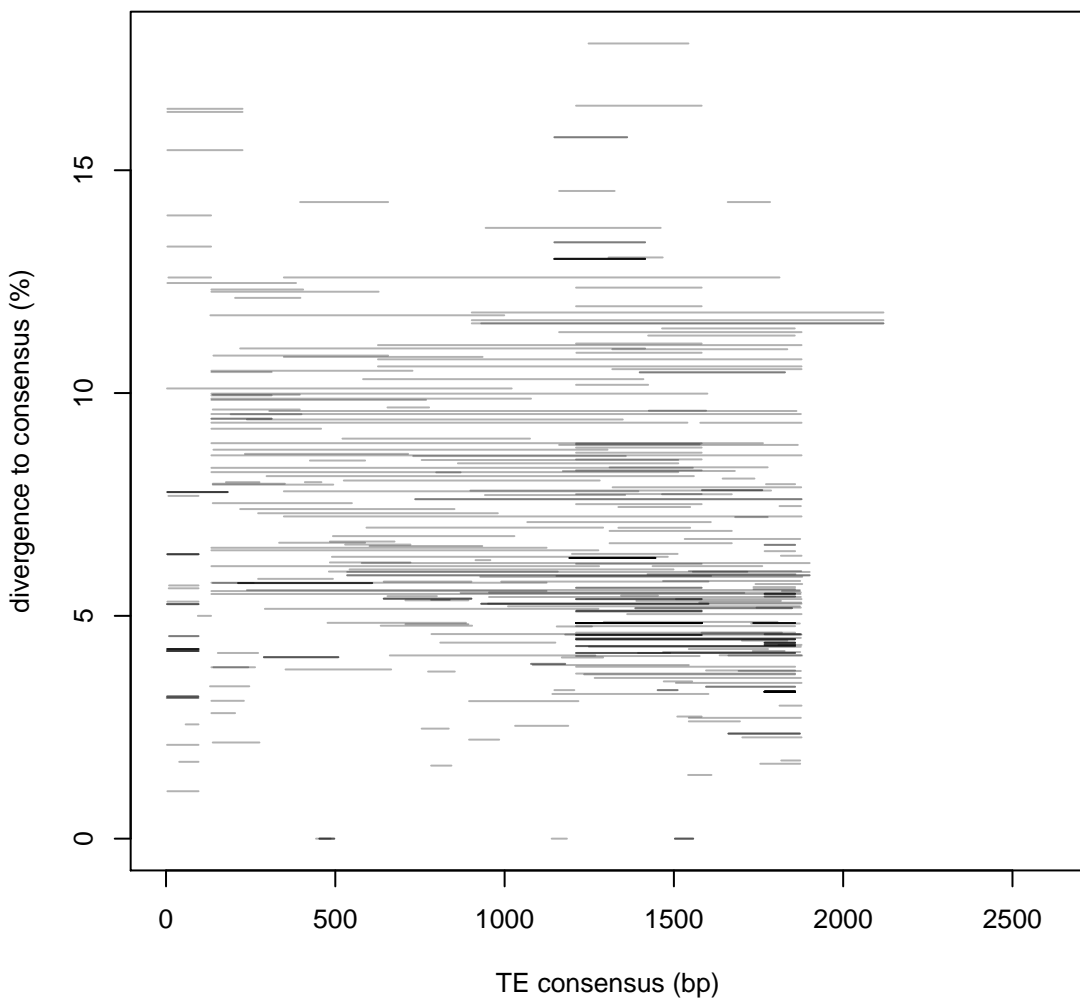
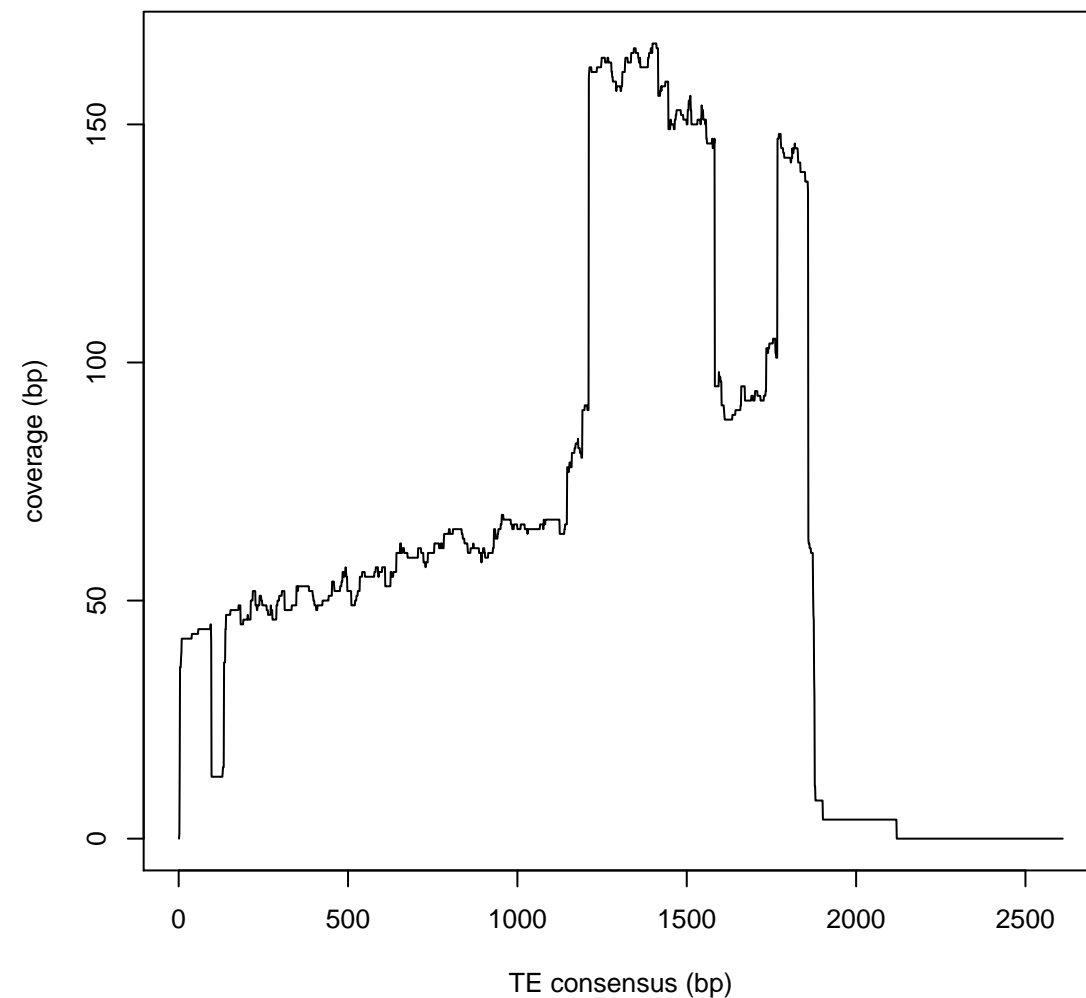


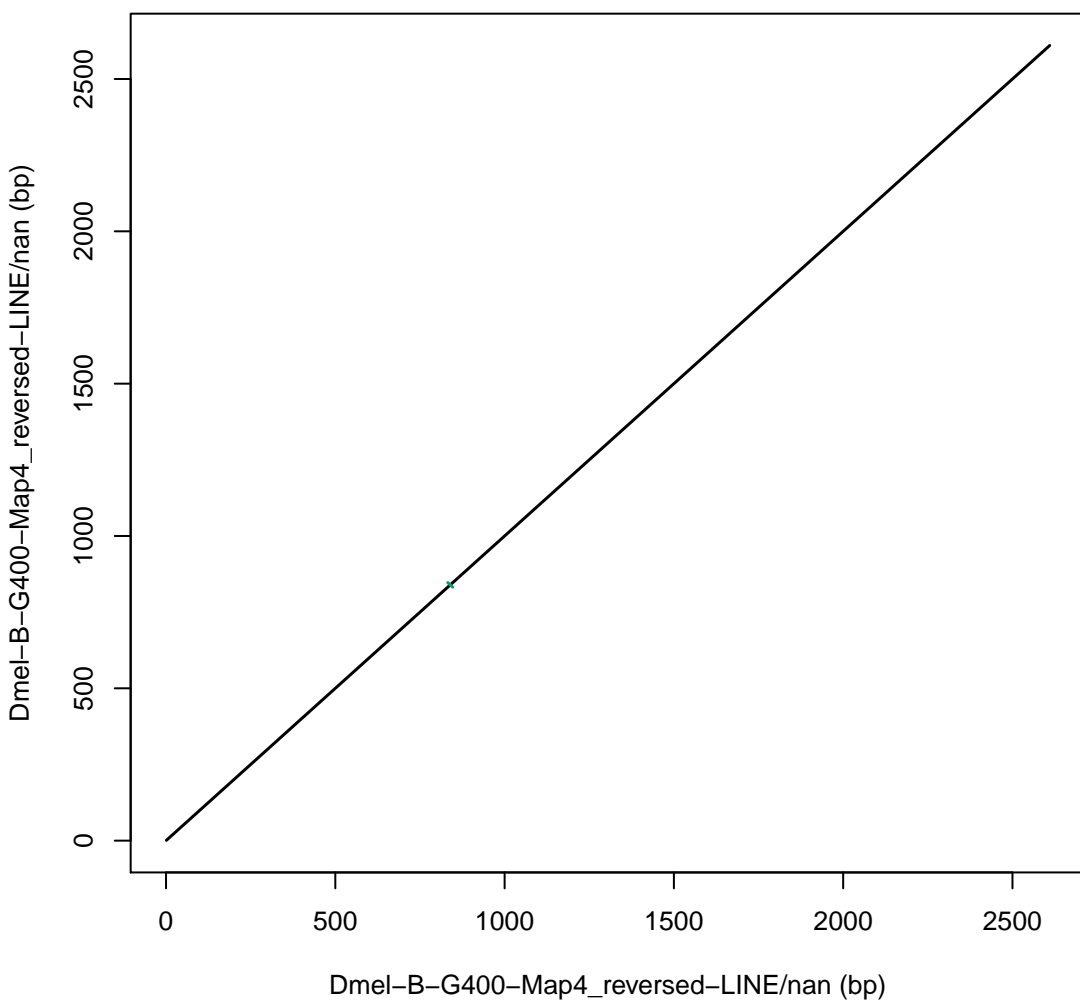
TE: Dmel-B-G400-Map4_reversed-LINE/nan
consensus size: 2610bp; fragments: 421; full length: 0 (>=2349bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

