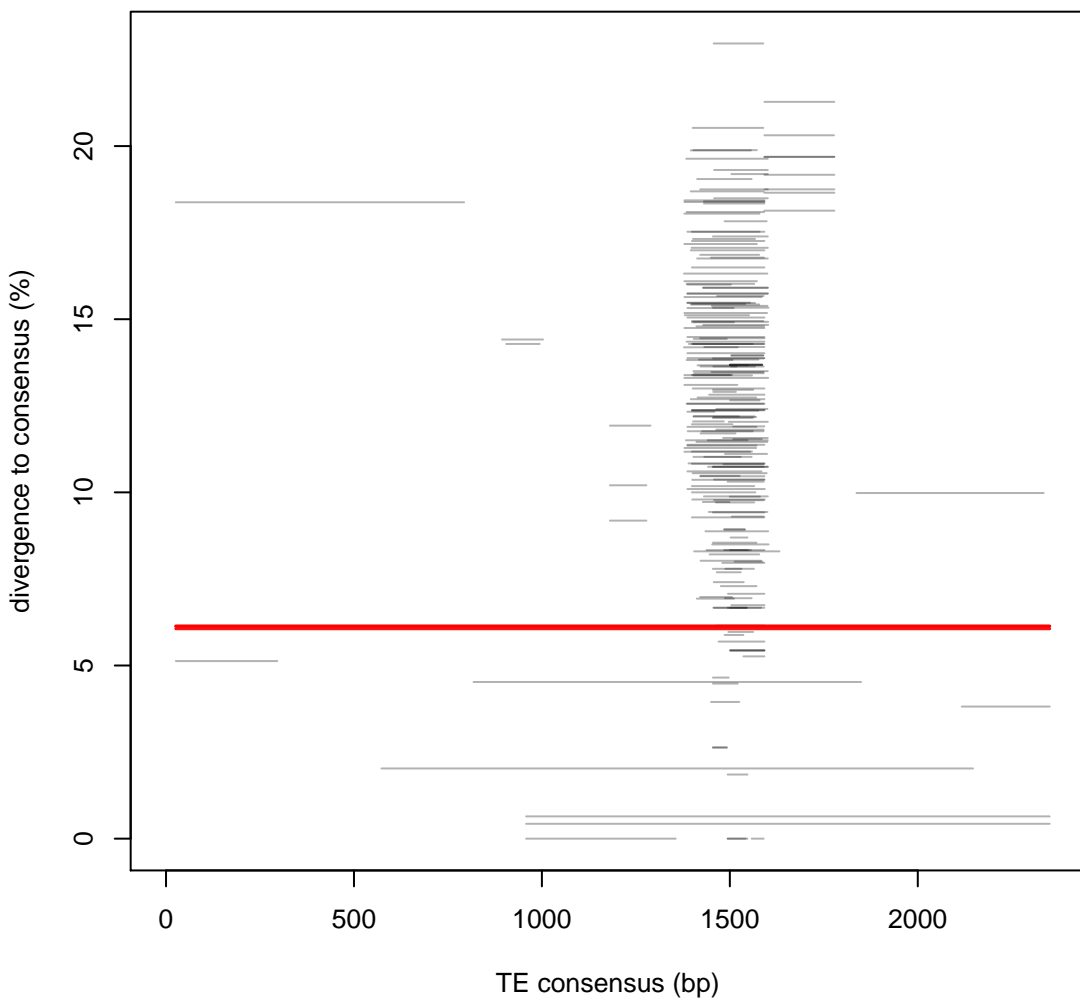
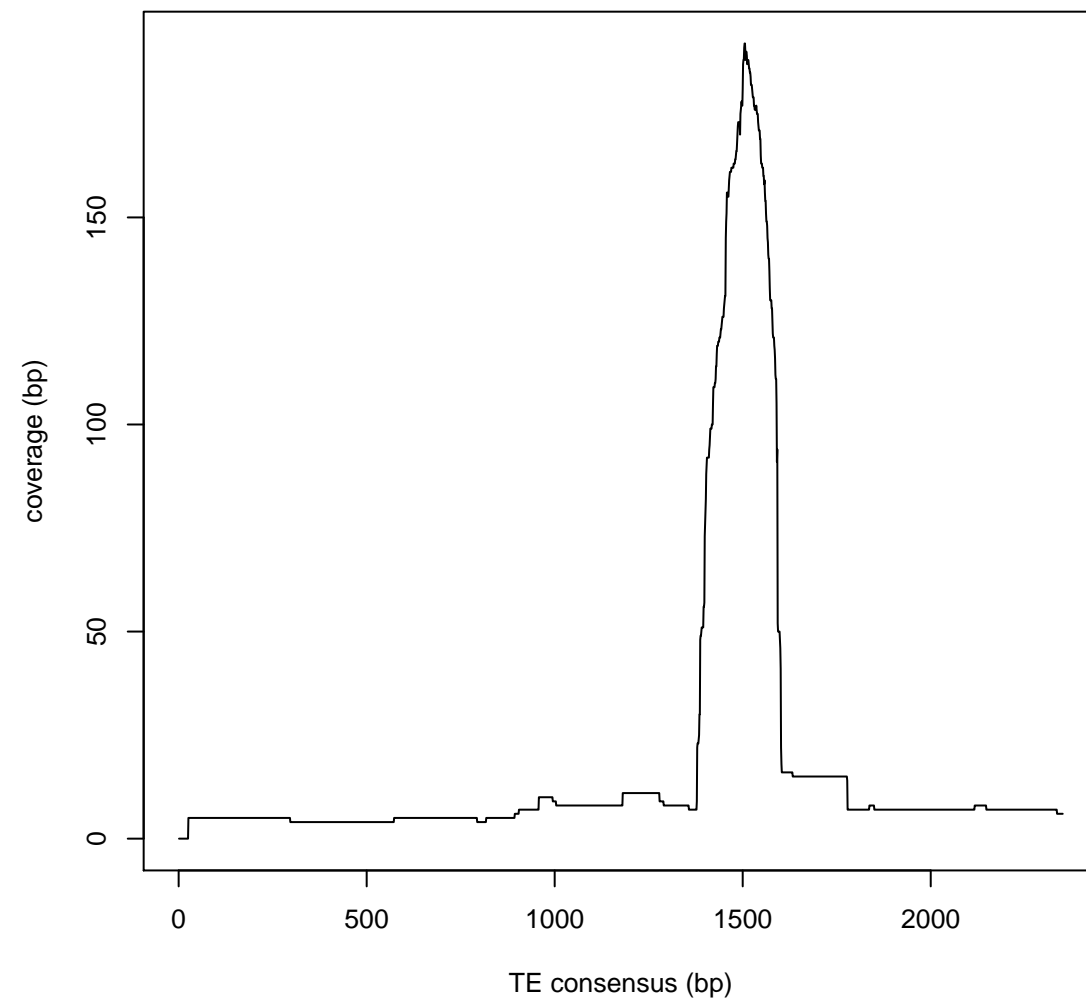


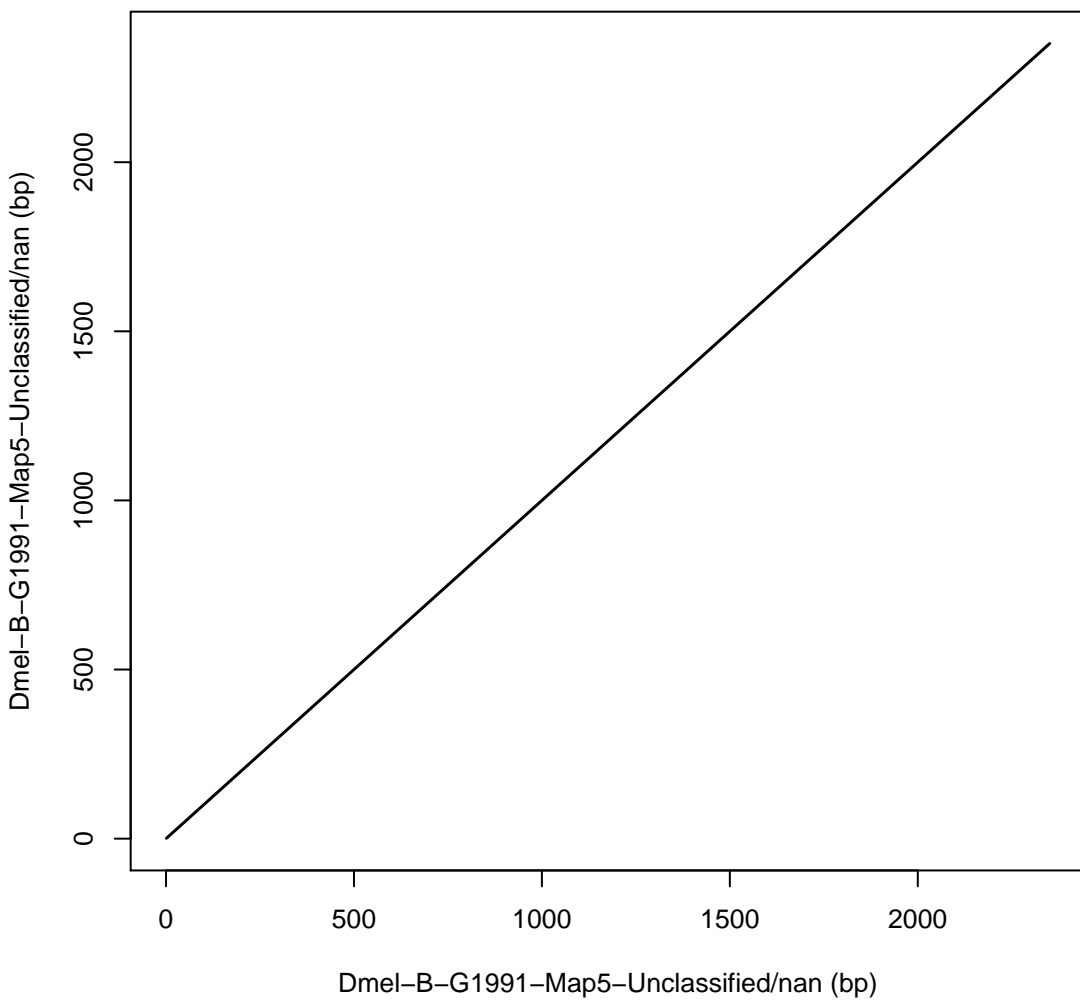
TE: Dmel-B-G1991-Map5-Unclassified/nan
consensus size: 2351bp; fragments: 224; full length: 3 (>=2115.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

