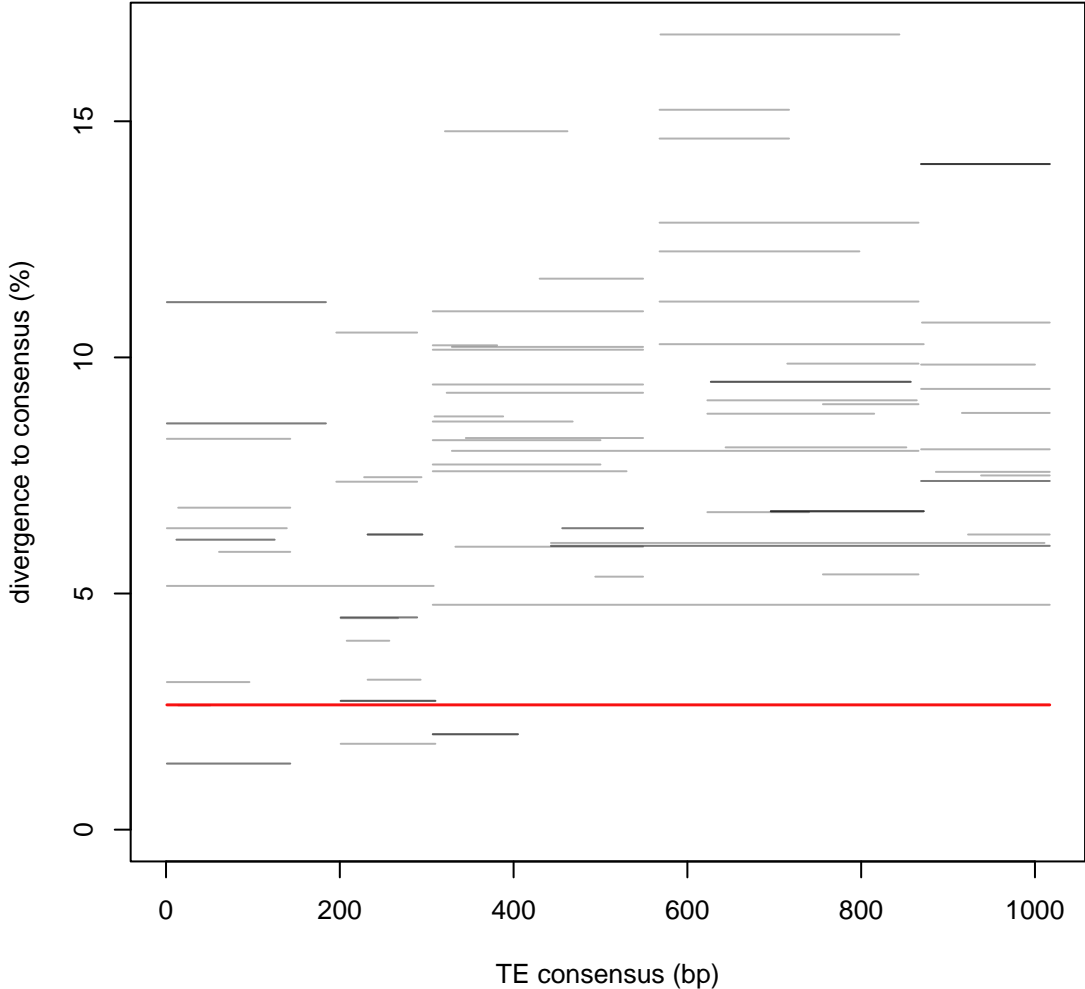
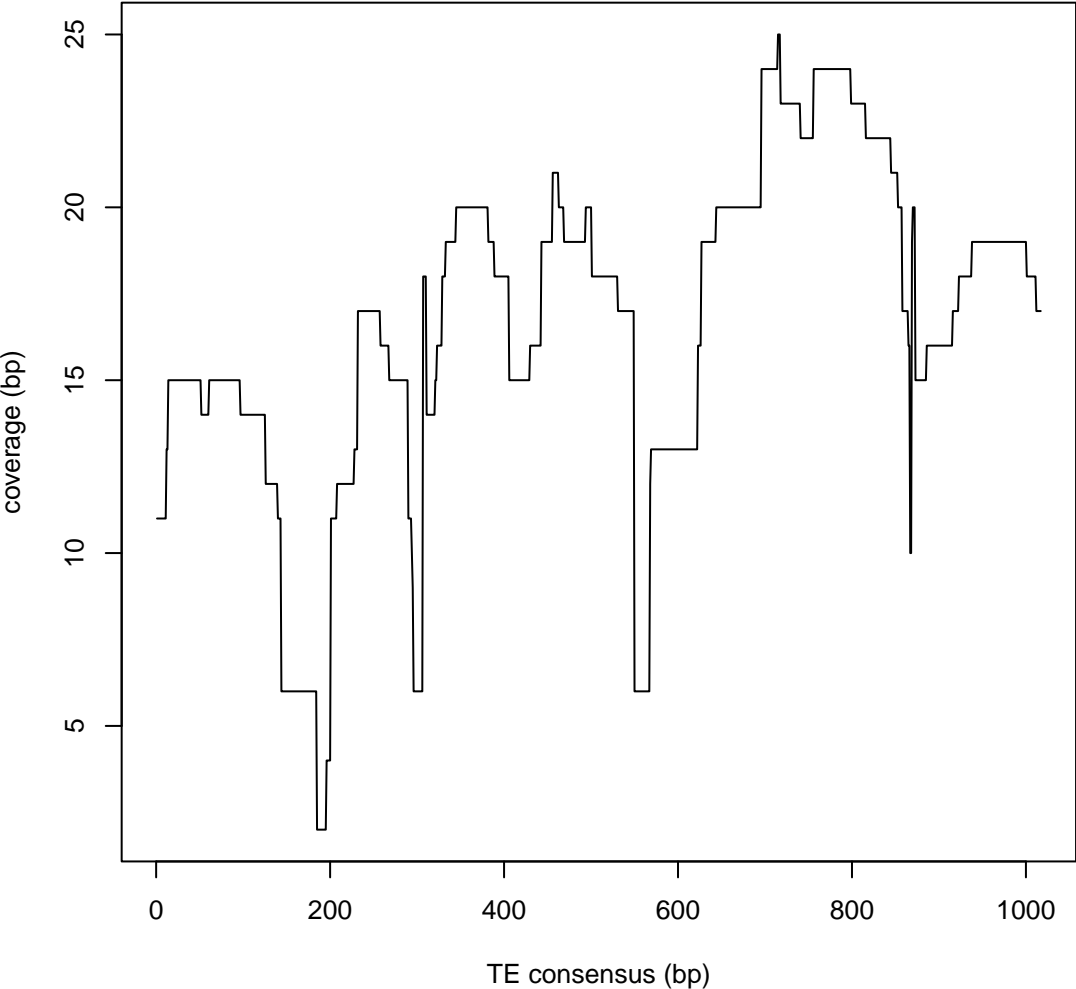


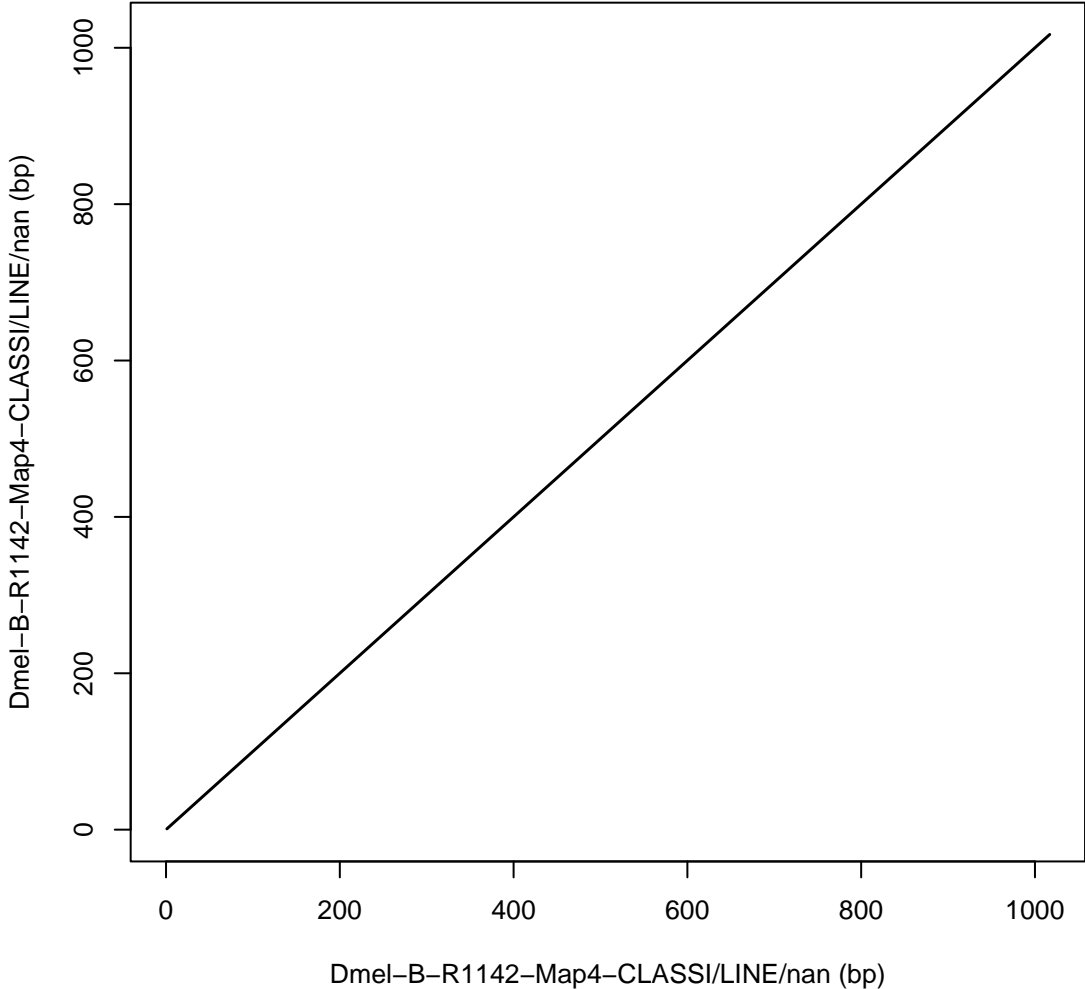
TE: Dmel-B-R1142-Map4-CLASSI/LINE/nan
consensus size: 1017bp; fragments: 92; full length: 1 (>=915.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

