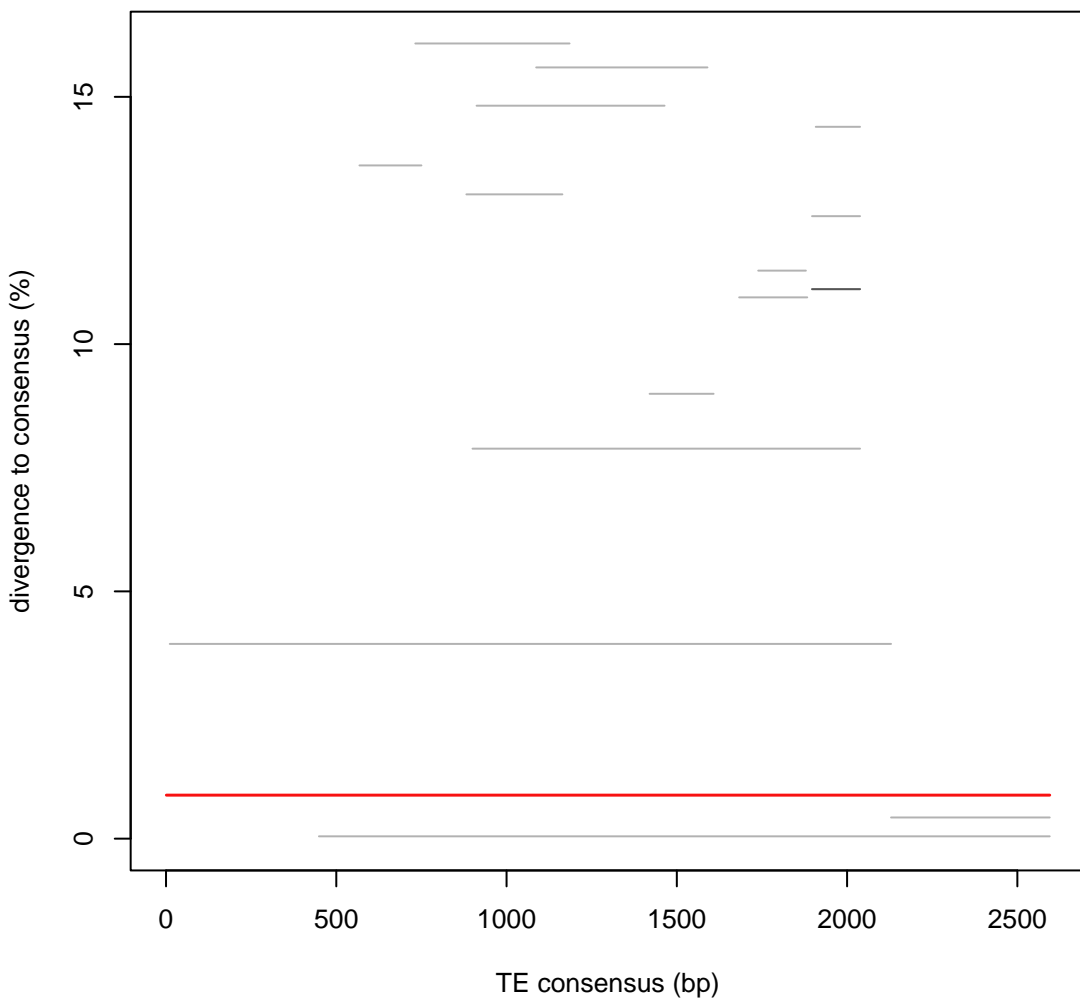
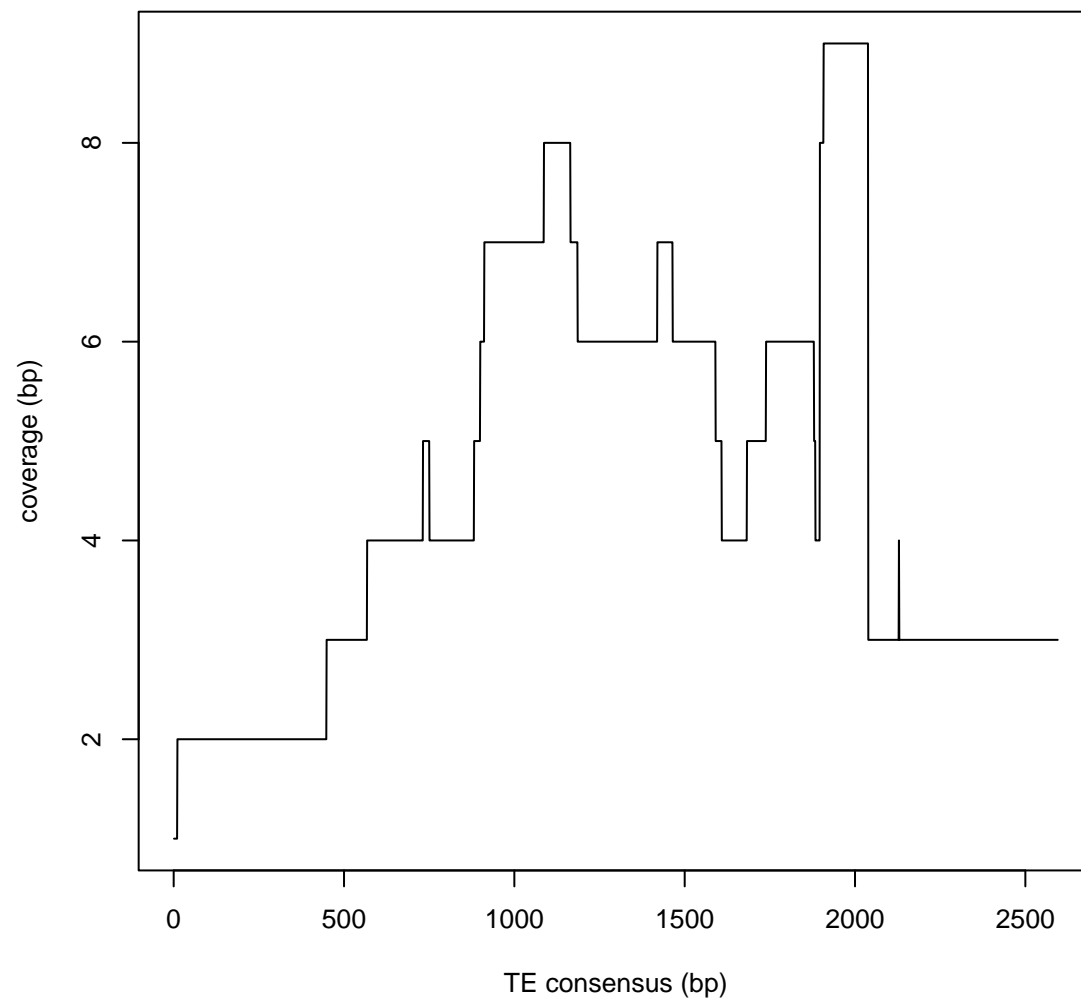


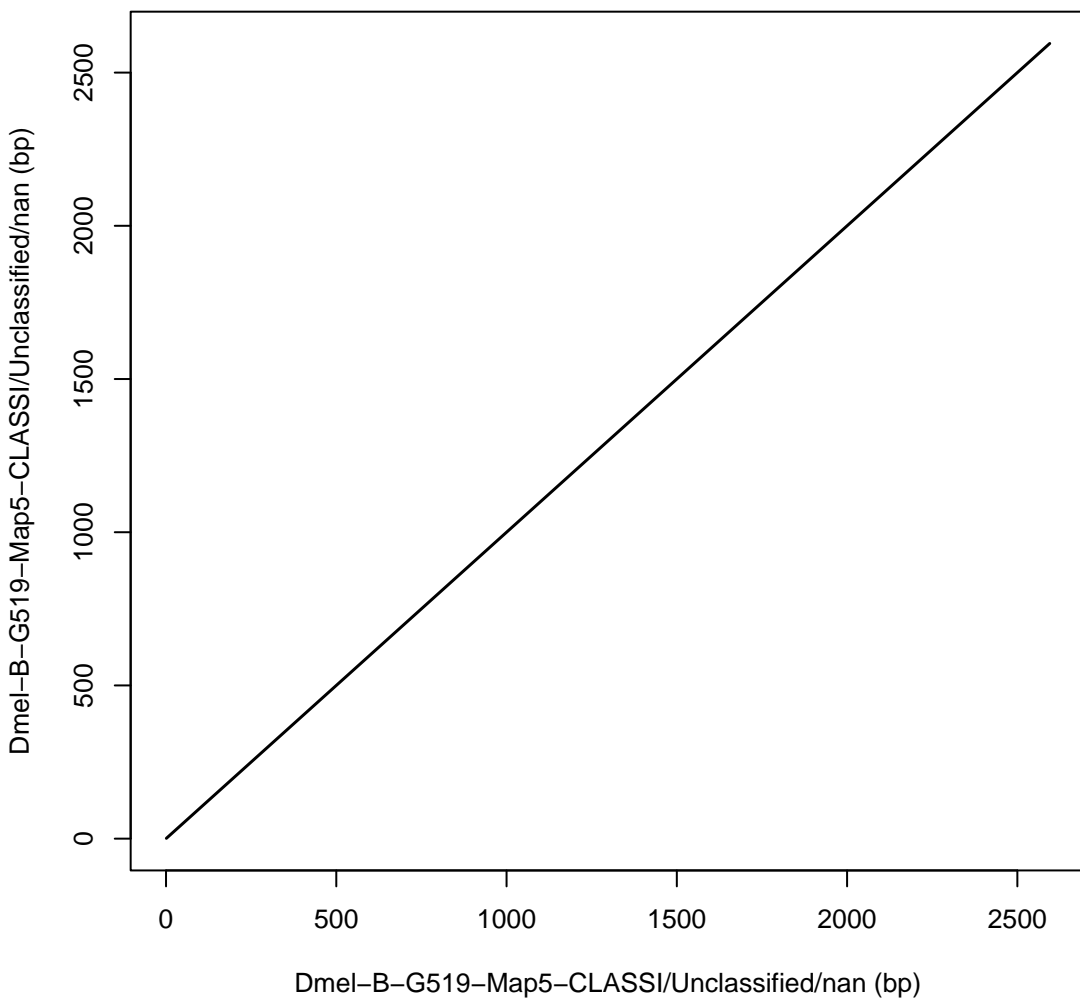
TE: Dmel-B-G519-Map5-CLASSI/Unclassified/nan
consensus size: 2595bp; fragments: 18; full length: 1 (>=2335.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

