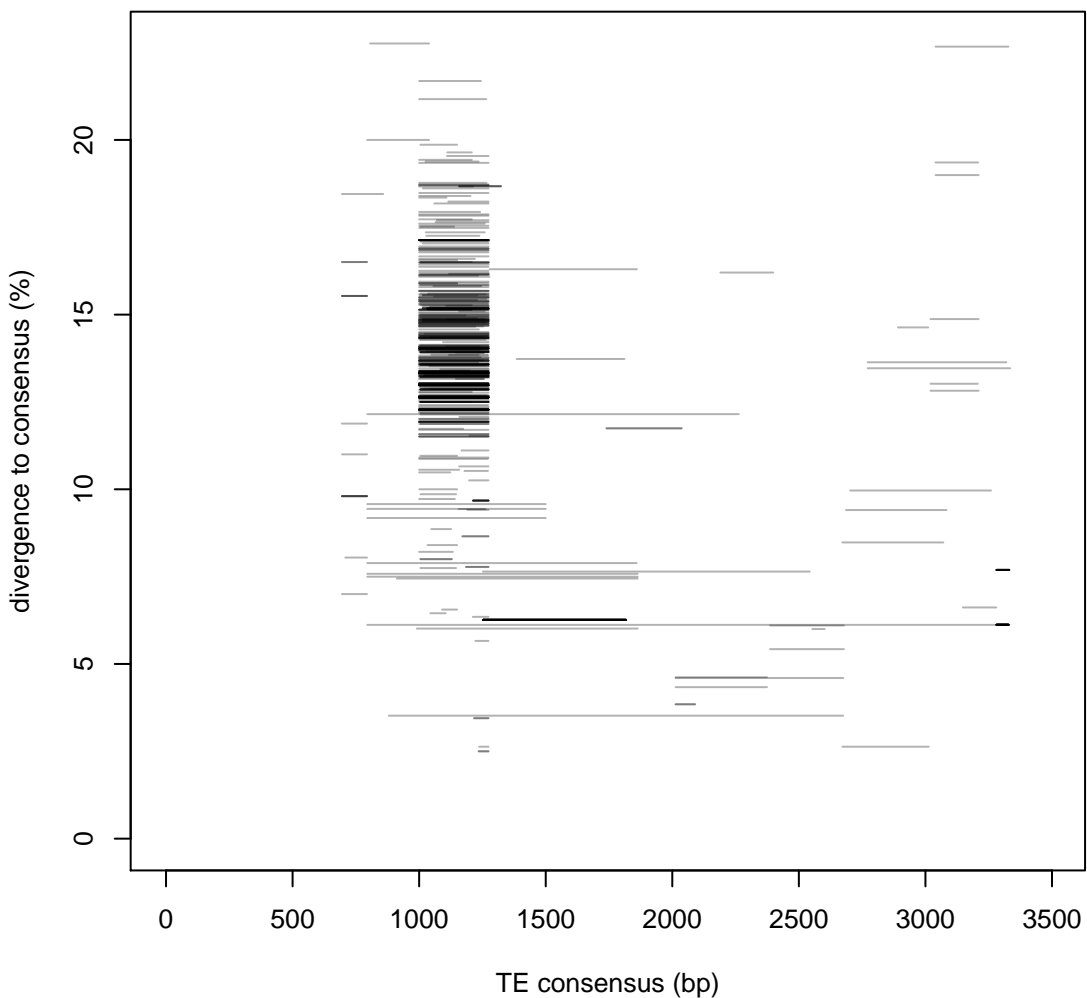
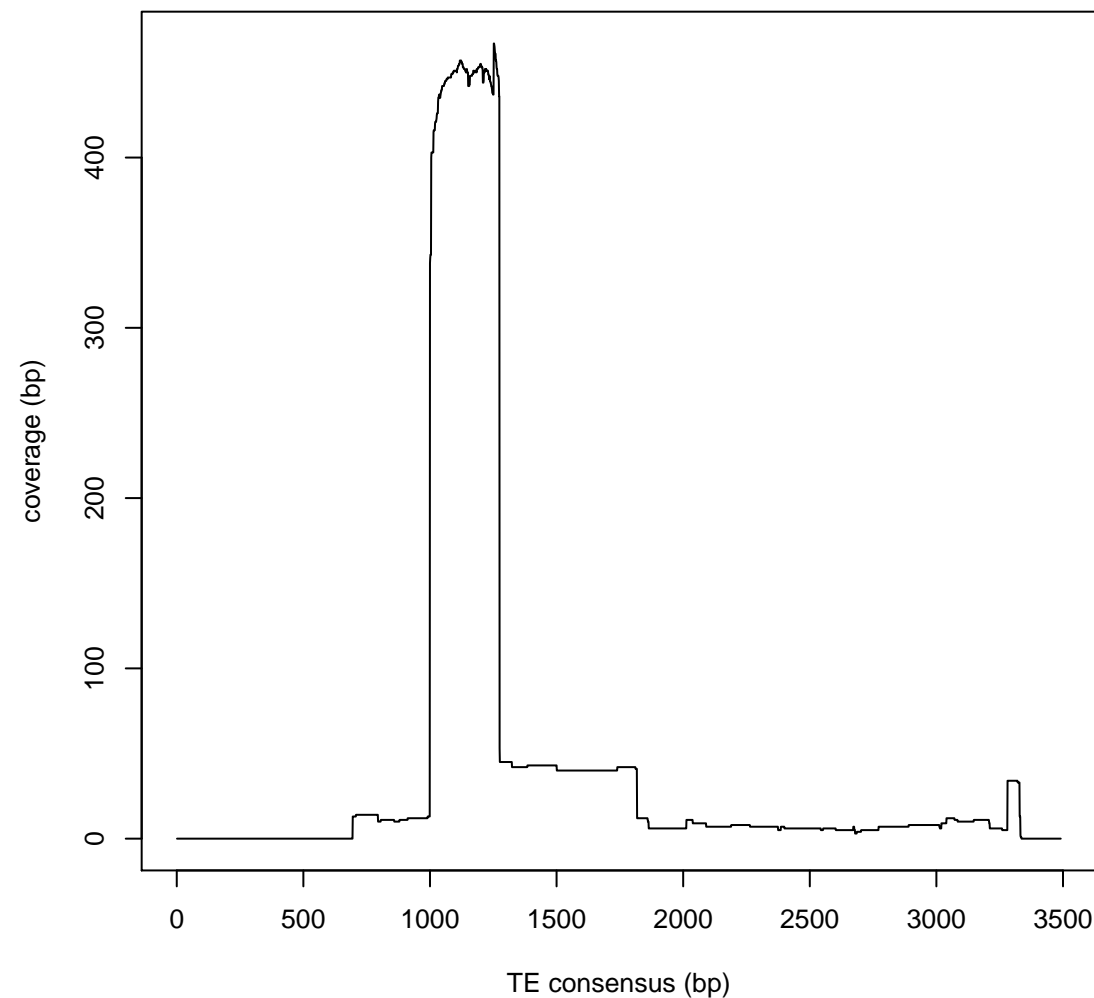


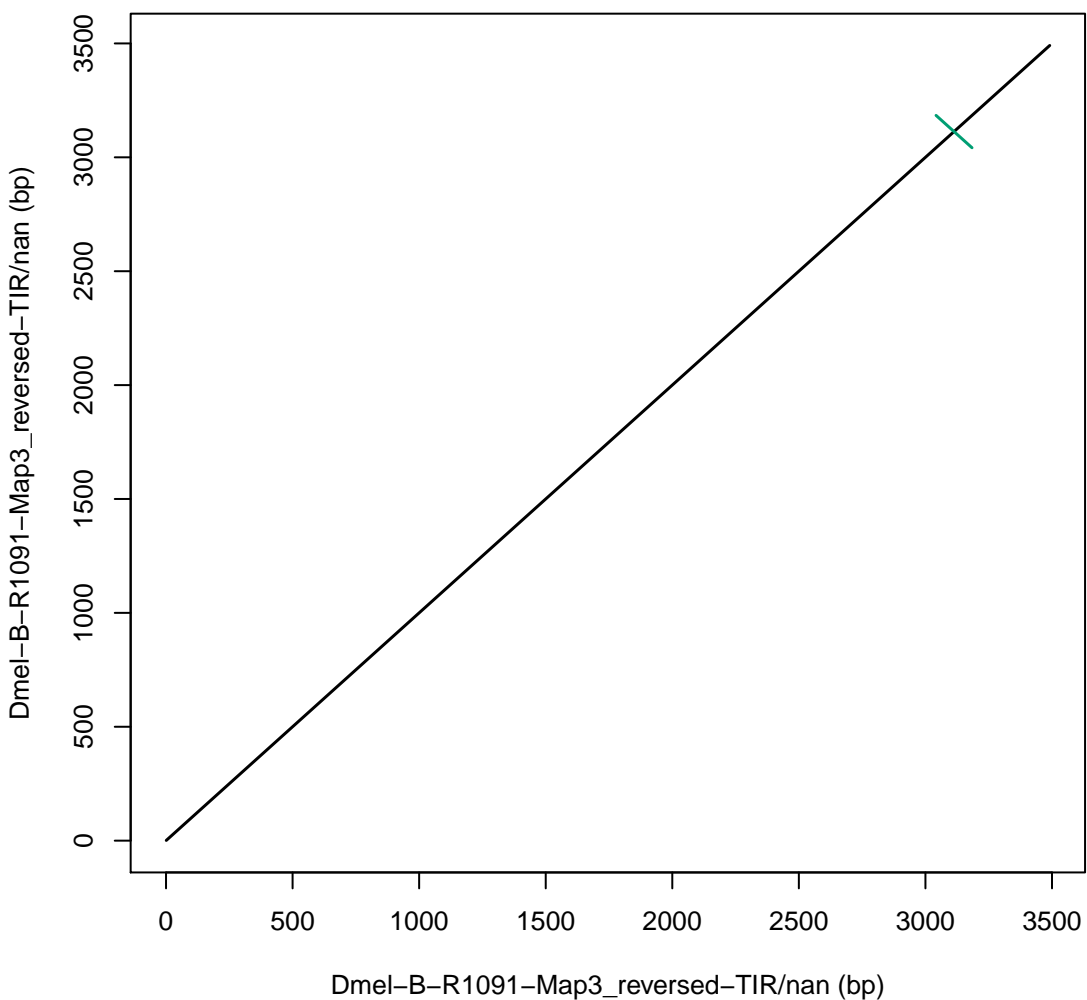
TE: Dmel-B-R1091-Map3_reversed-TIR/nan
consensus size: 3491bp; fragments: 593; full length: 0 (>=3141.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

