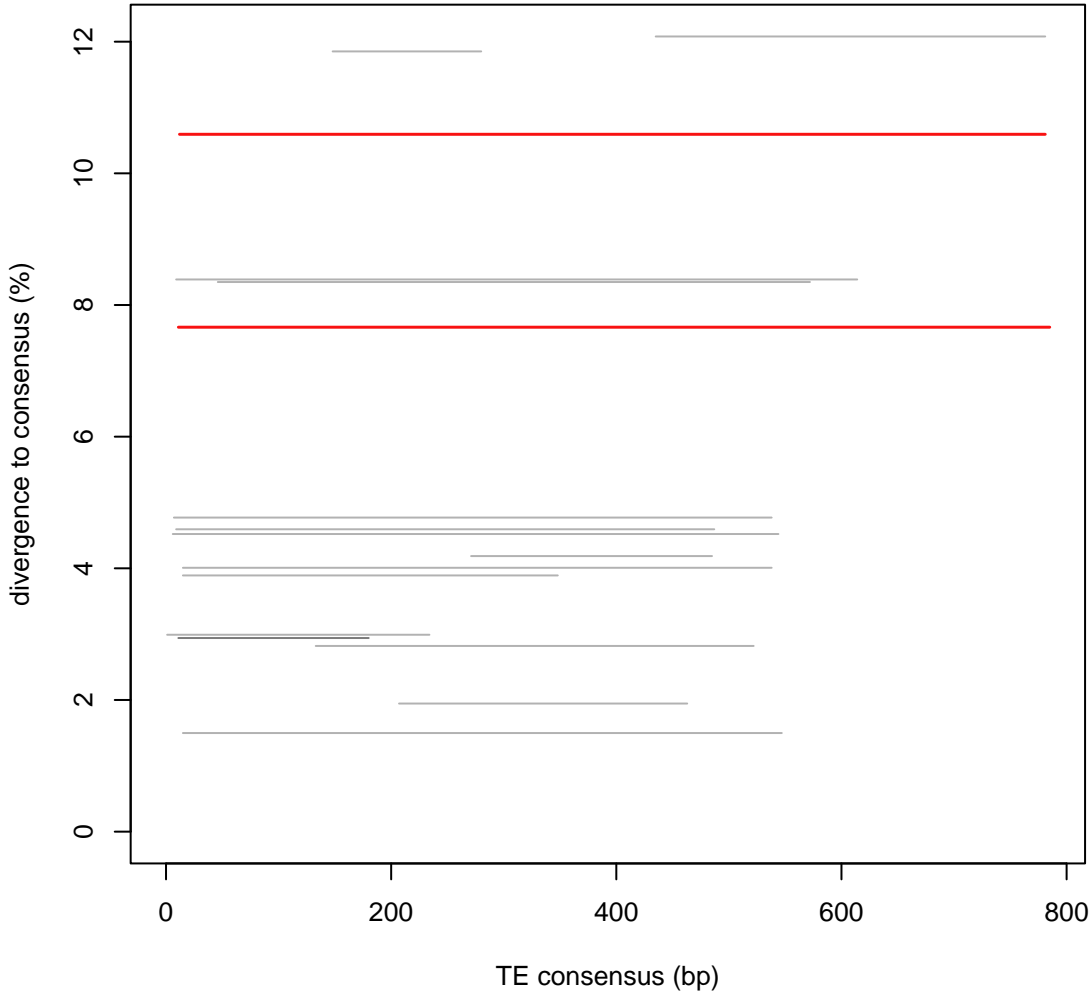
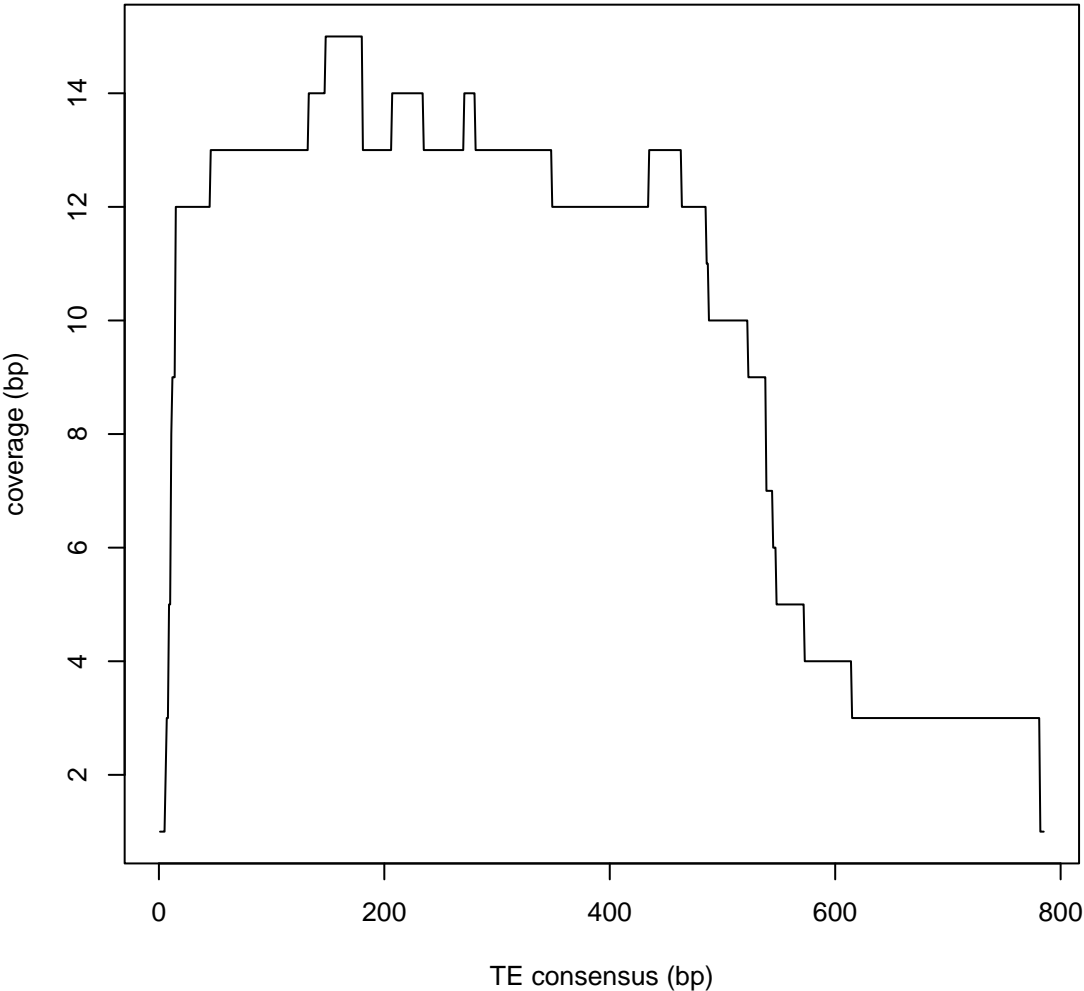


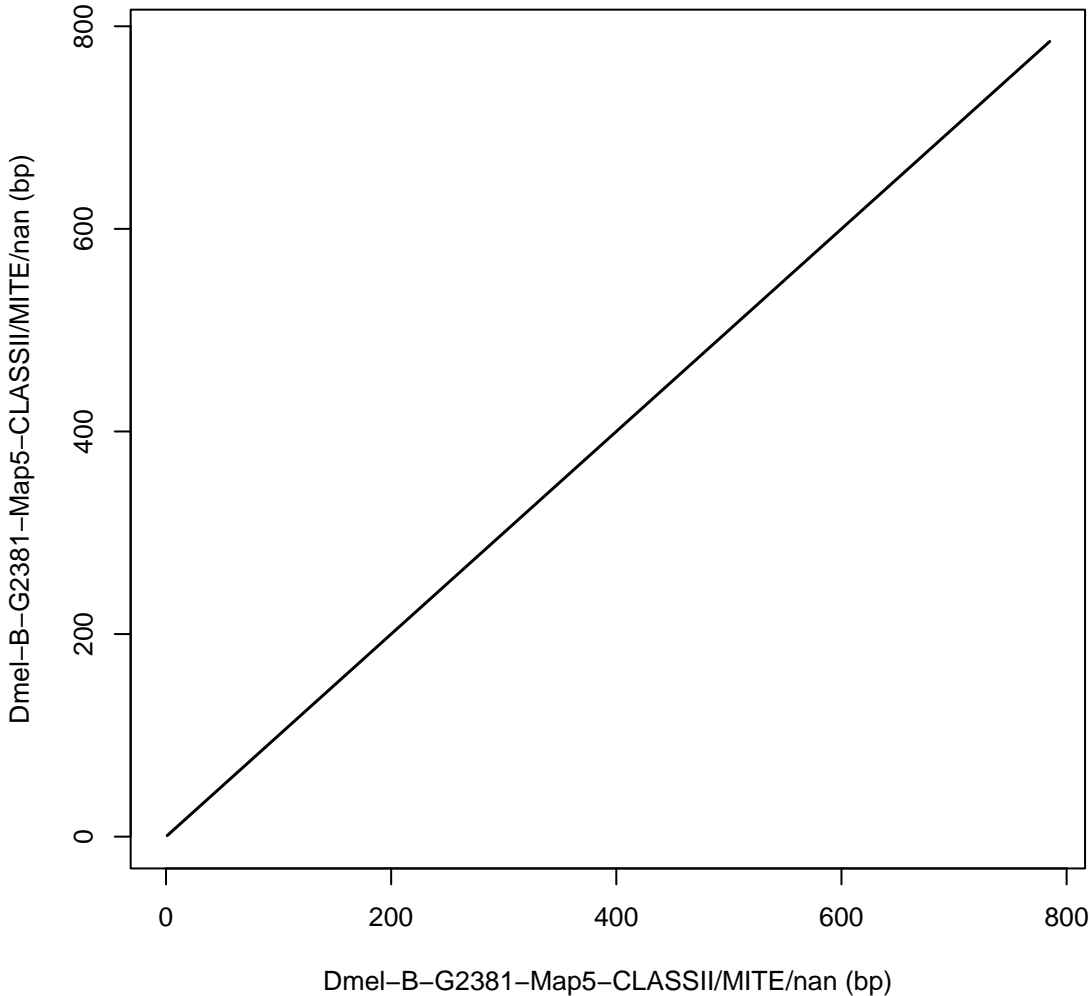
TE: Dmel-B-G2381-Map5-CLASSII/MITE/nan
consensus size: 785bp; fragments: 18; full length: 2 (>=706.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

