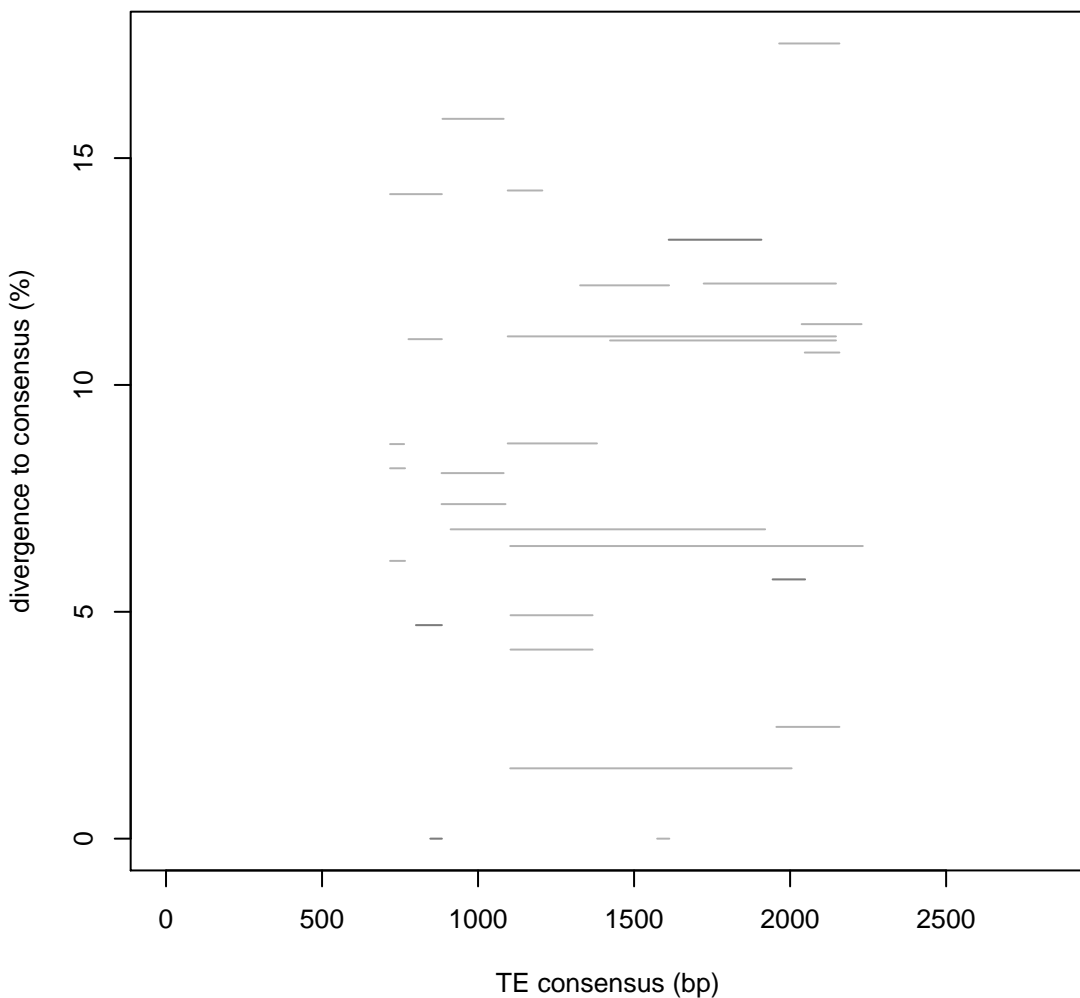
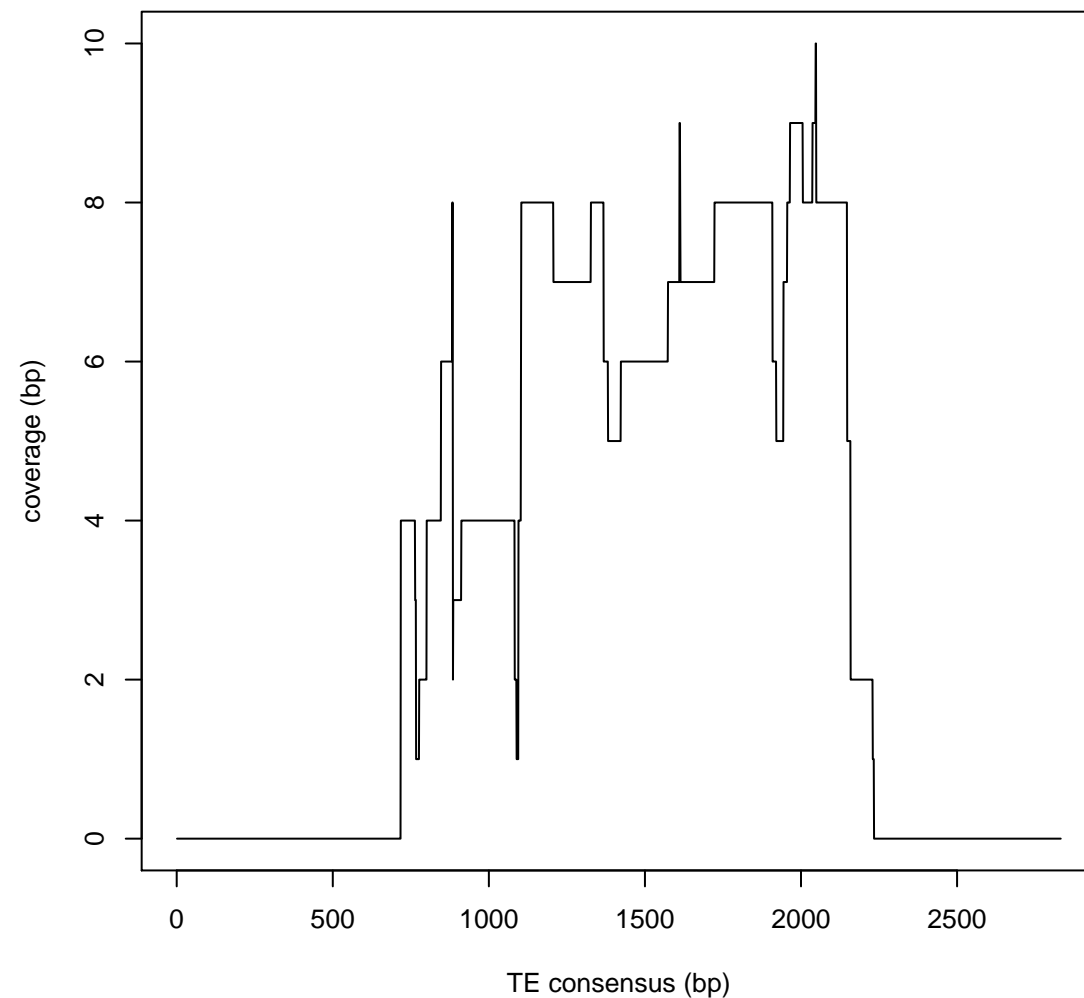


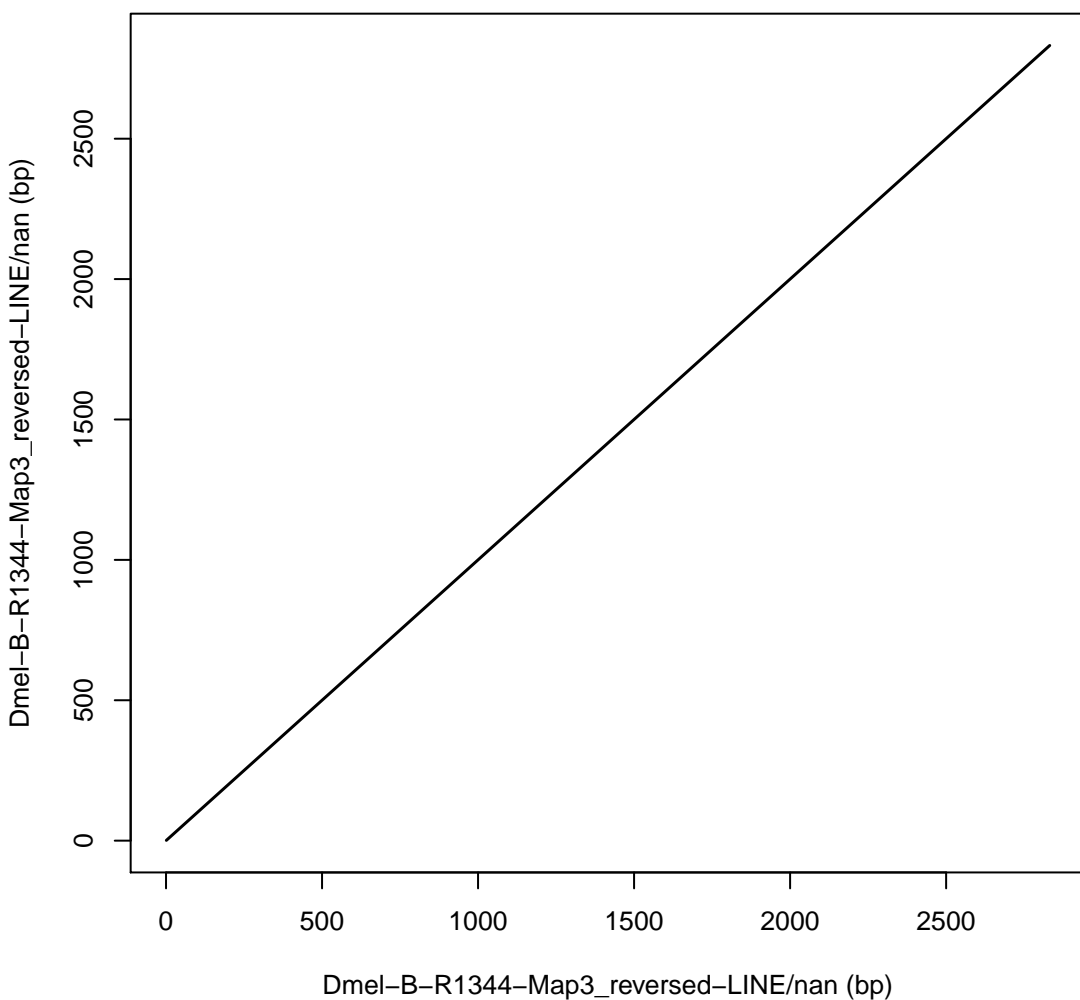
TE: Dmel-B-R1344-Map3_reversed-LINE/nan
consensus size: 2832bp; fragments: 32; full length: 0 (>=2548.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

