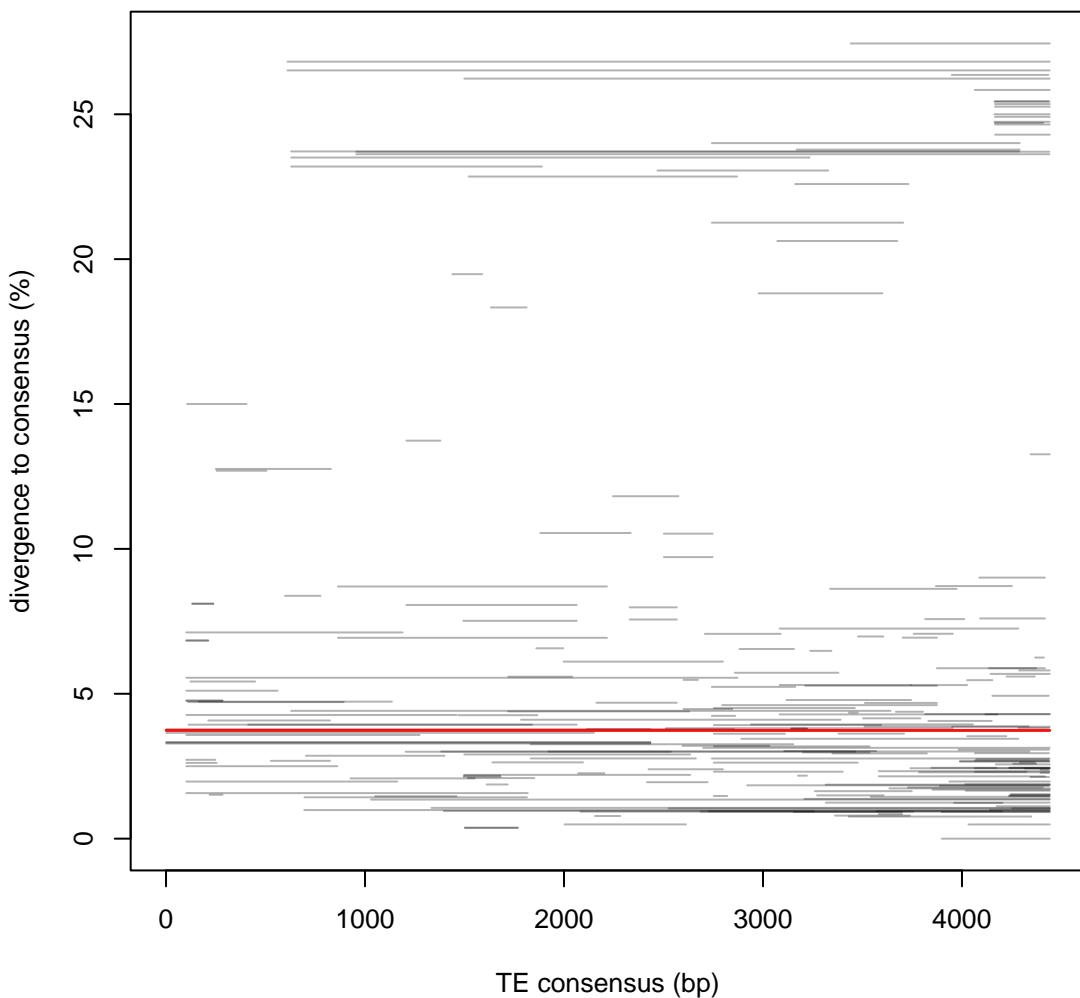
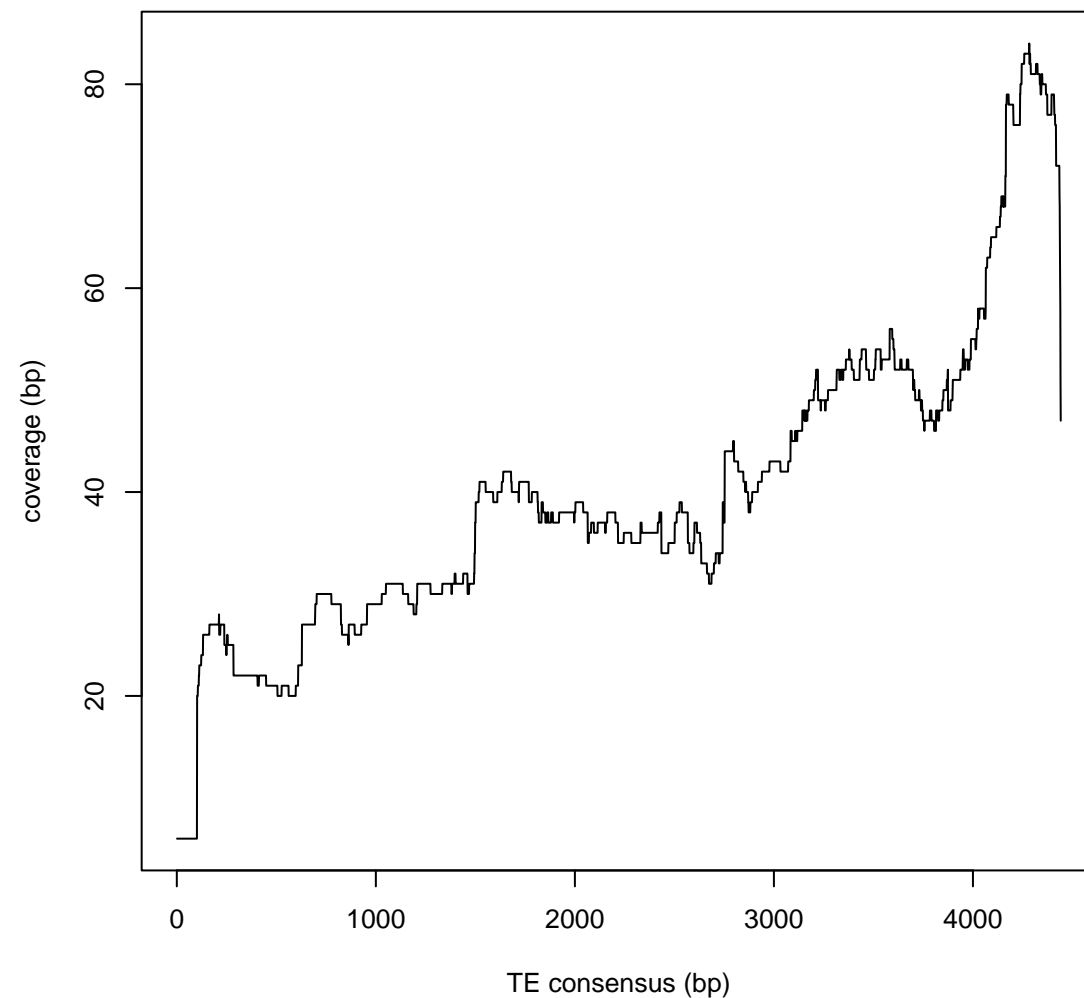


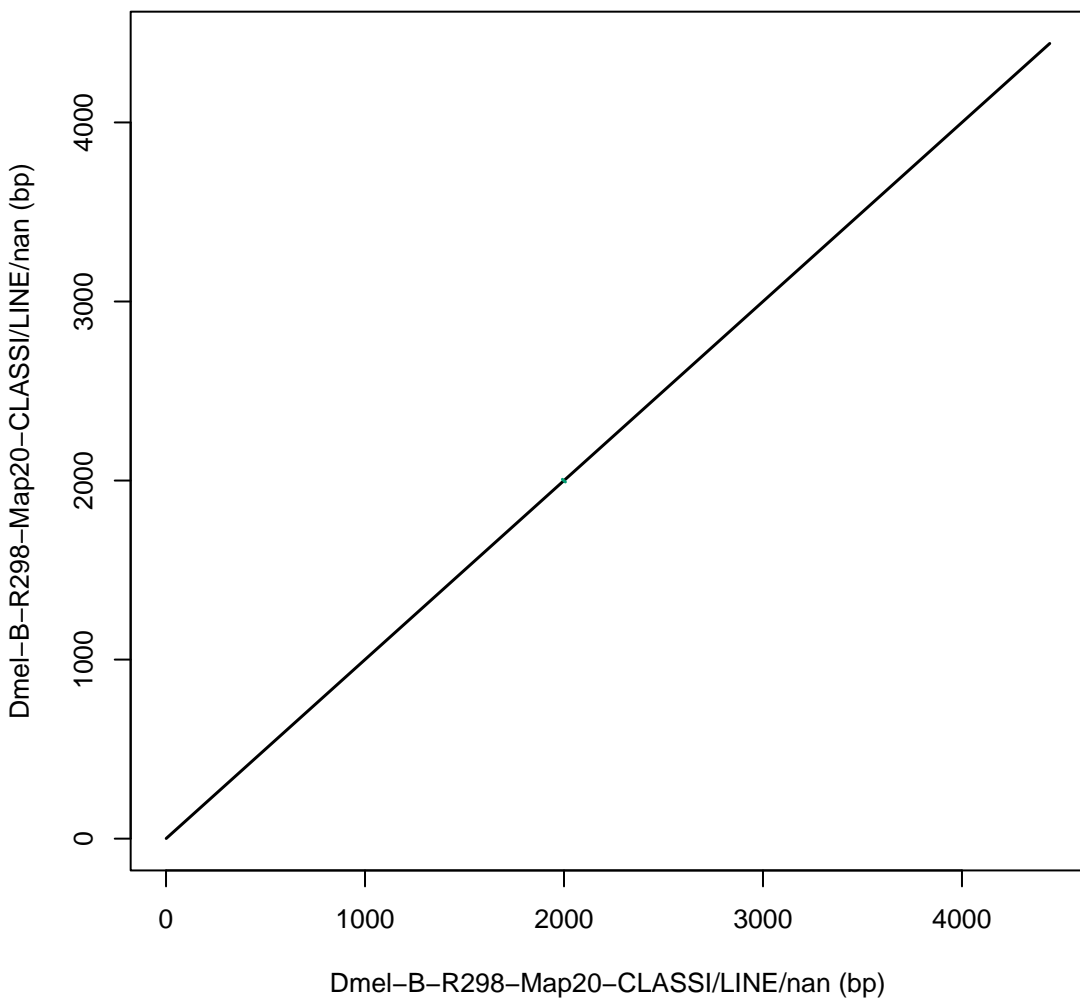
TE: Dmel-B-R298-Map20-CLASSI/LINE/nan  
consensus size: 4441bp; fragments: 231; full length: 1 (>=3996.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

