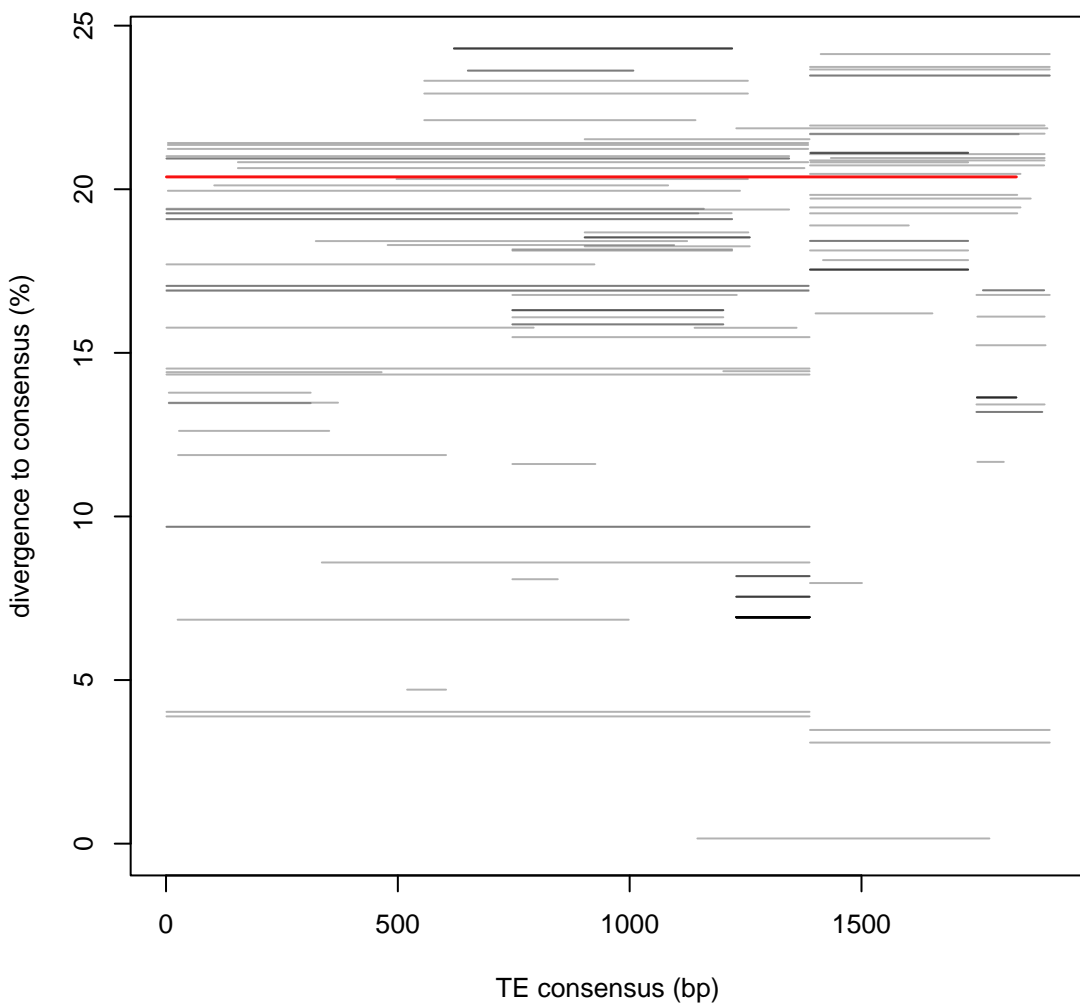
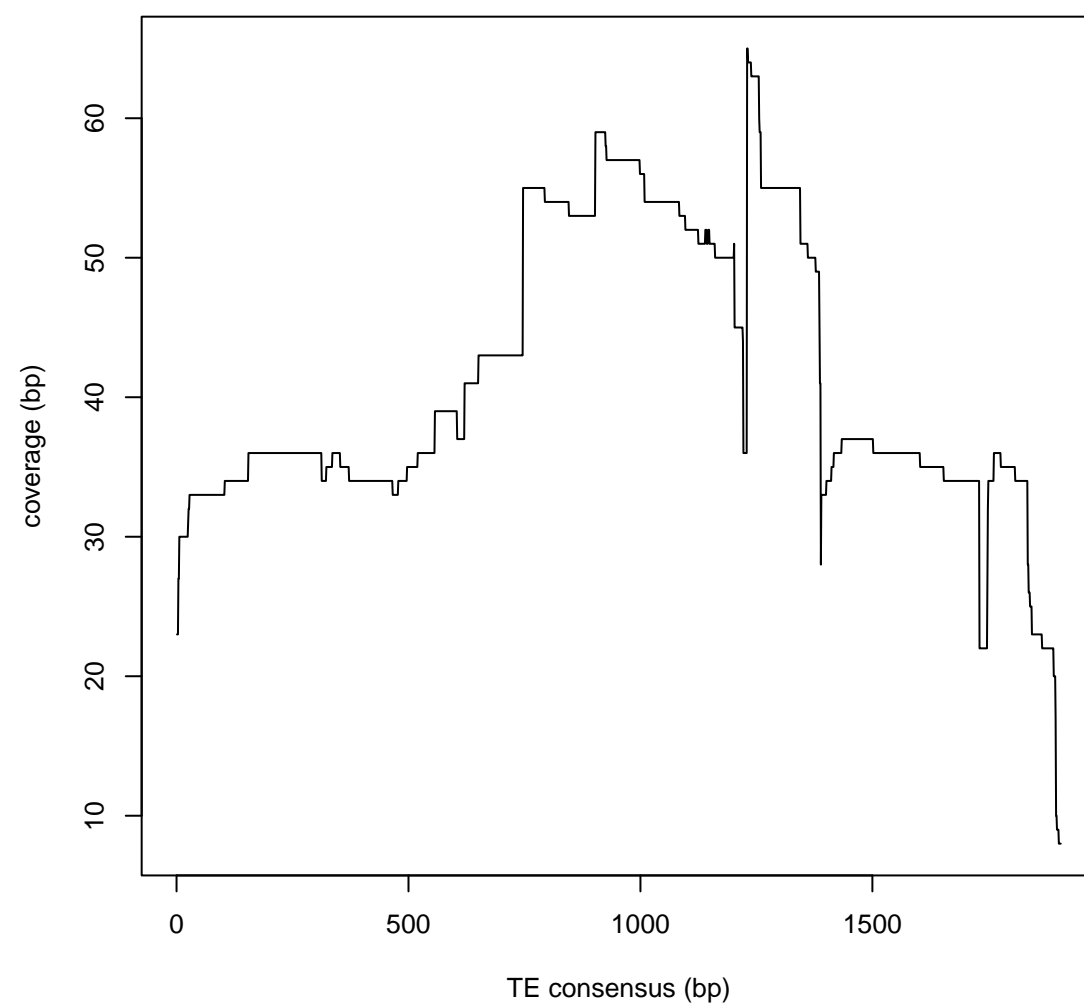


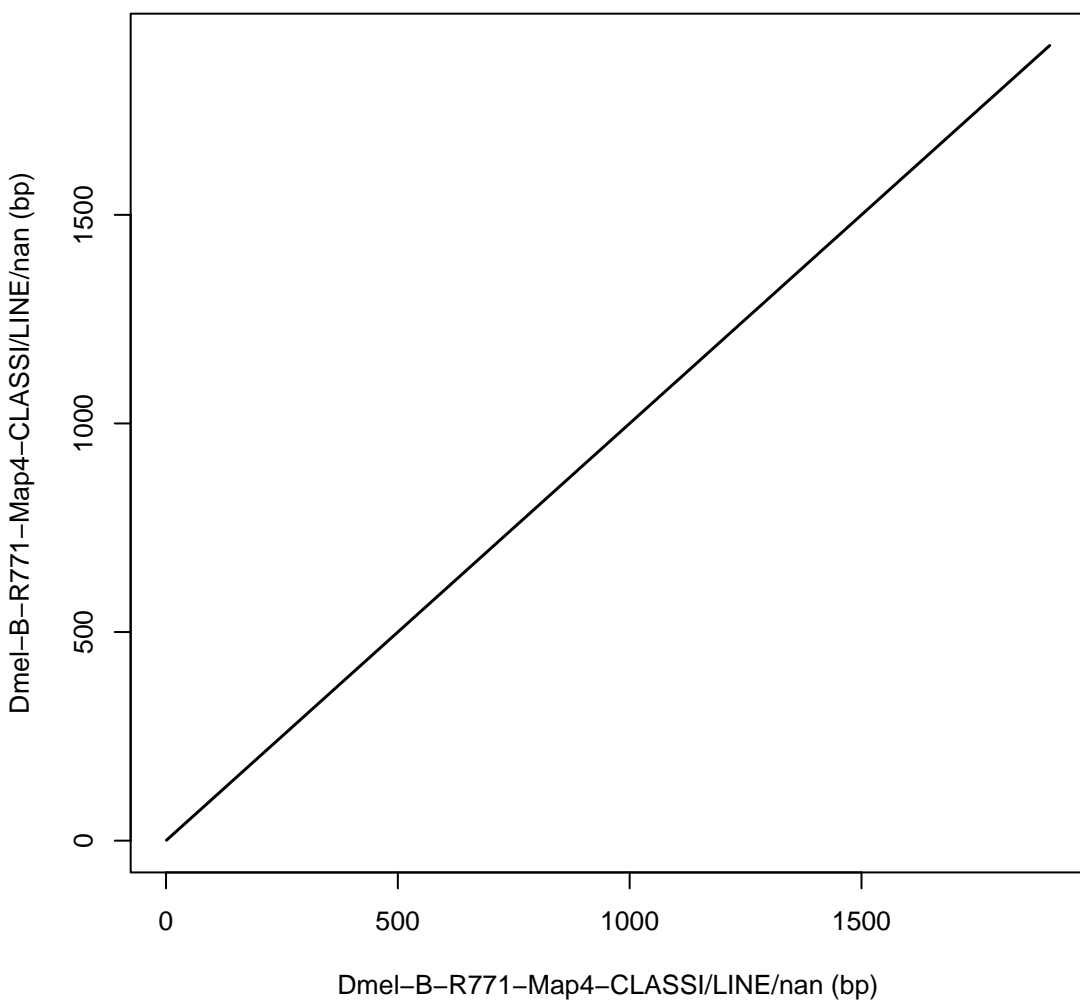
TE: Dmel-B-R771-Map4-CLASSI/LINE/nan
consensus size: 1906bp; fragments: 148; full length: 1 (>=1715.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

