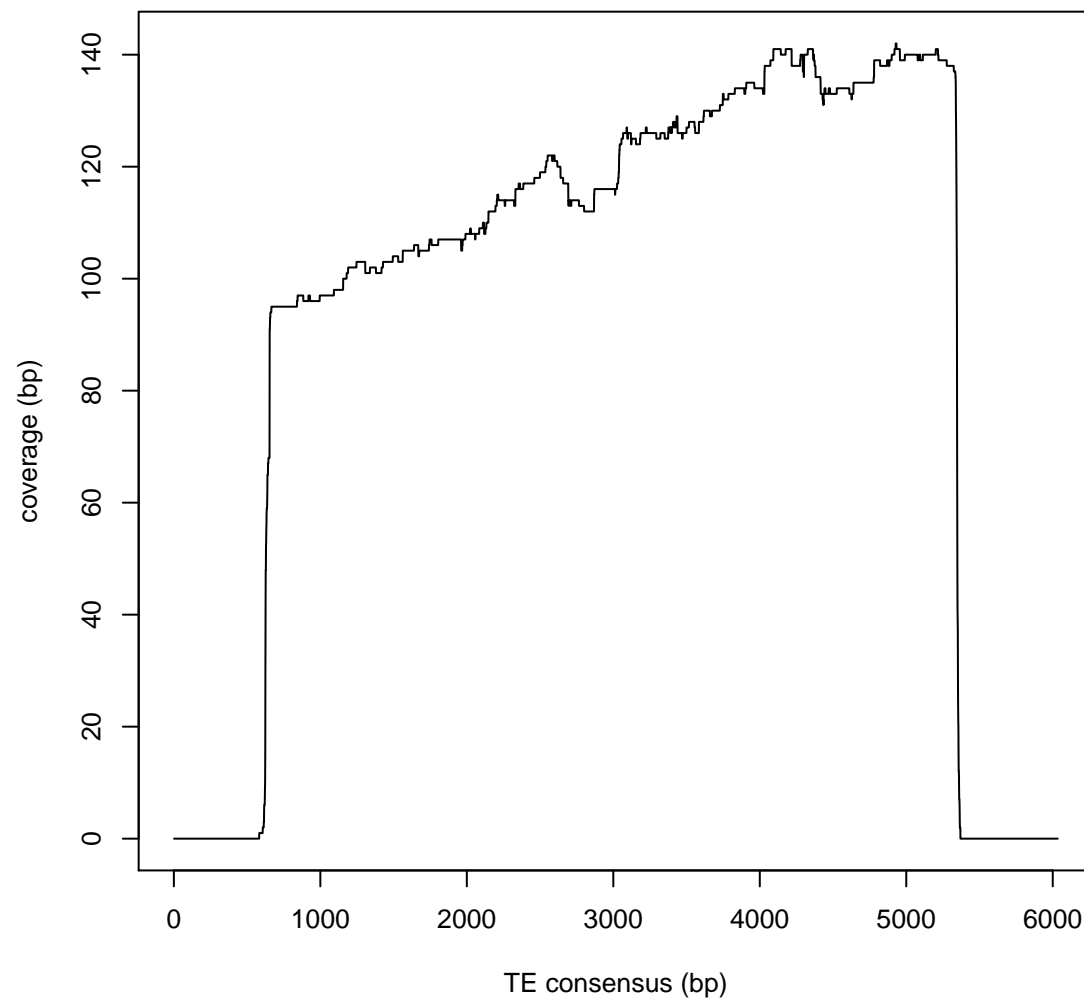


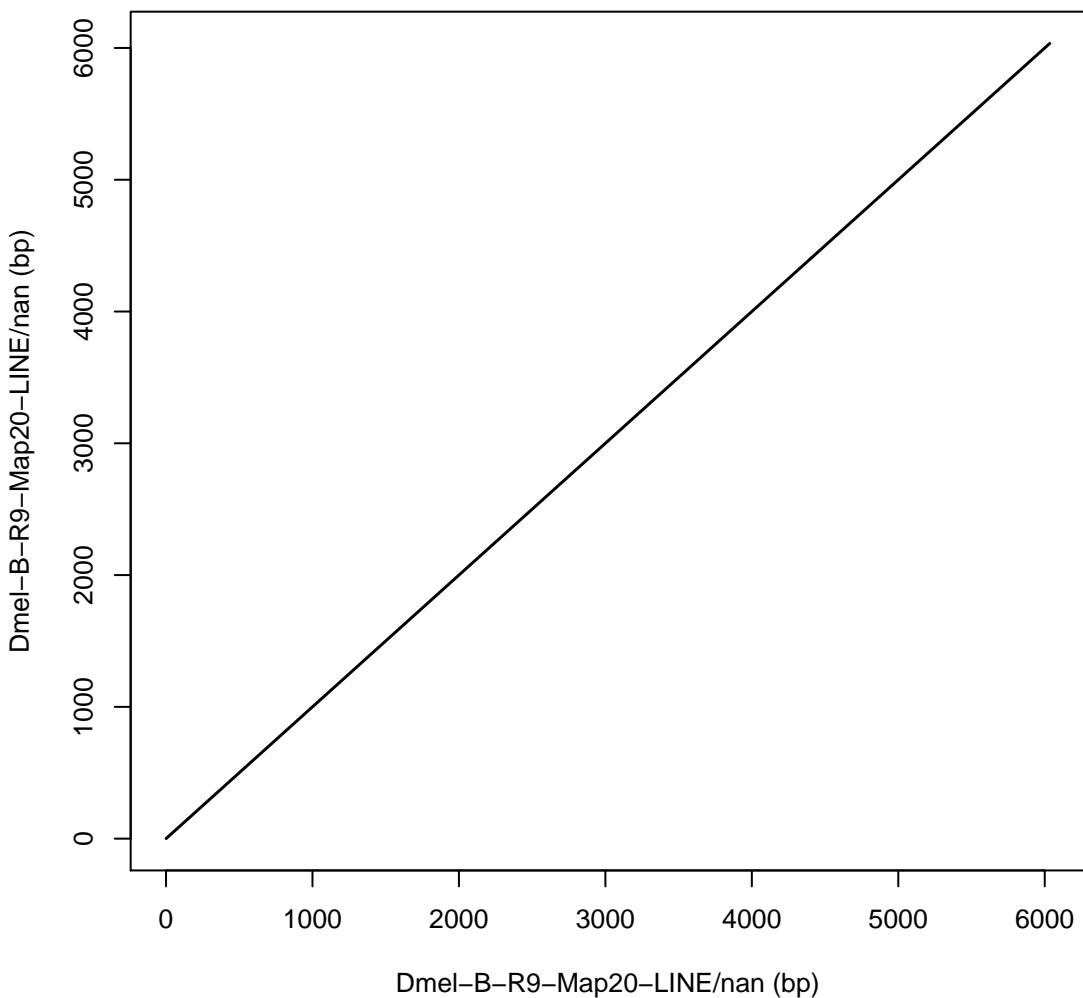
TE: Dmel-B-R9-Map20-LINE/nan
consensus size: 6034bp; fragments: 227; full length: 0 (>=5430.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

