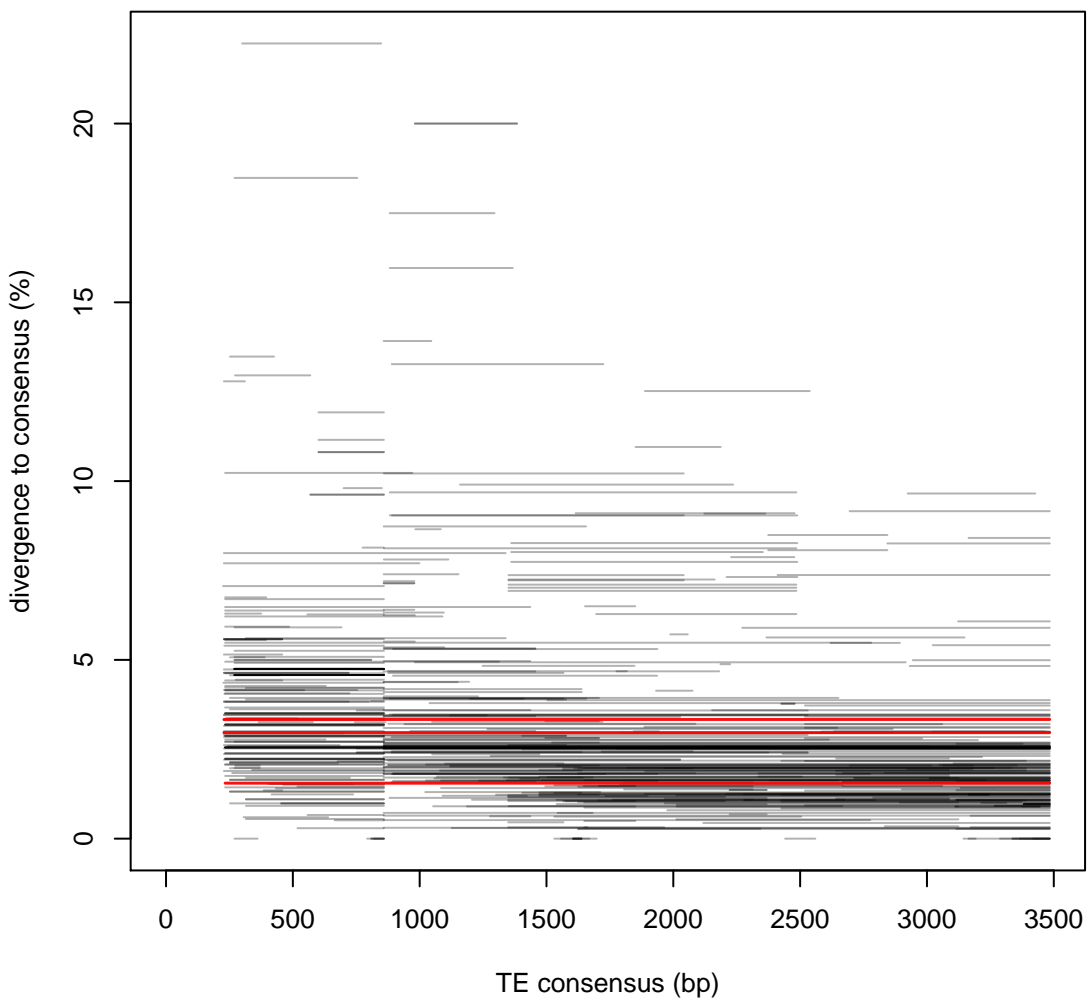
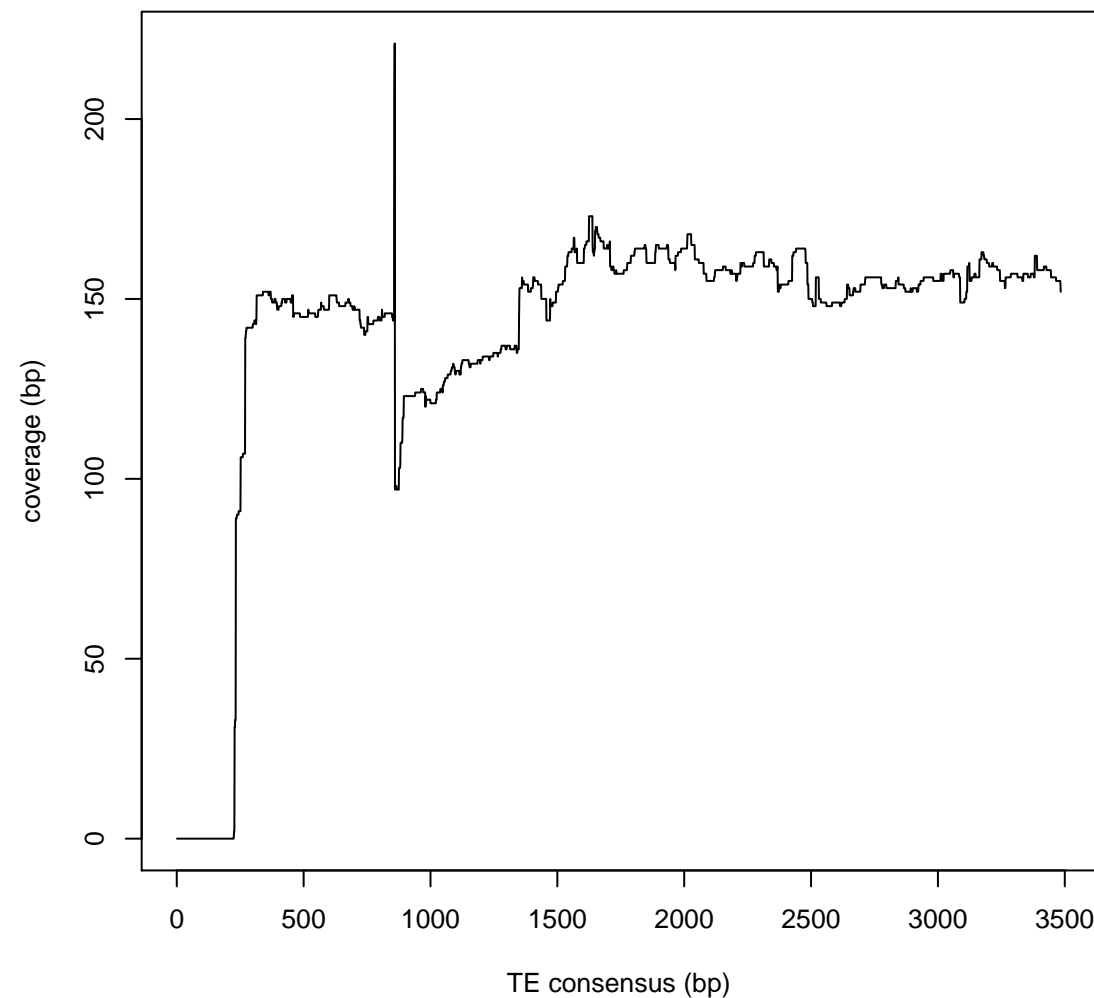


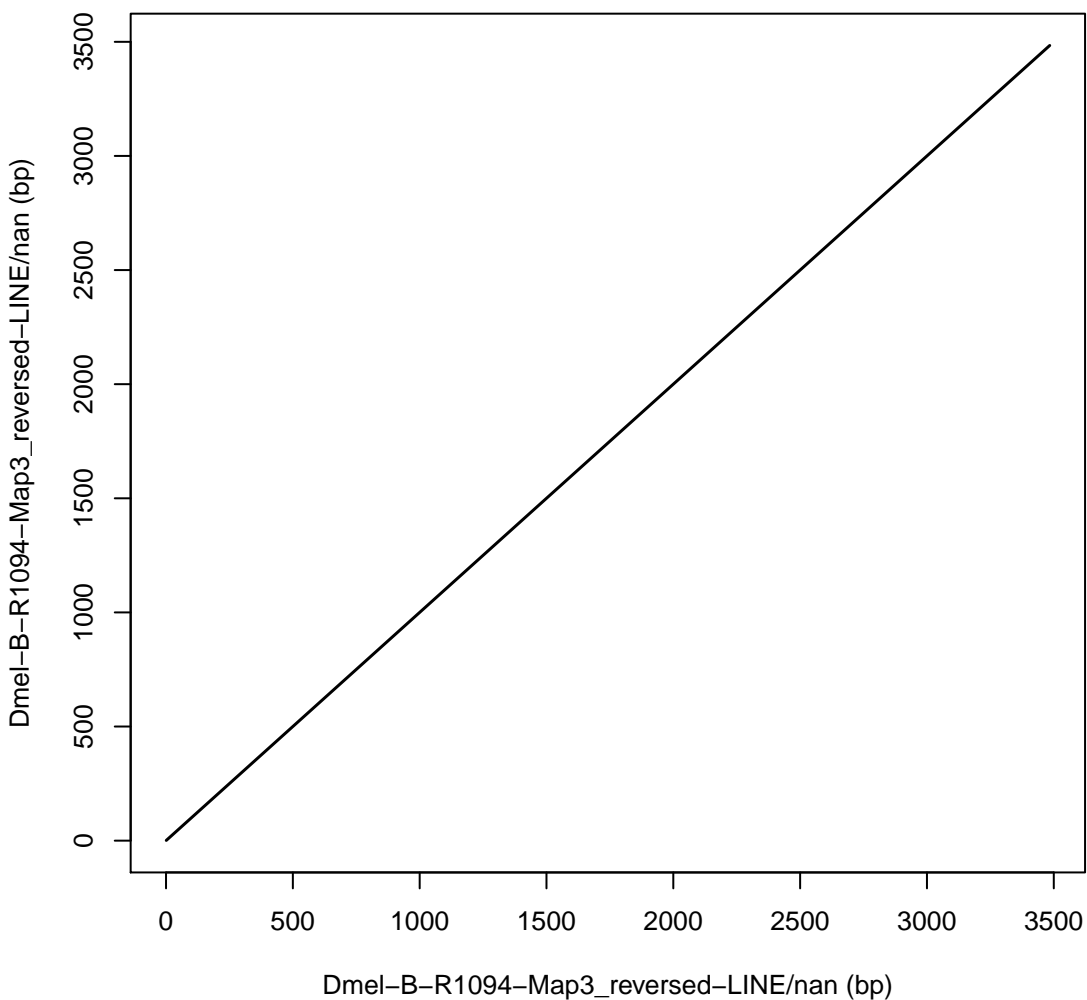
TE: Dmel-B-R1094-Map3_reversed-LINE/nan
consensus size: 3484bp; fragments: 534; full length: 3 (>=3135.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

