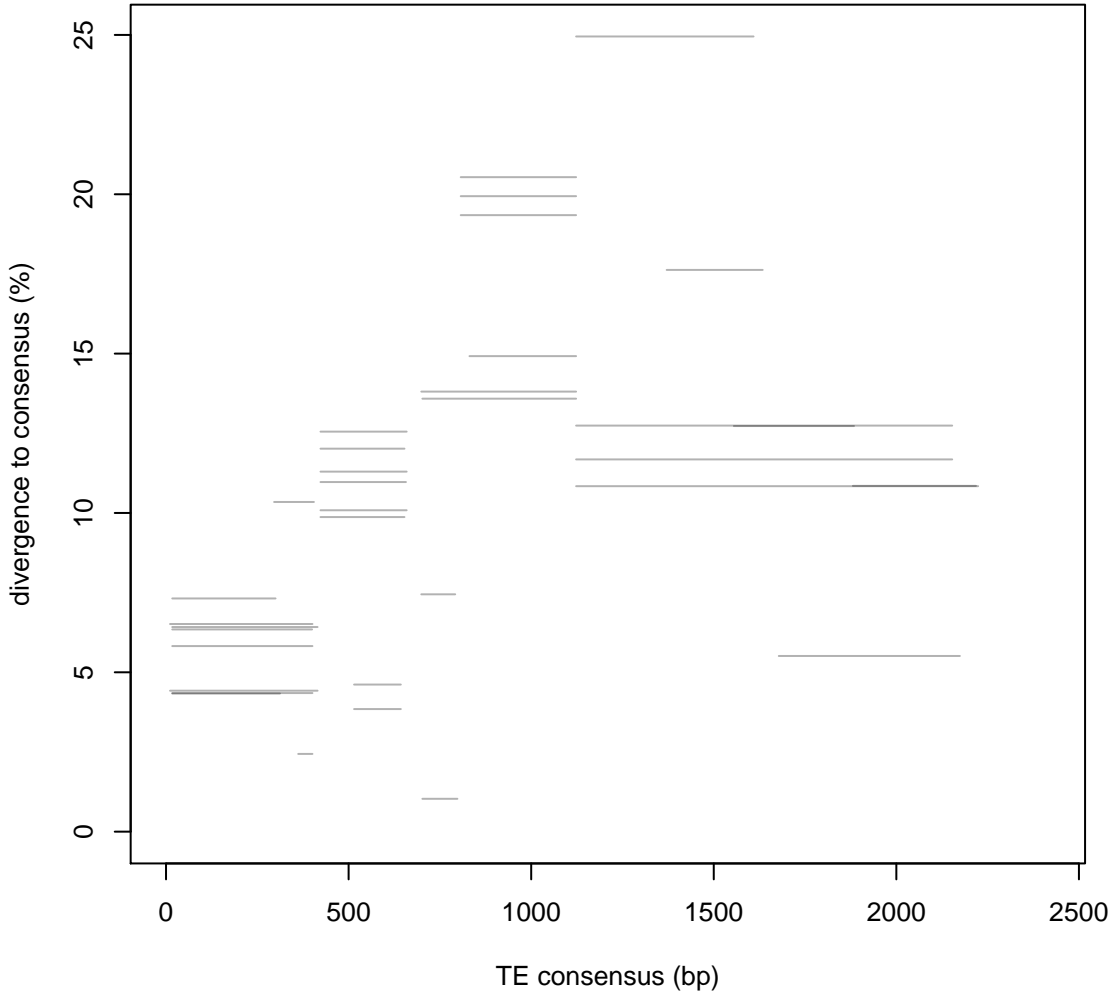
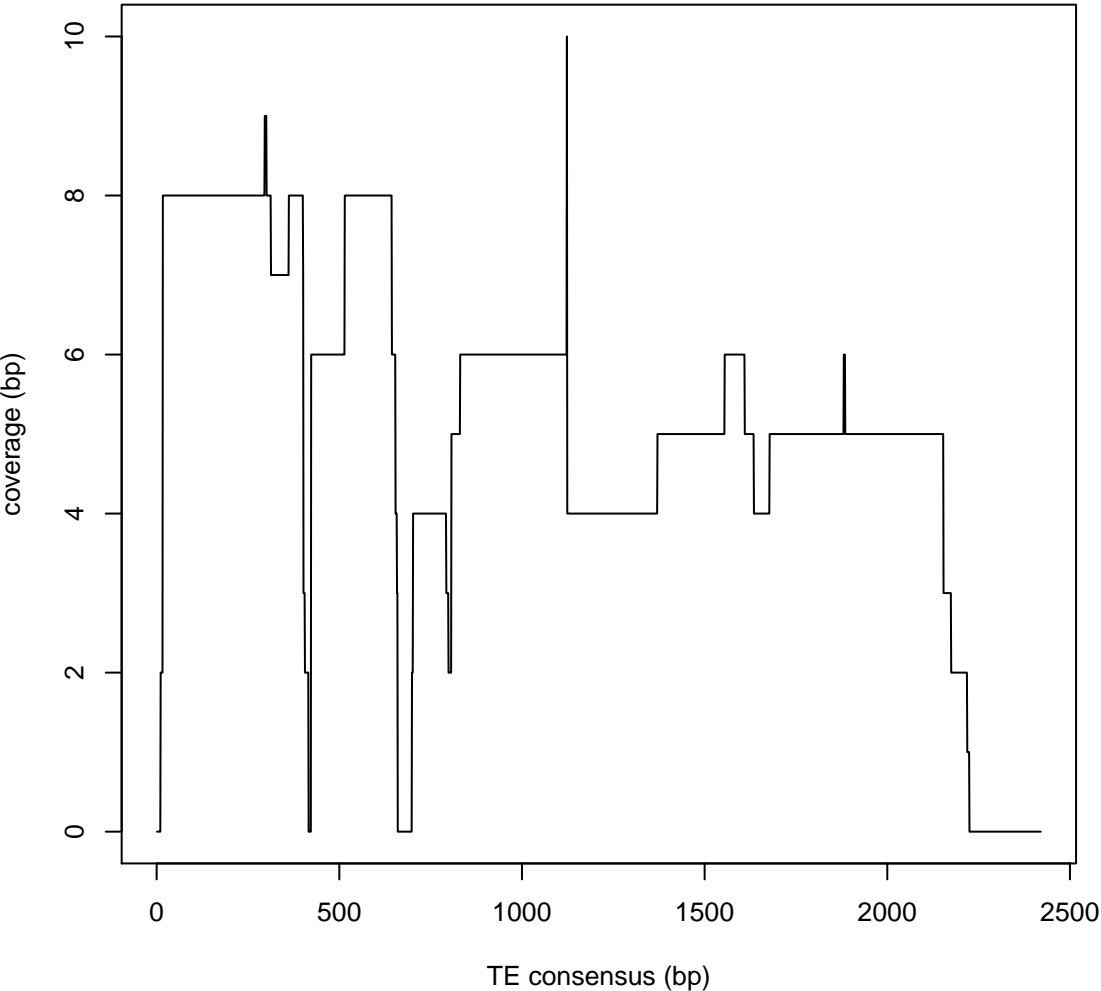


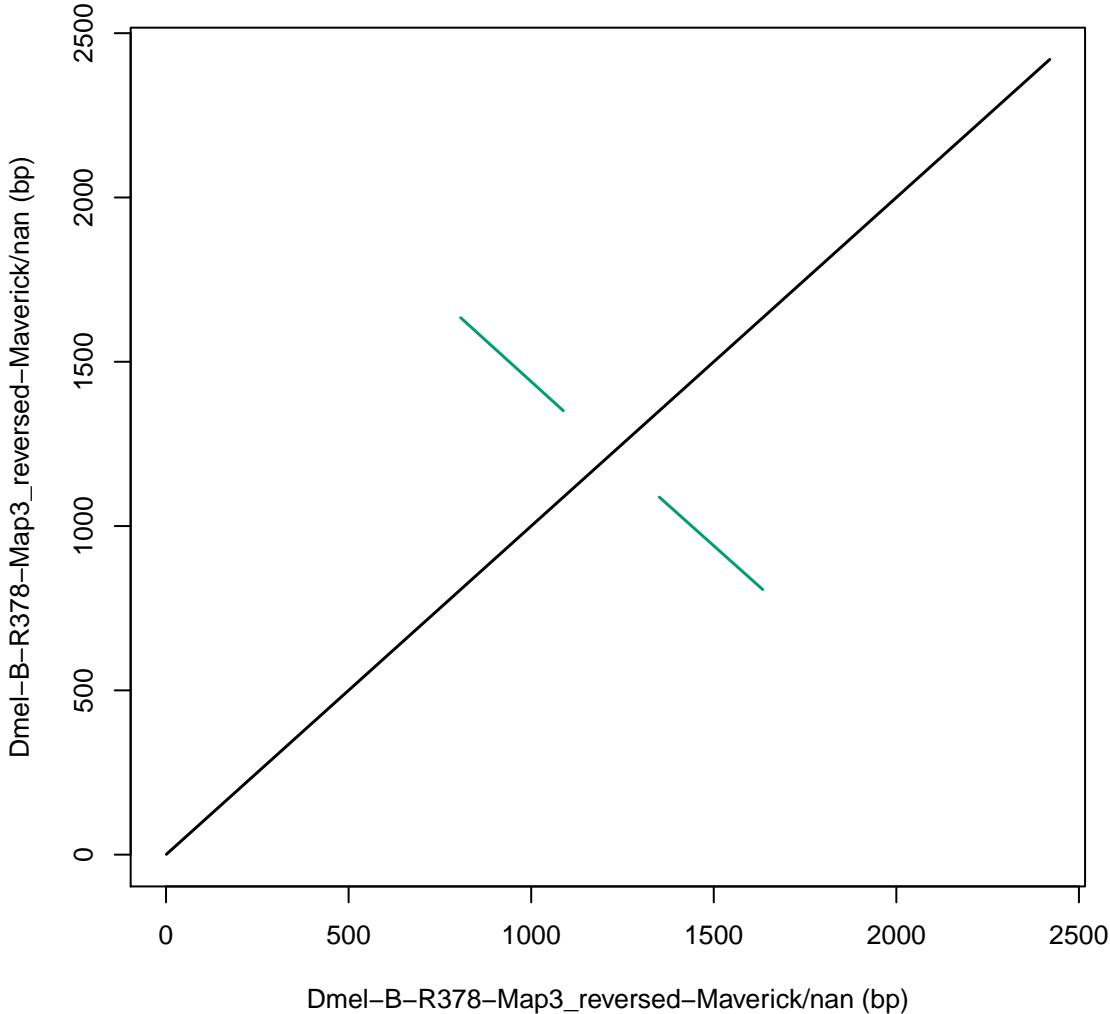
TE: Dmel-B-R378-Map3_reversed-Maverick/nan
consensus size: 2420bp; fragments: 34; full length: 0 (>=2178bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

