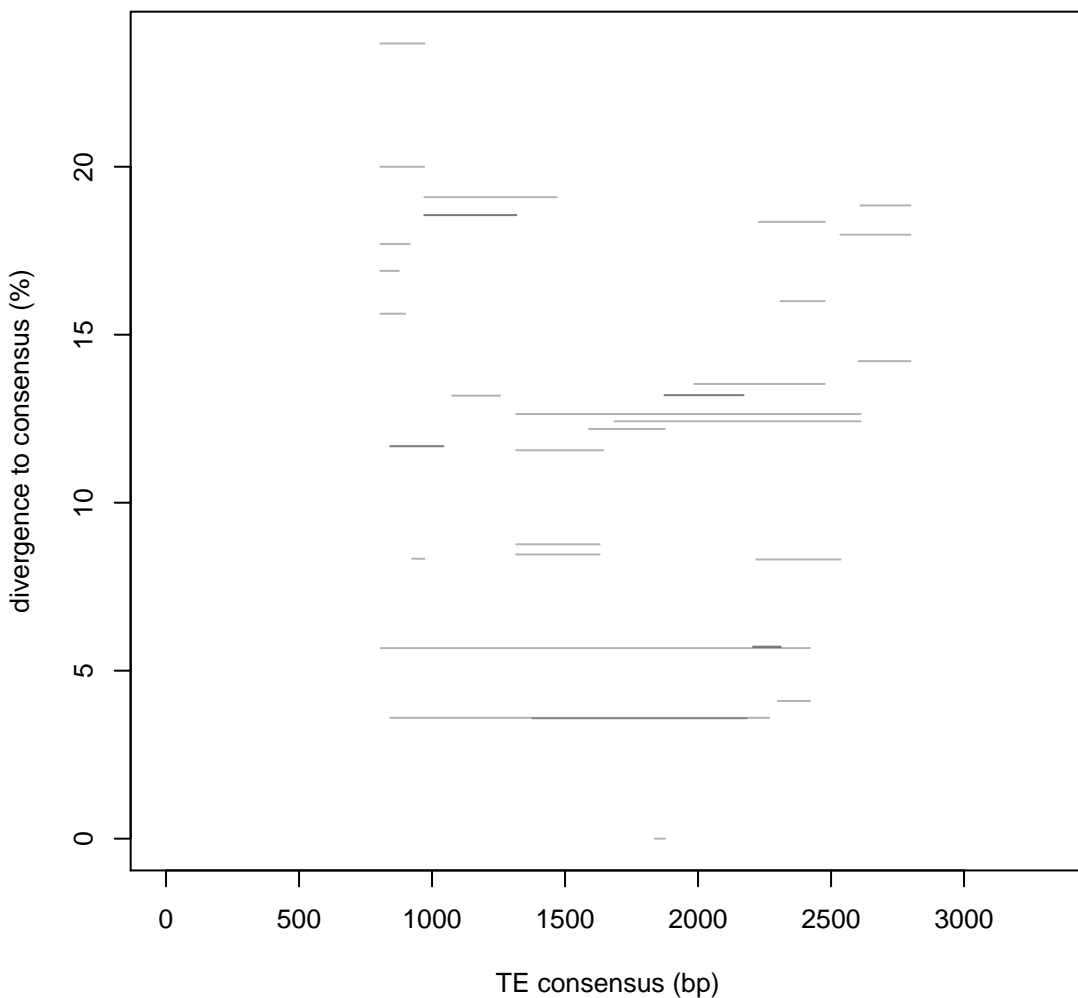
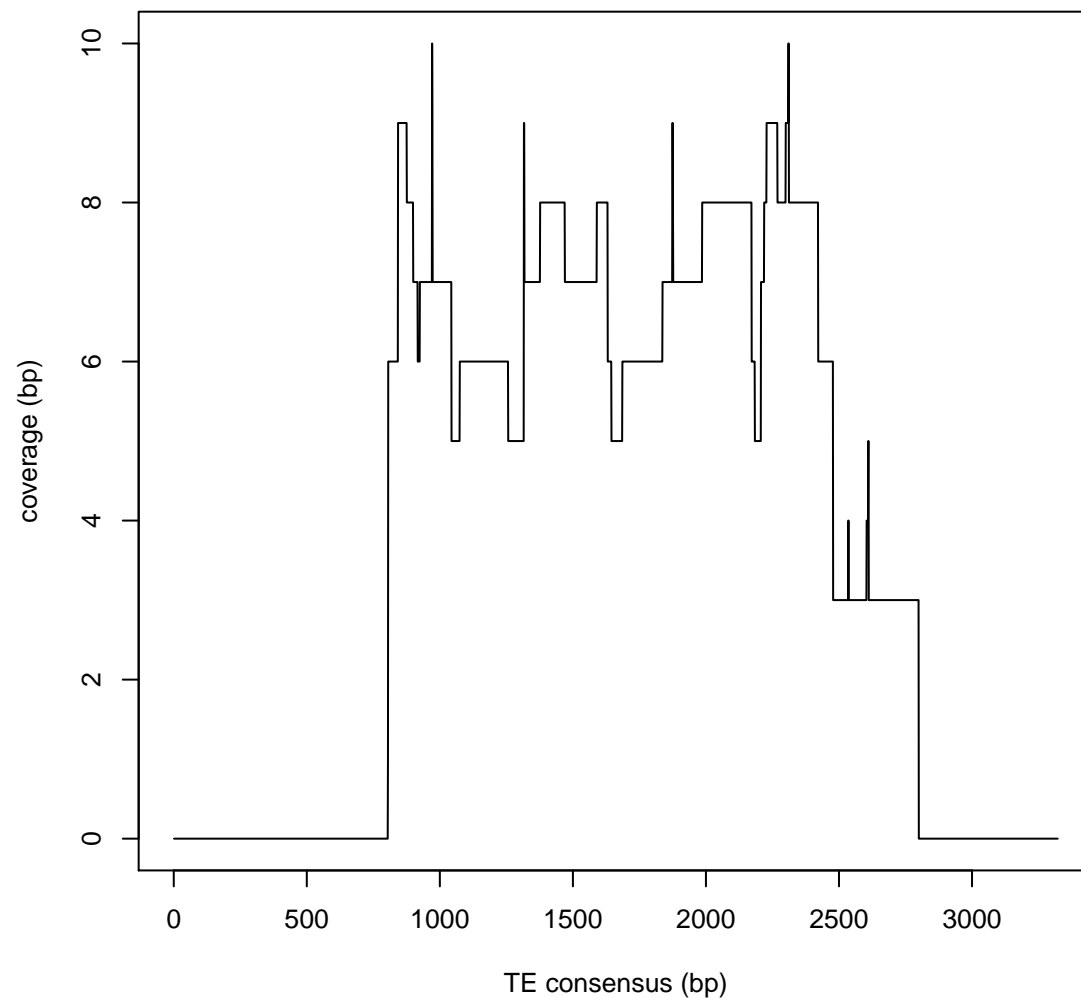


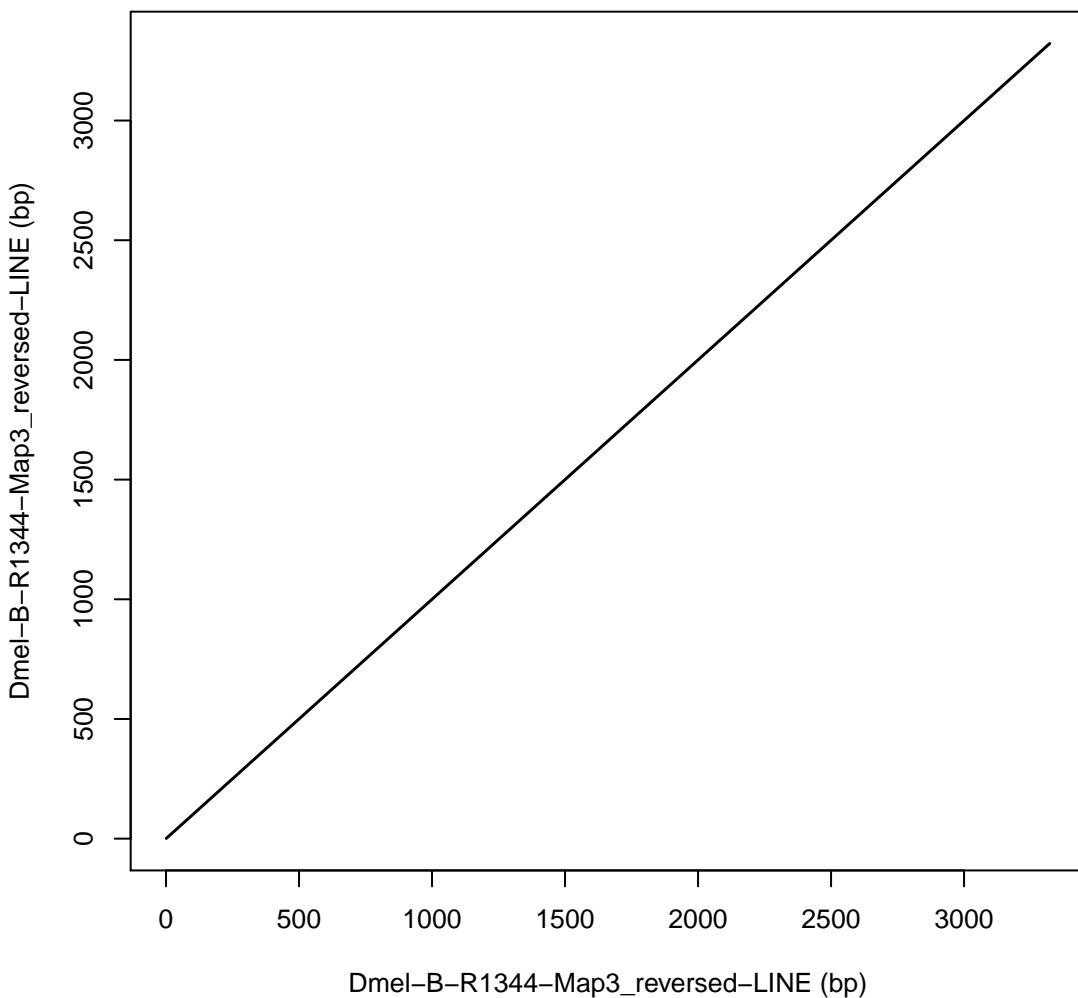
TE: Dmel-B-R1344-Map3_reversed-LINE
consensus size: 3322bp; fragments: 34; full length: 0 (>=2989.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

