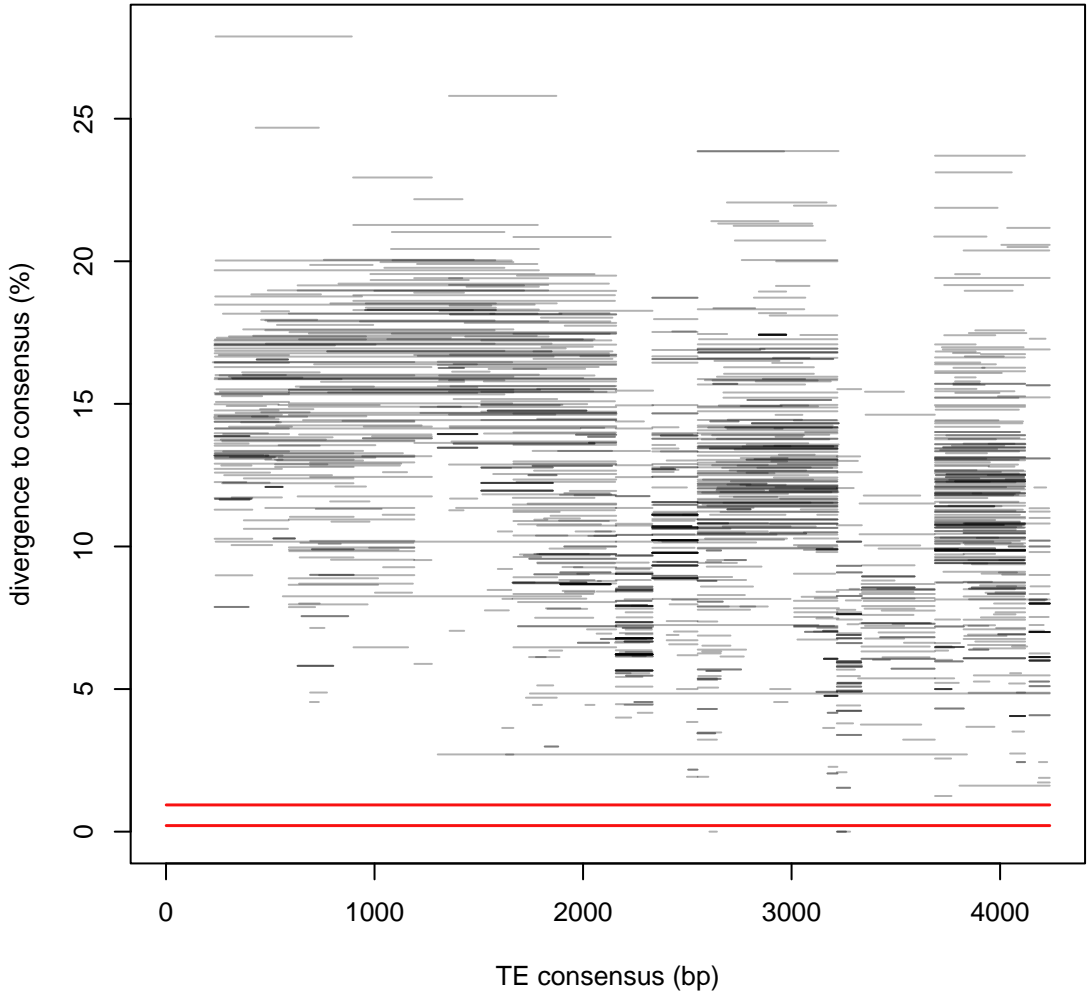
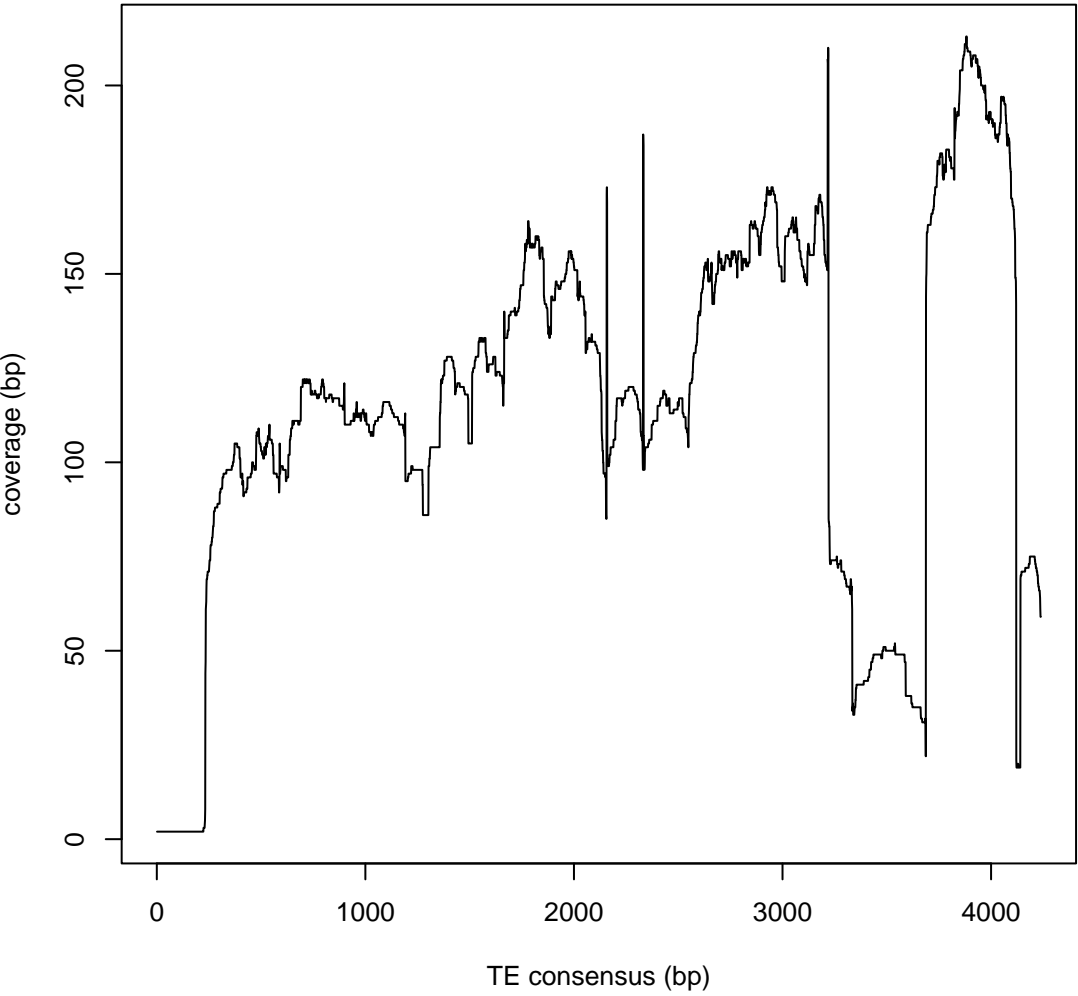


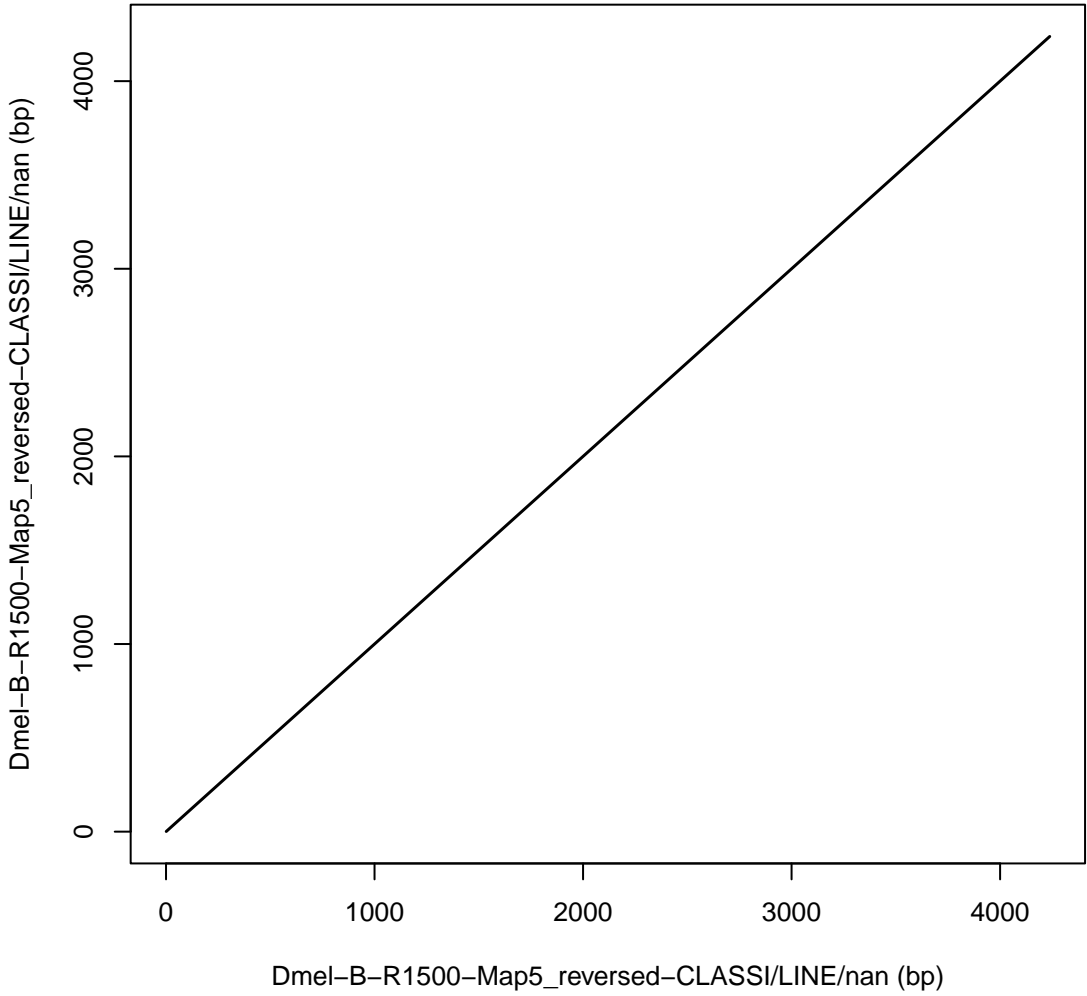
TE: Dmel-B-R1500-Map5_reversed-CLASSI/LINE/nan
consensus size: 4238bp; fragments: 1444; full length: 2 (>=3814.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

