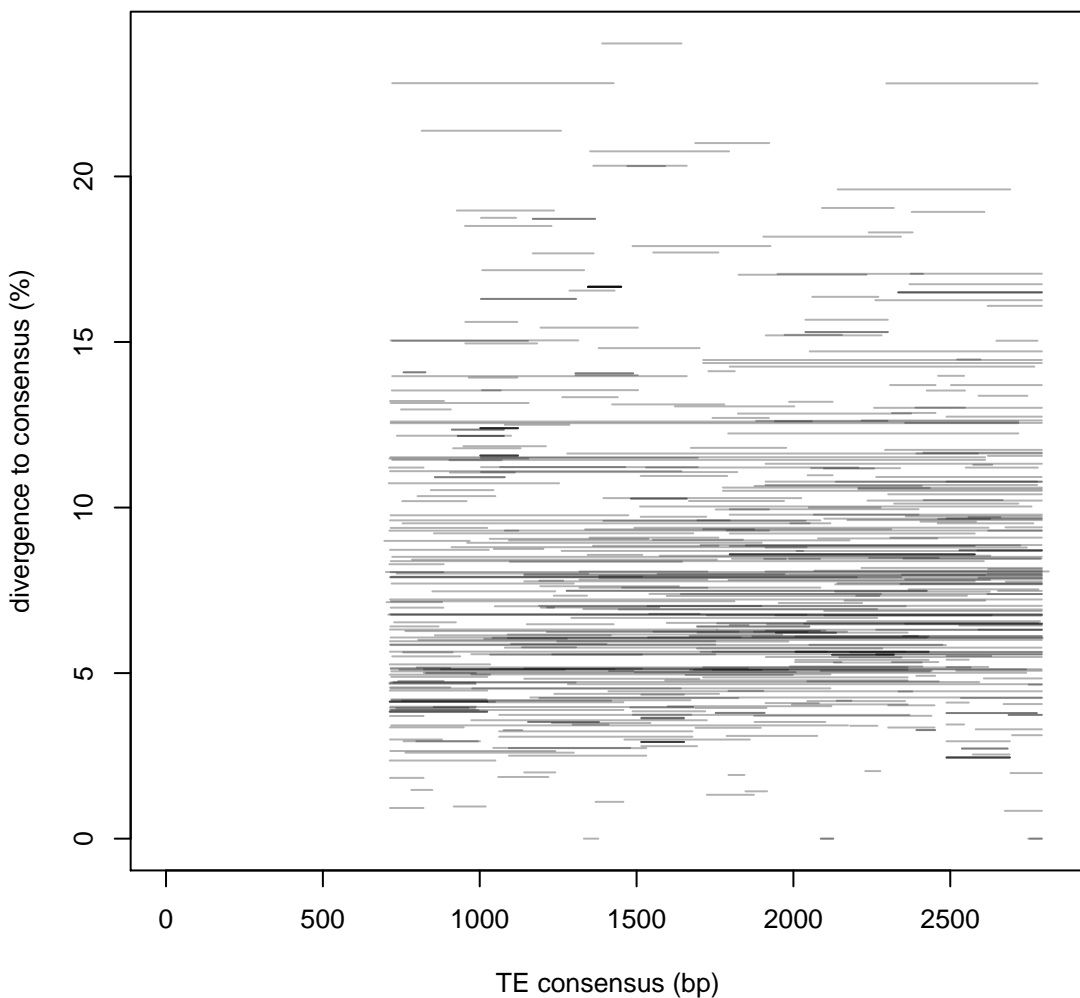
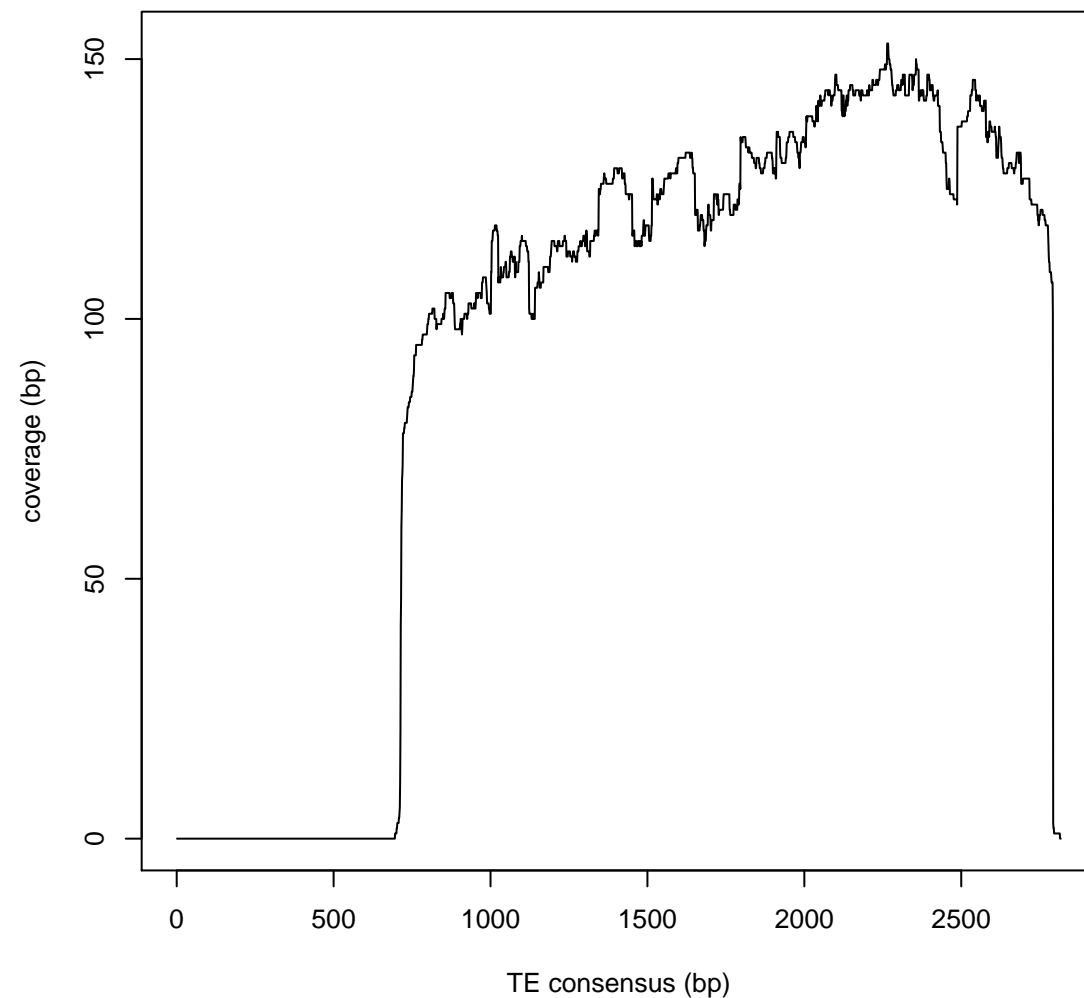


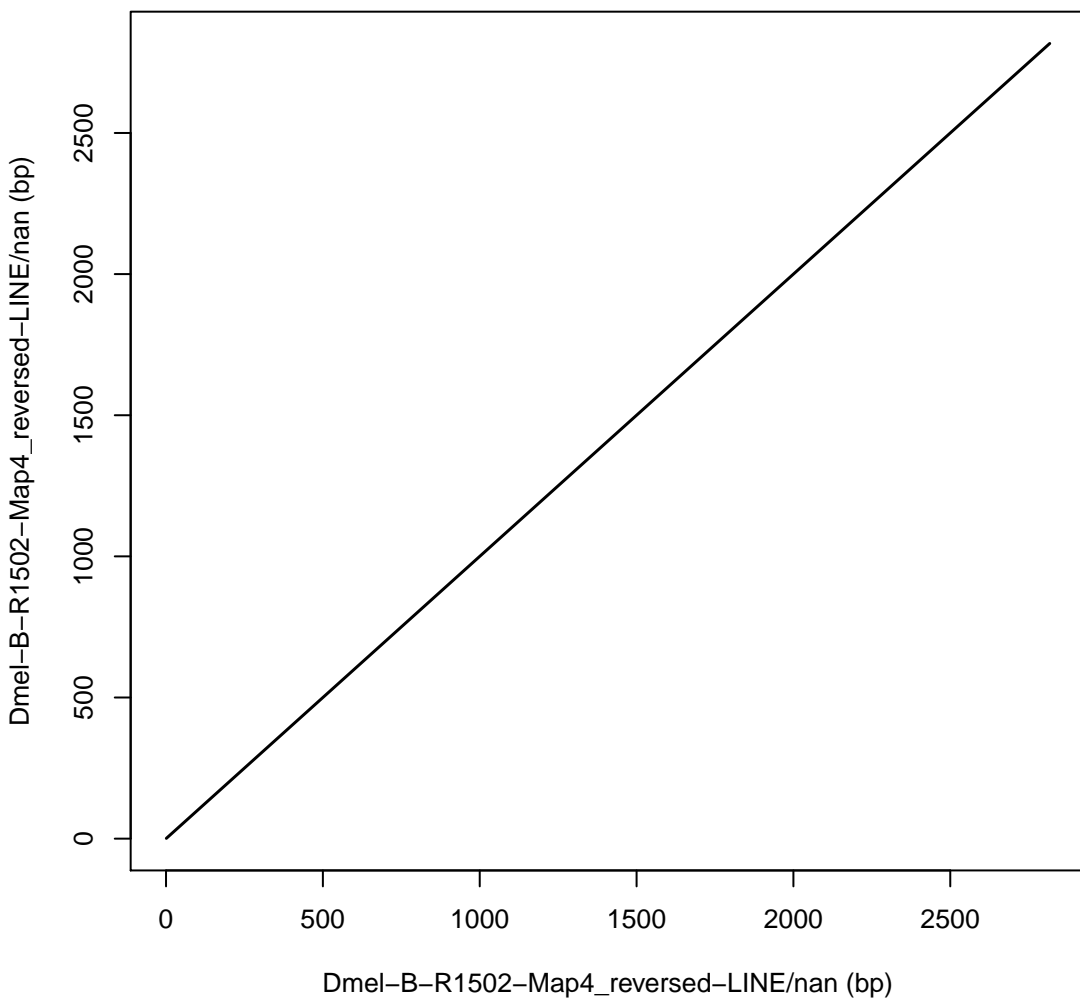
TE: Dmel-B-R1502-Map4\_reversed-LINE/nan  
consensus size: 2817bp; fragments: 523; full length: 0 (>=2535.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

