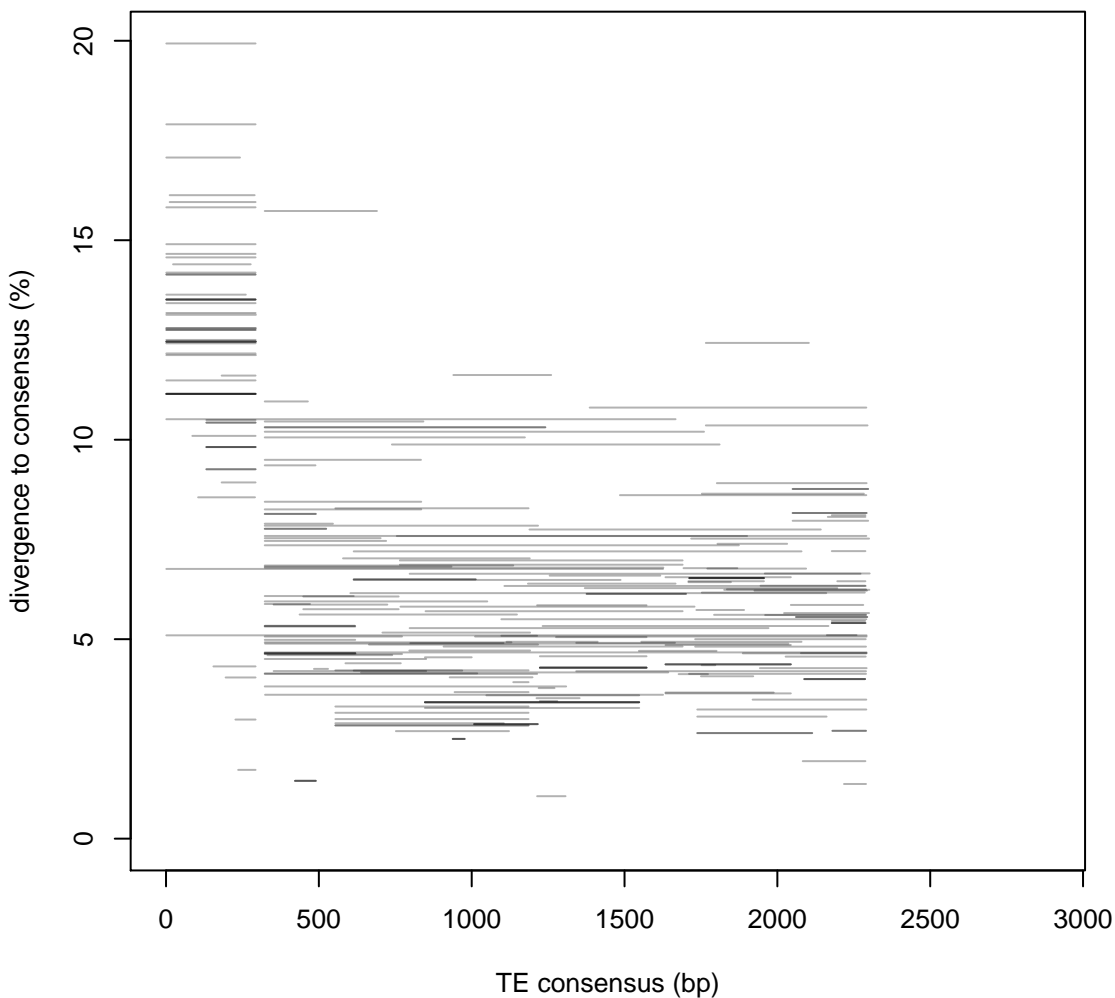
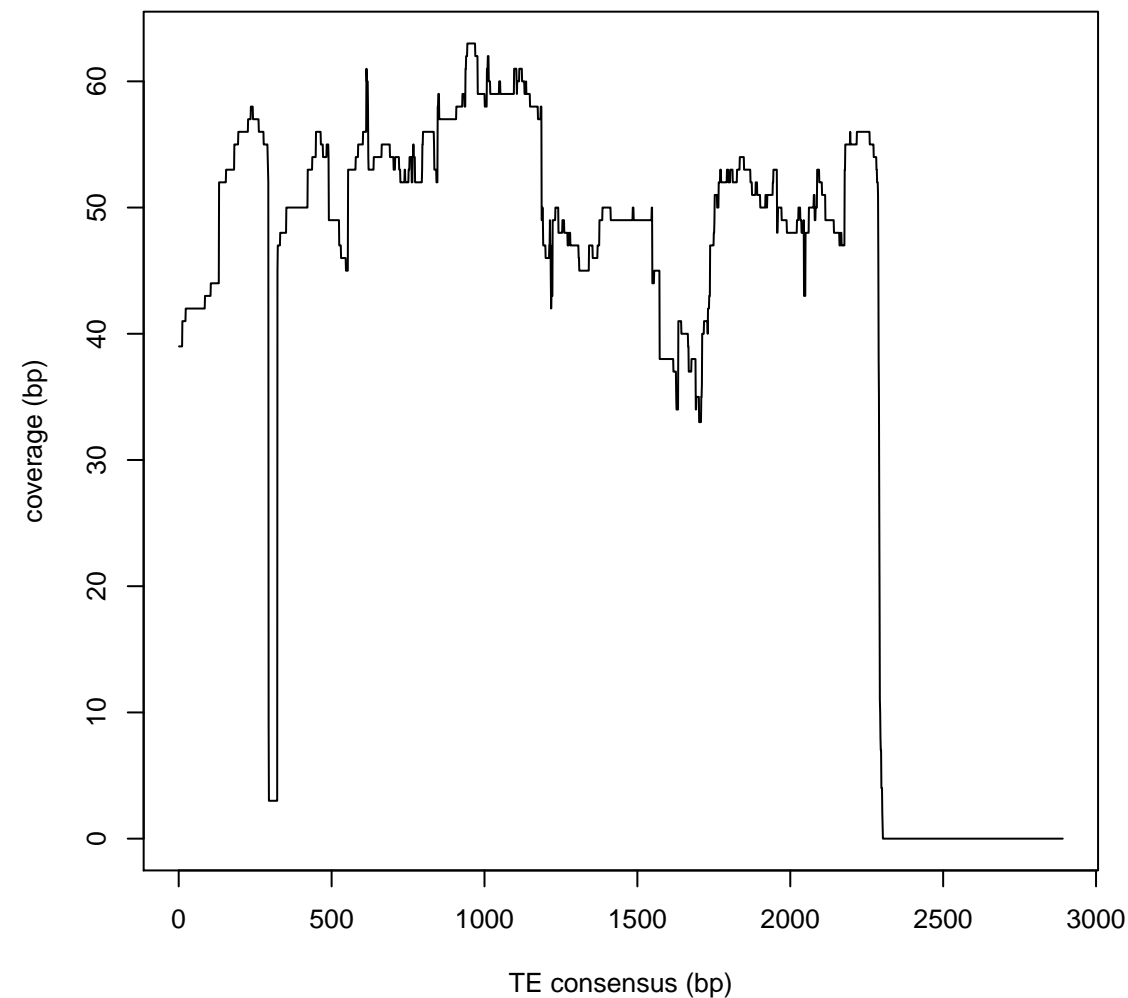


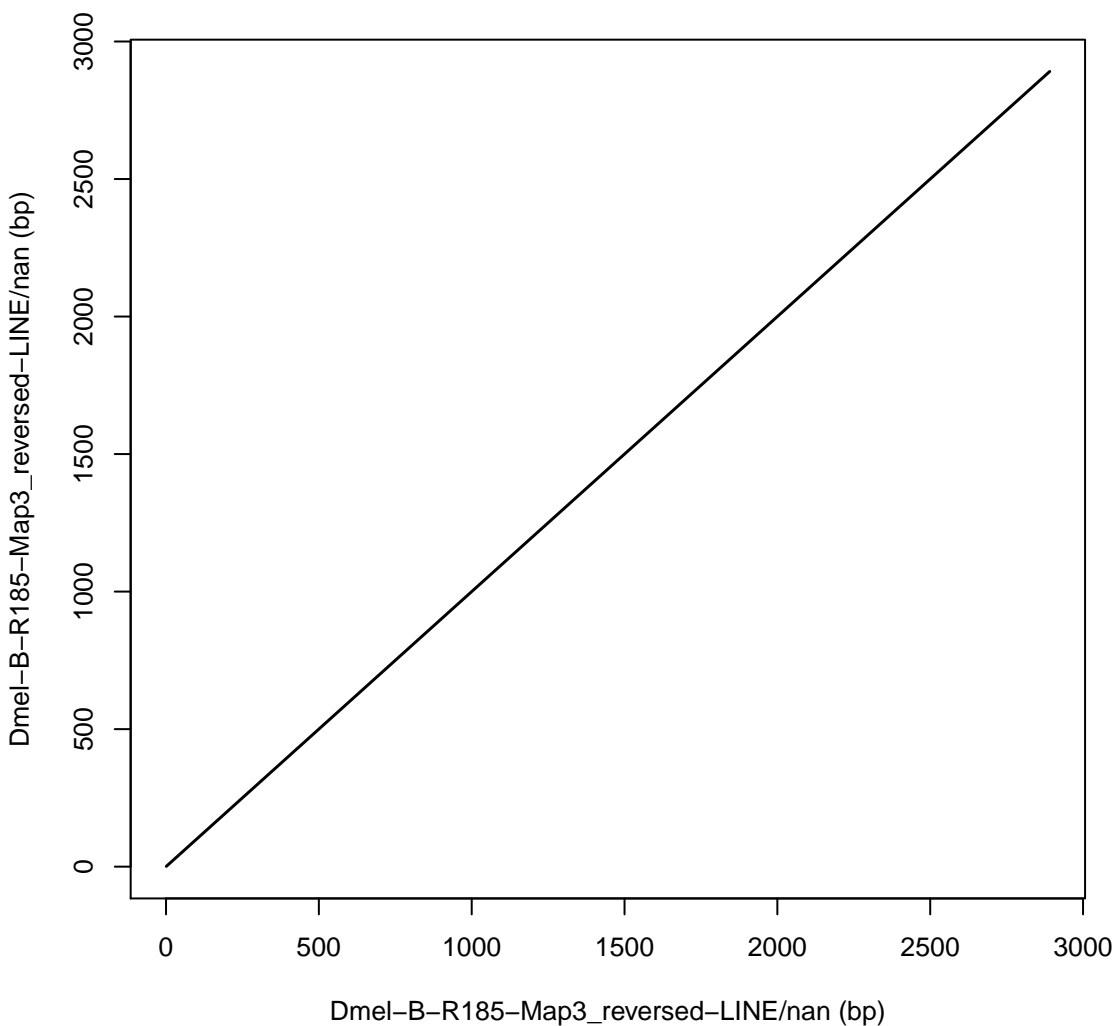
TE: Dmel-B-R185-Map3_reversed-LINE/nan
consensus size: 2891bp; fragments: 259; full length: 0 (>=2601.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

