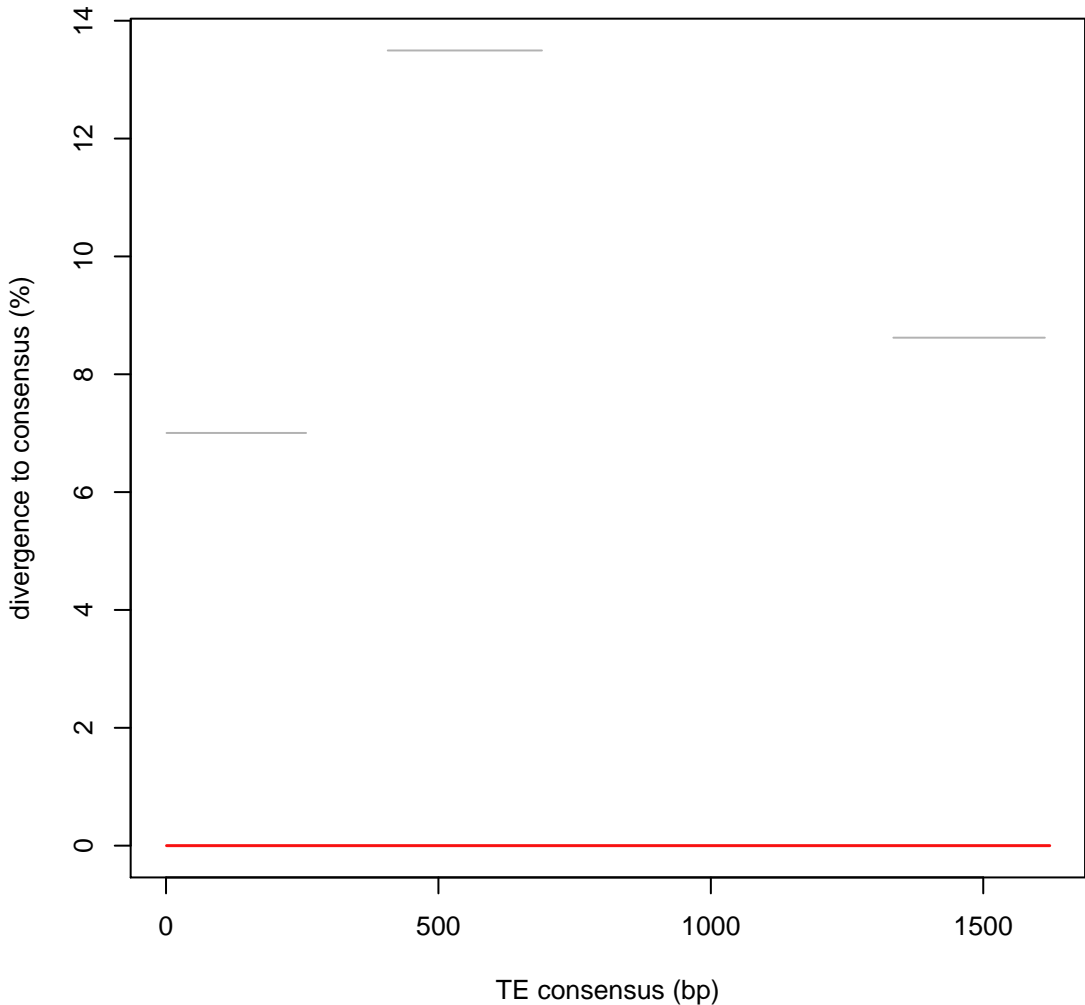
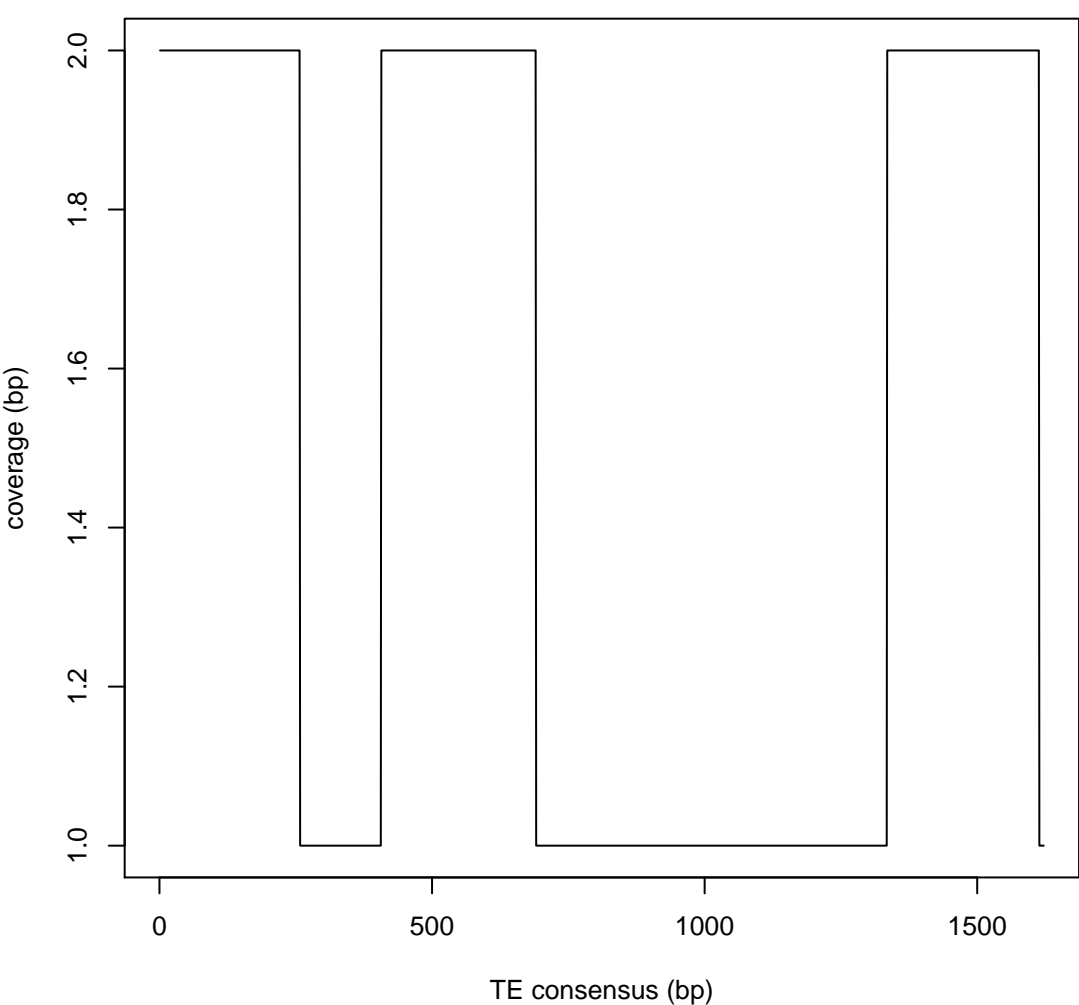


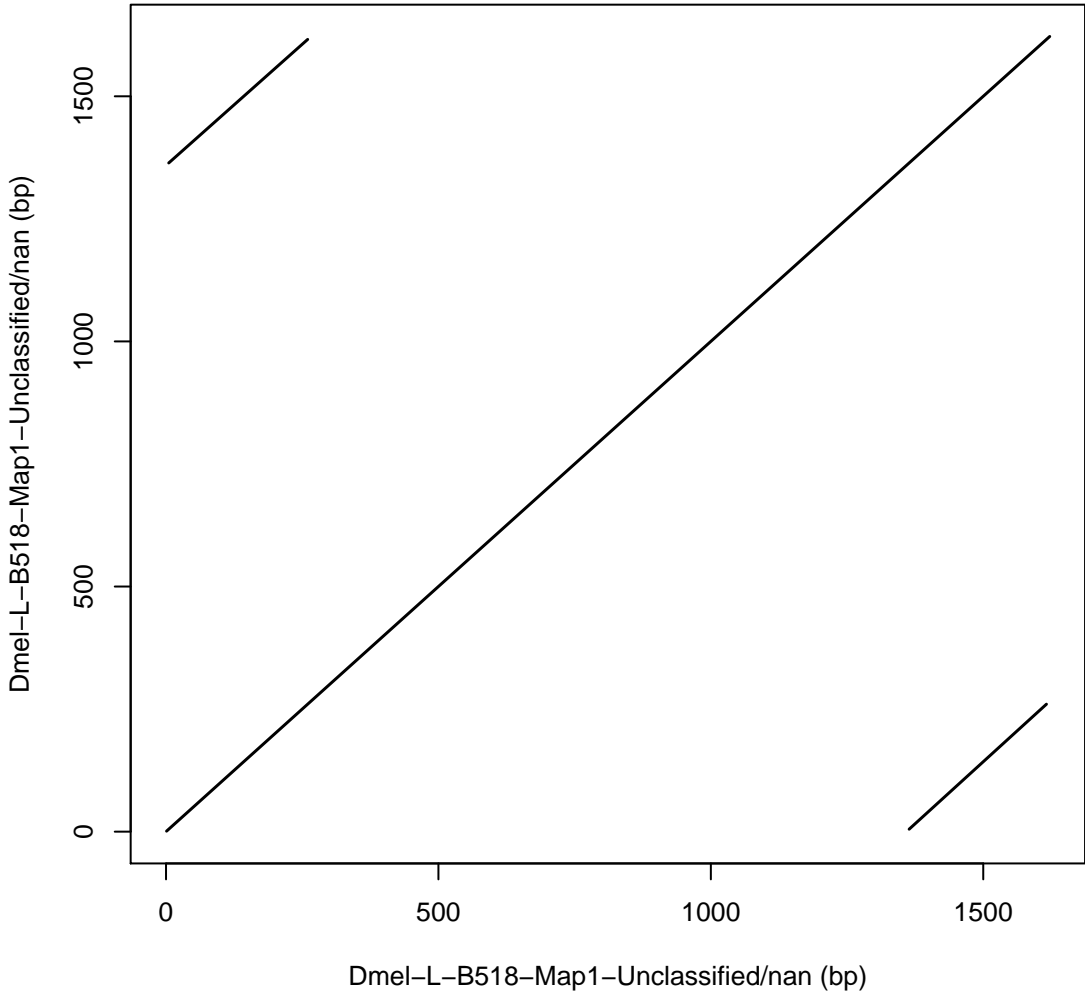
TE: Dmel-L-B518-Map1-Unclassified/nan  
consensus size: 1622bp; fragments: 4; full length: 1 (>=1459.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

