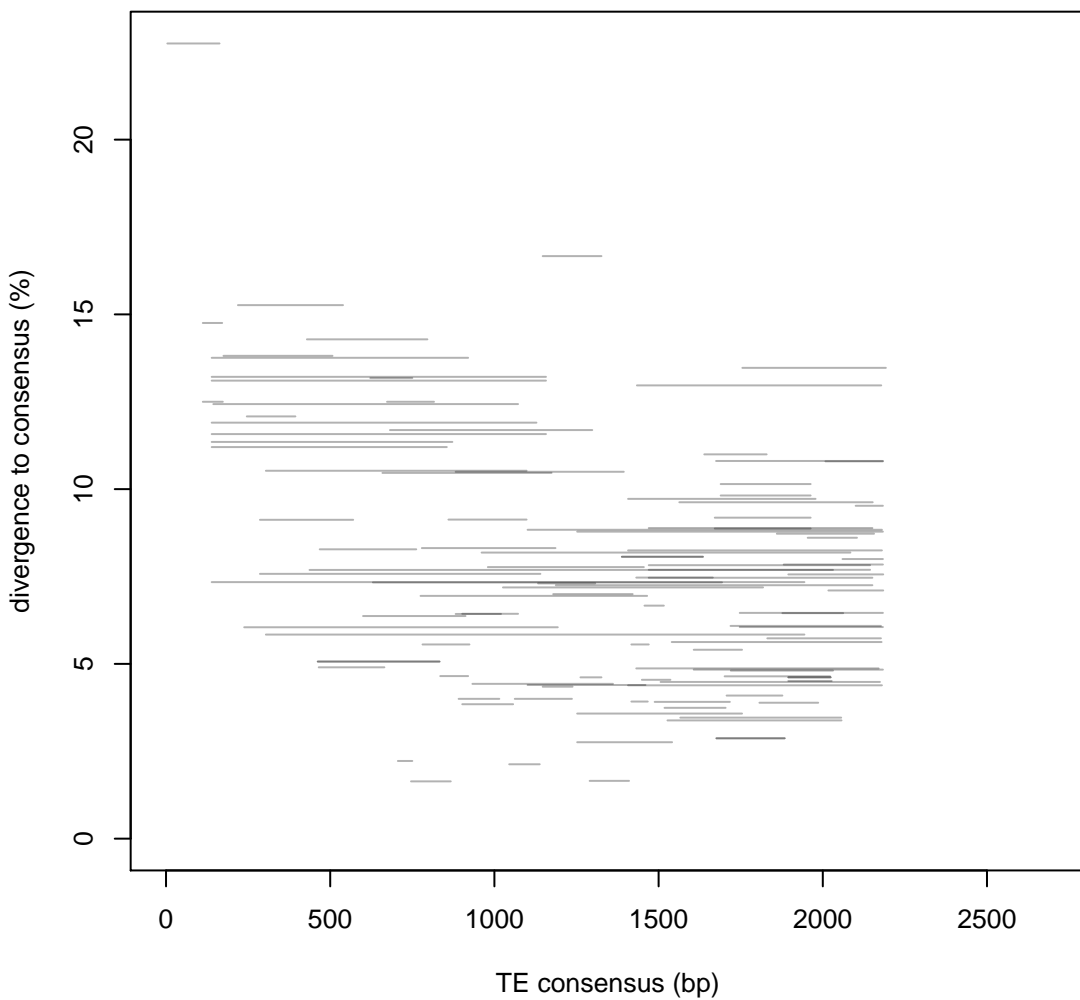
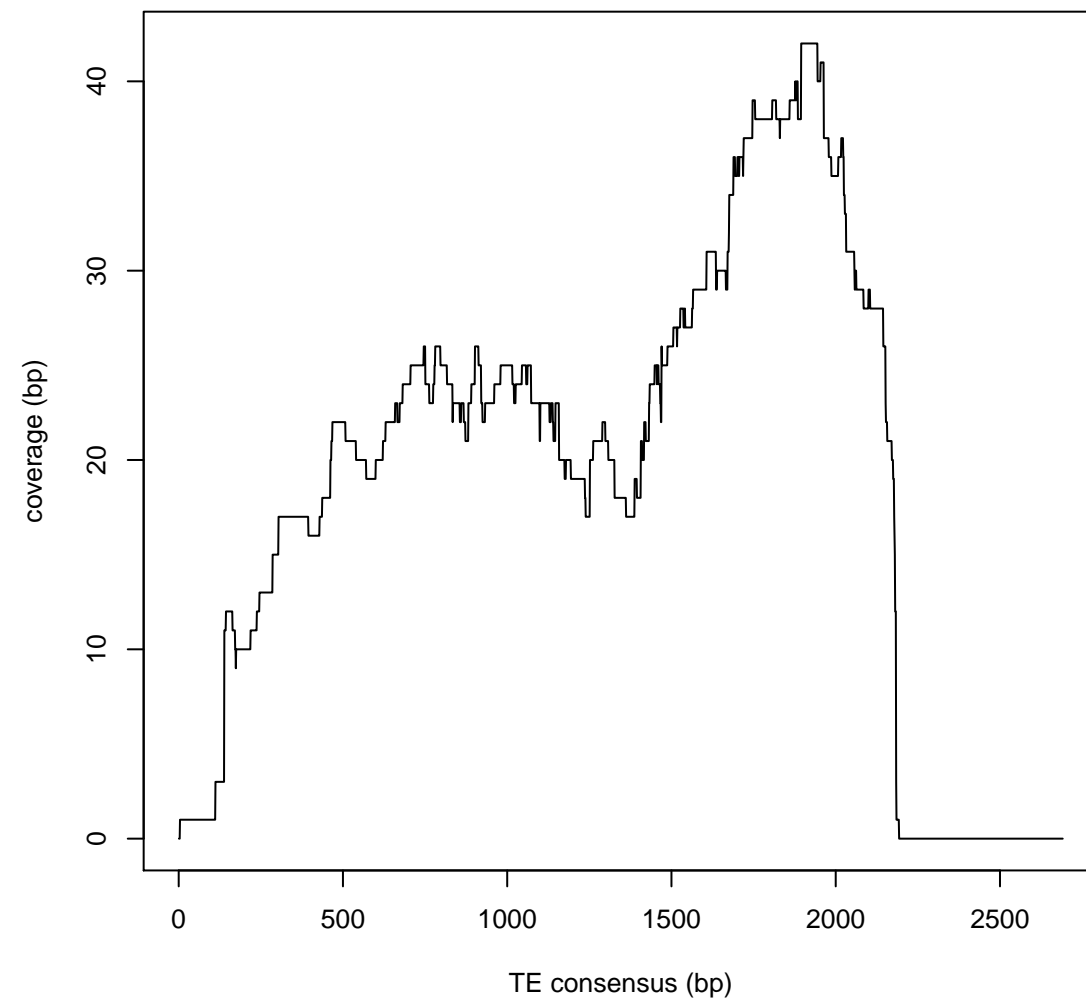


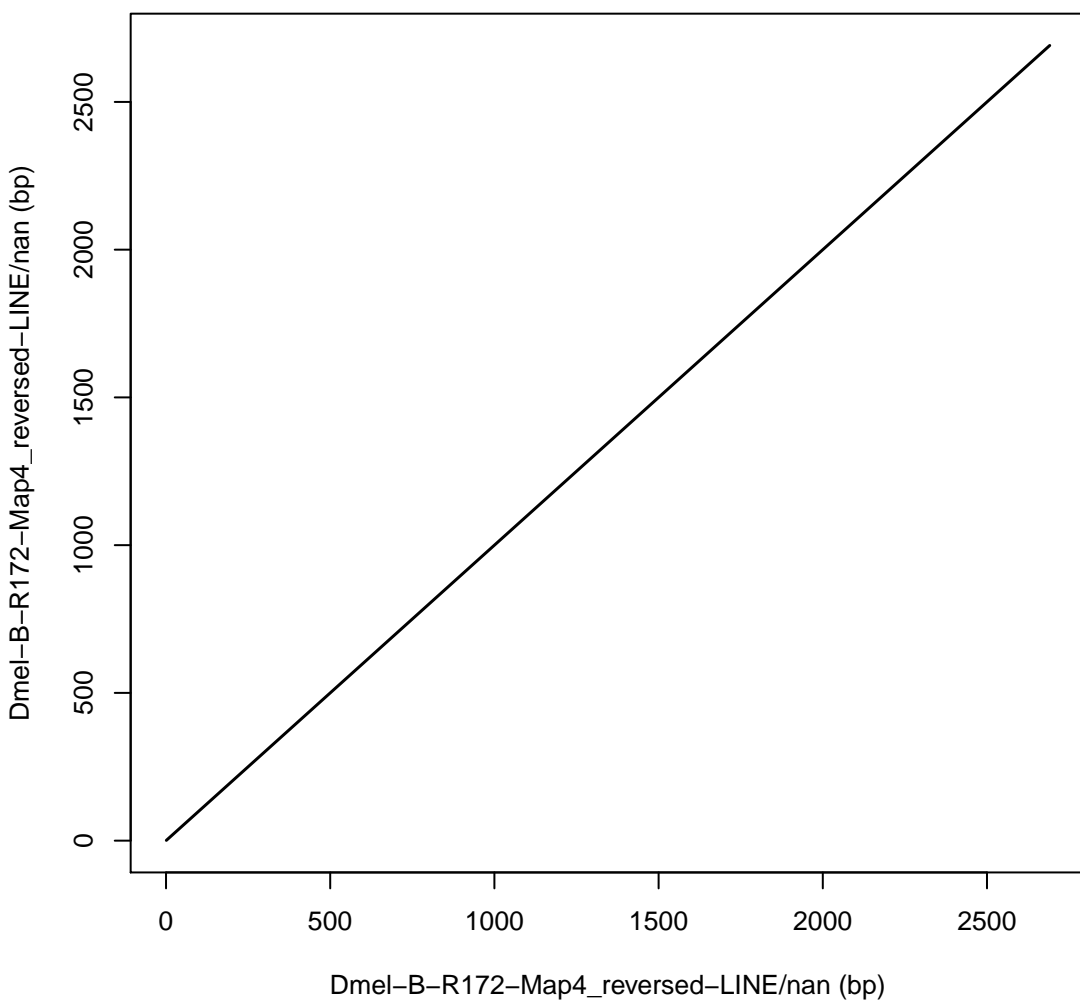
TE: Dmel-B-R172-Map4_reversed-LINE/nan
consensus size: 2691bp; fragments: 116; full length: 0 (>=2421.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

