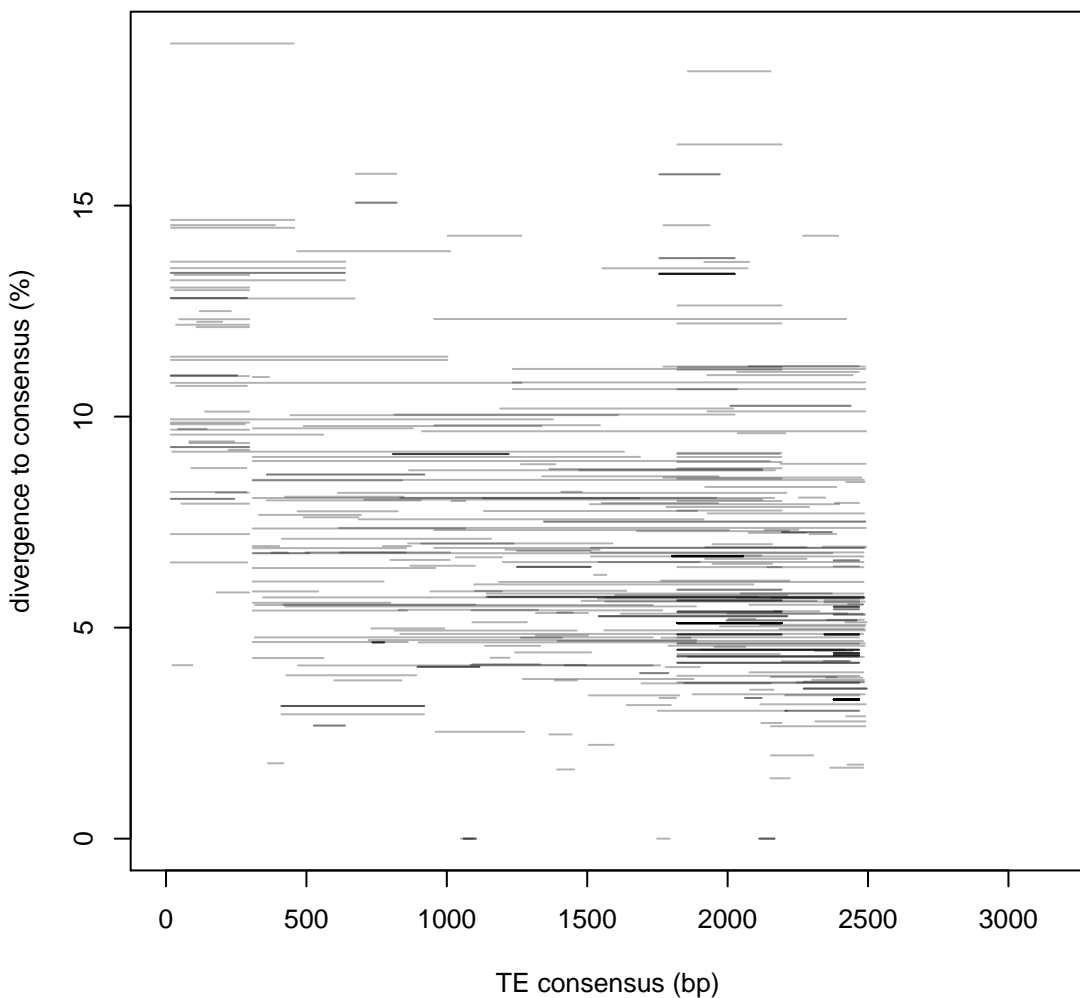
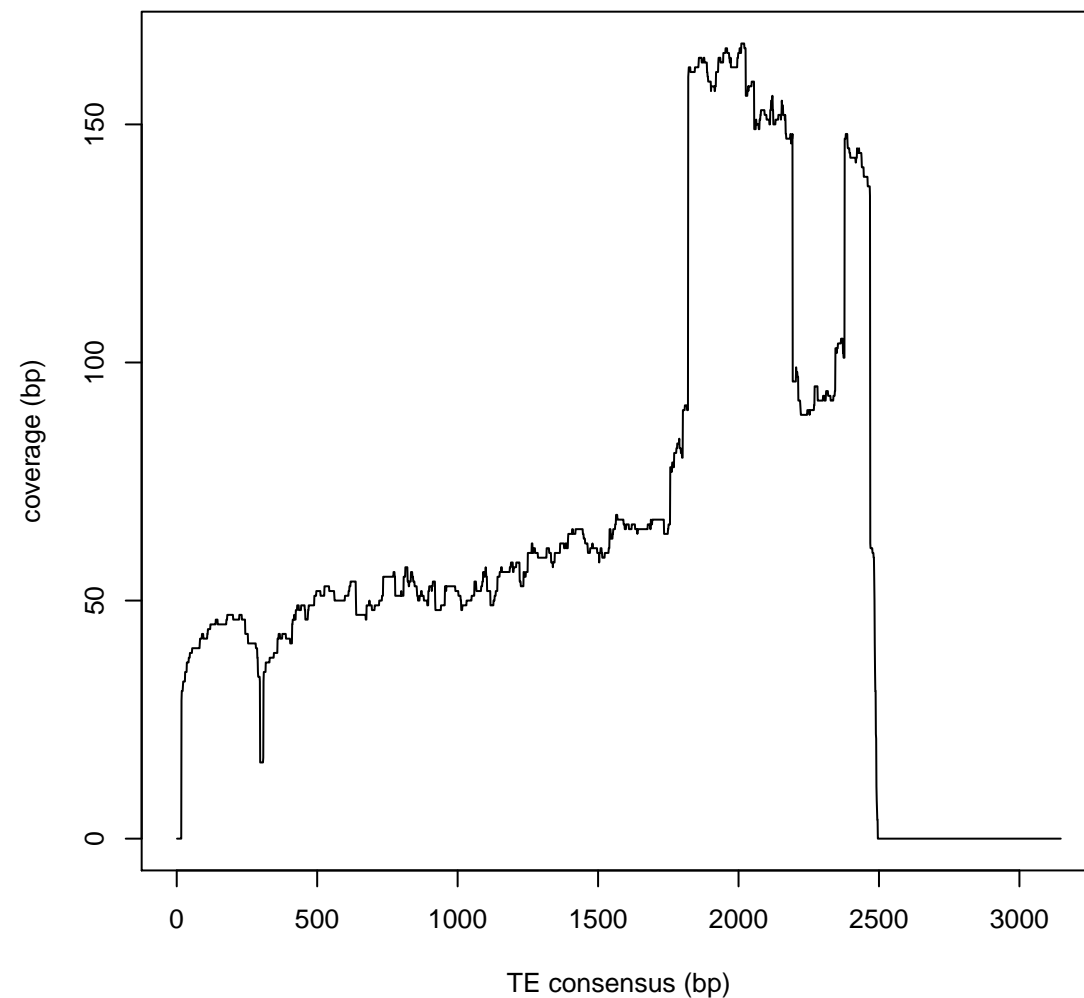


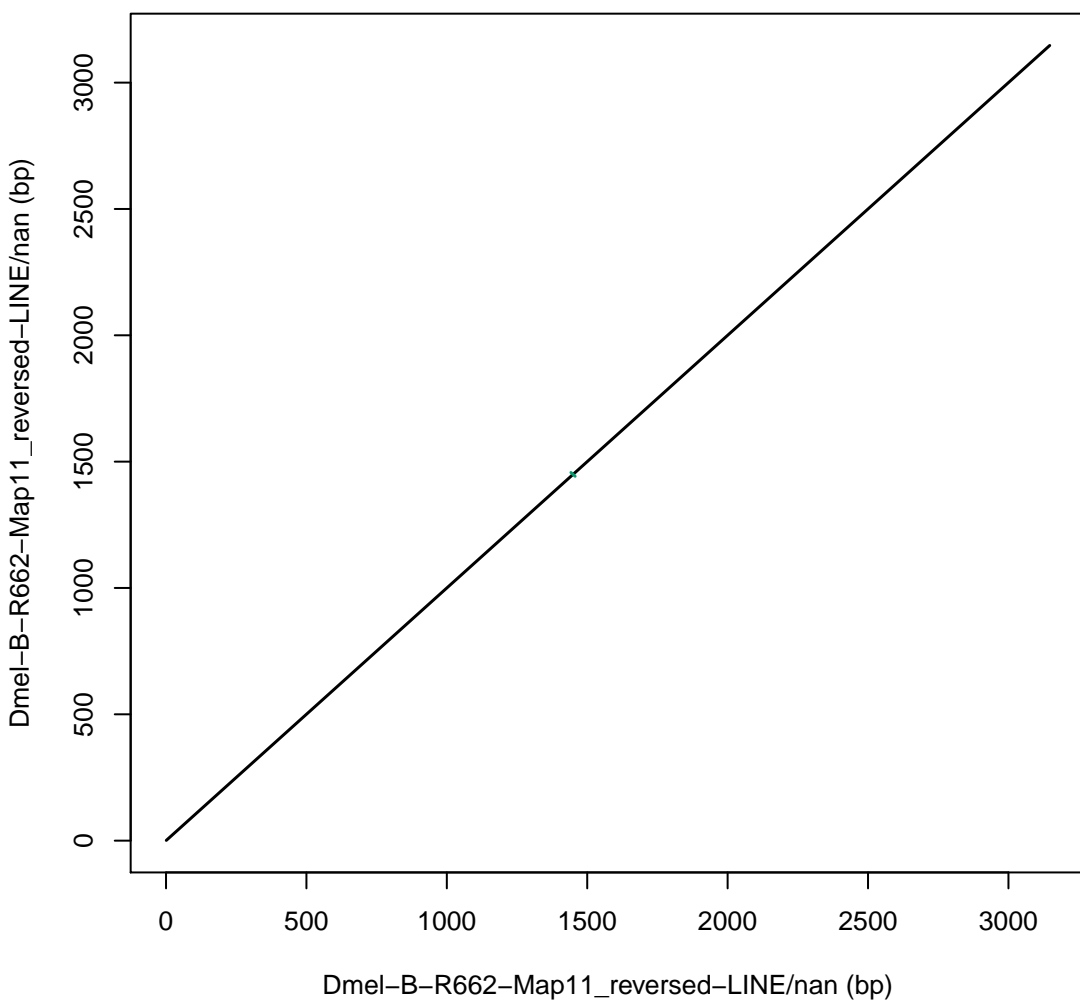
TE: Dmel-B-R662-Map11_reversed-LINE/nan
consensus size: 3147bp; fragments: 449; full length: 0 (>=2832.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

