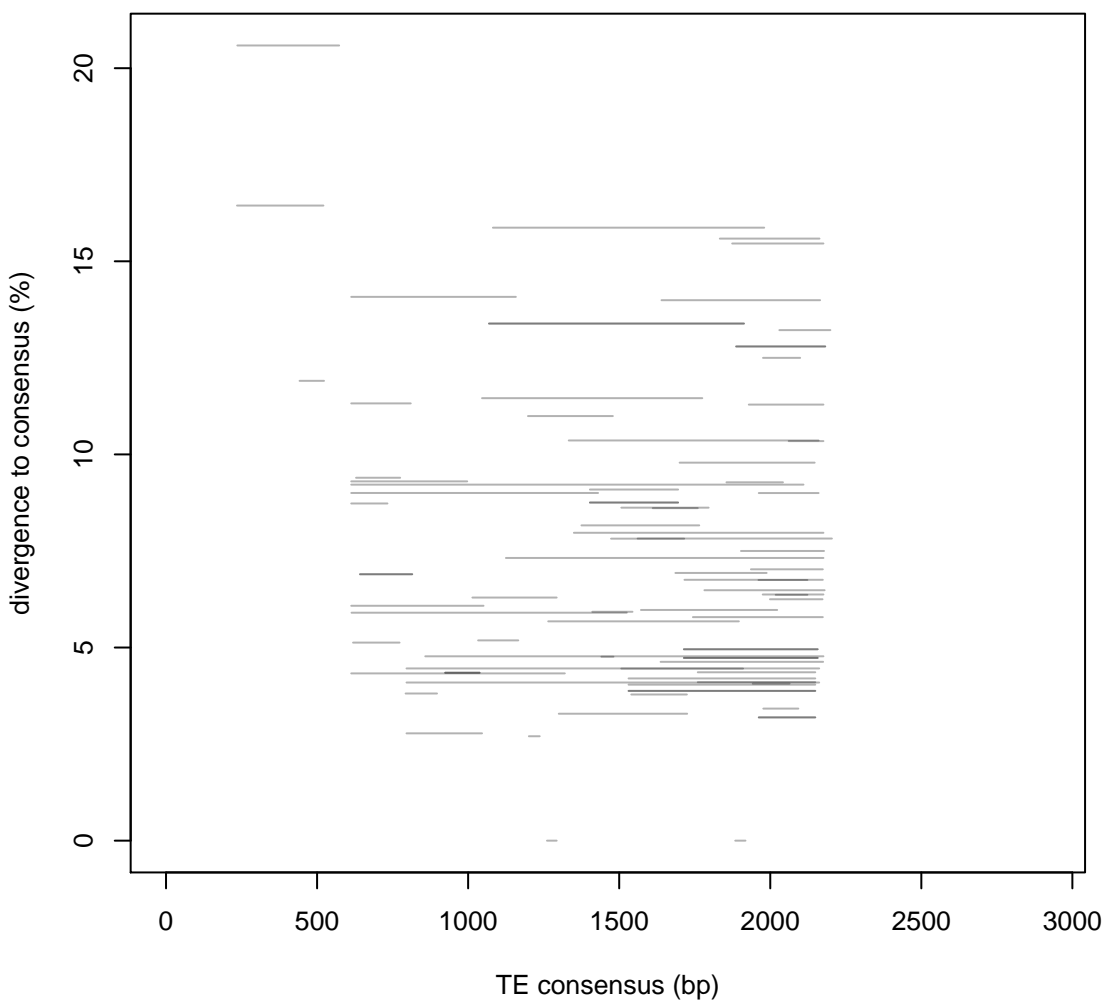
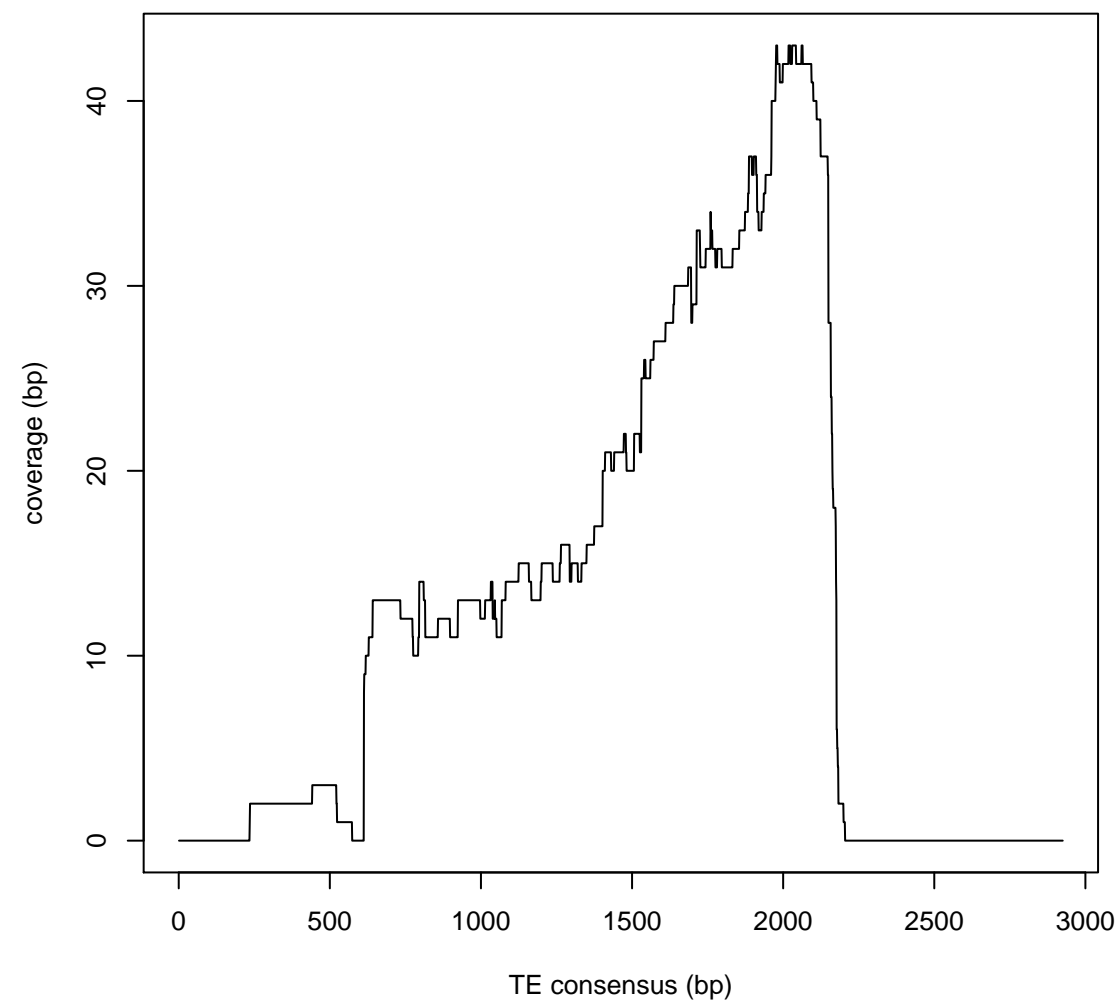


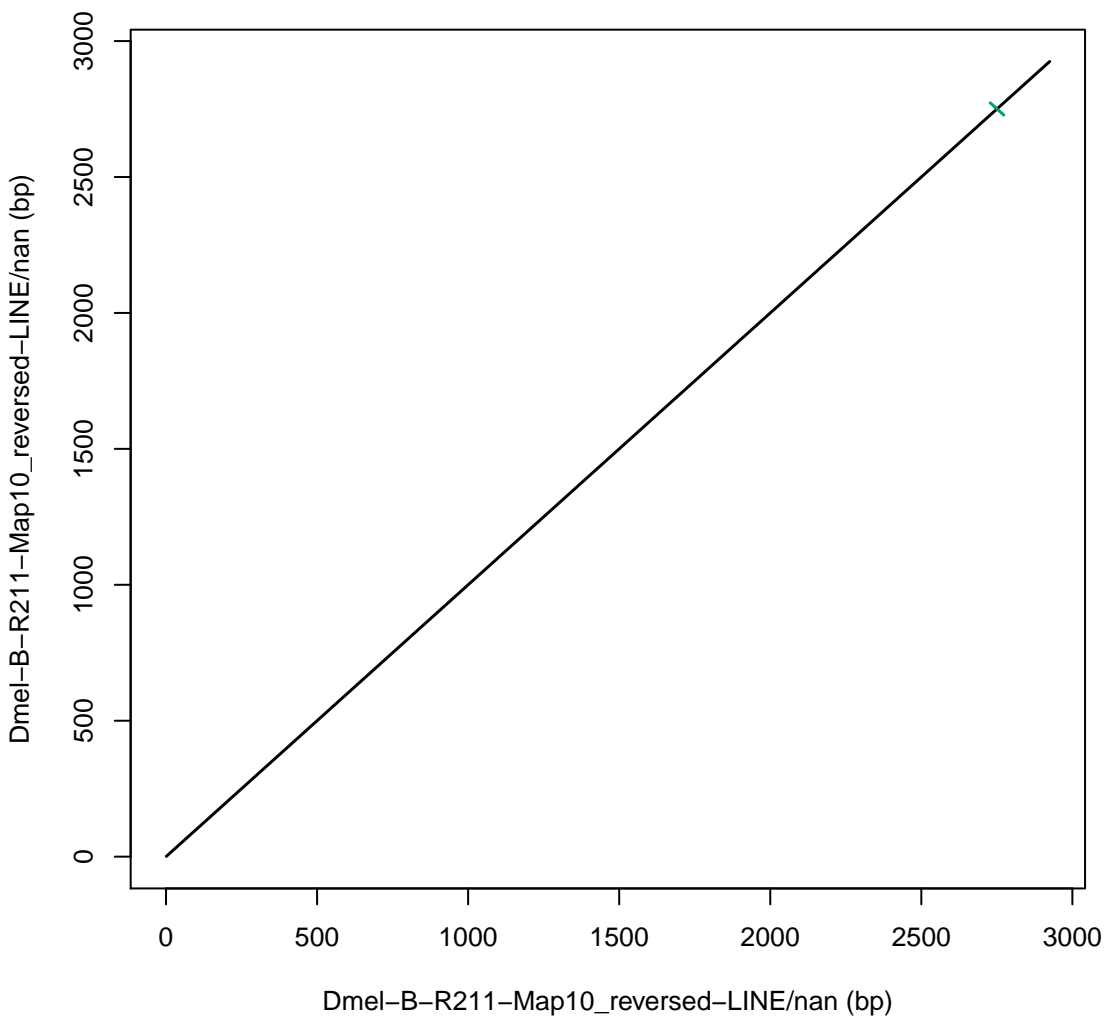
TE: Dmel-B-R211-Map10_reversed-LINE/nan
consensus size: 2925bp; fragments: 88; full length: 0 (>=2632.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

