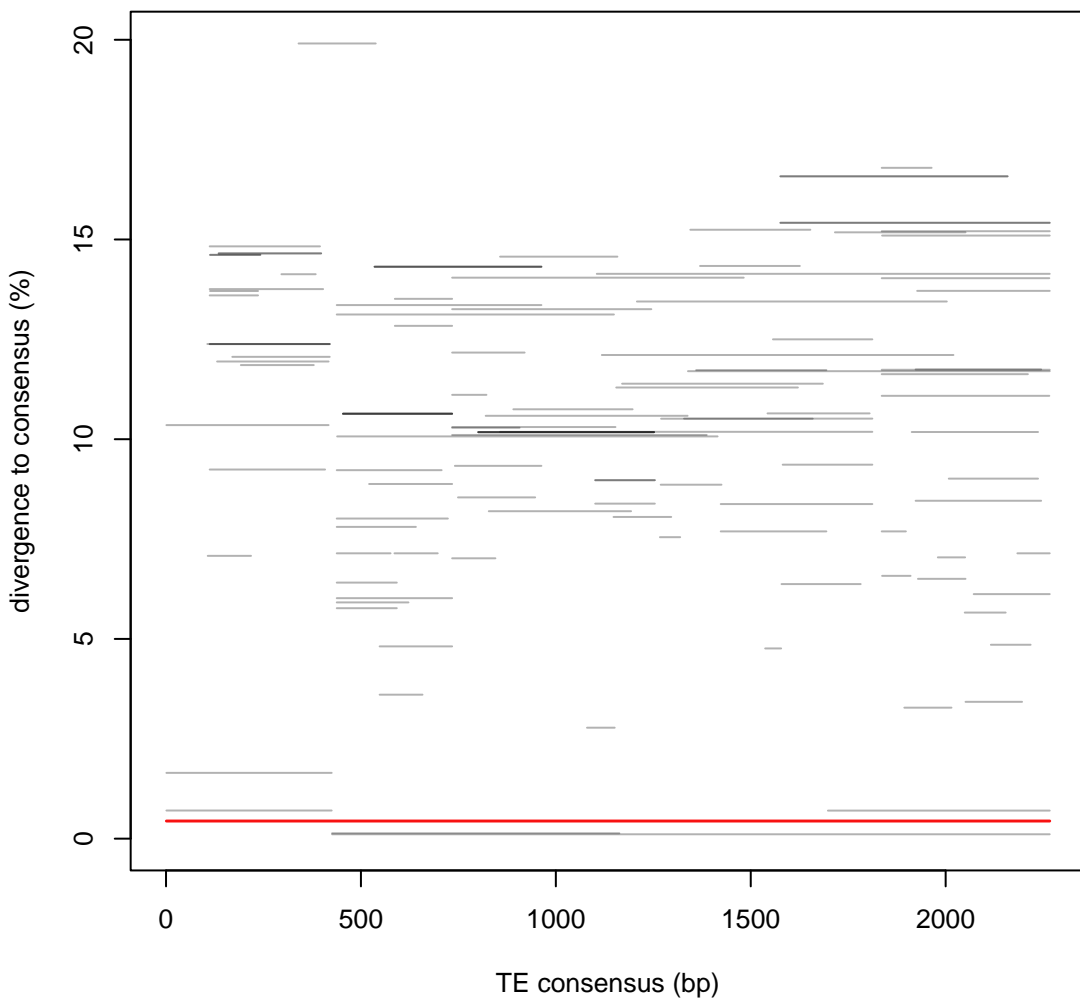
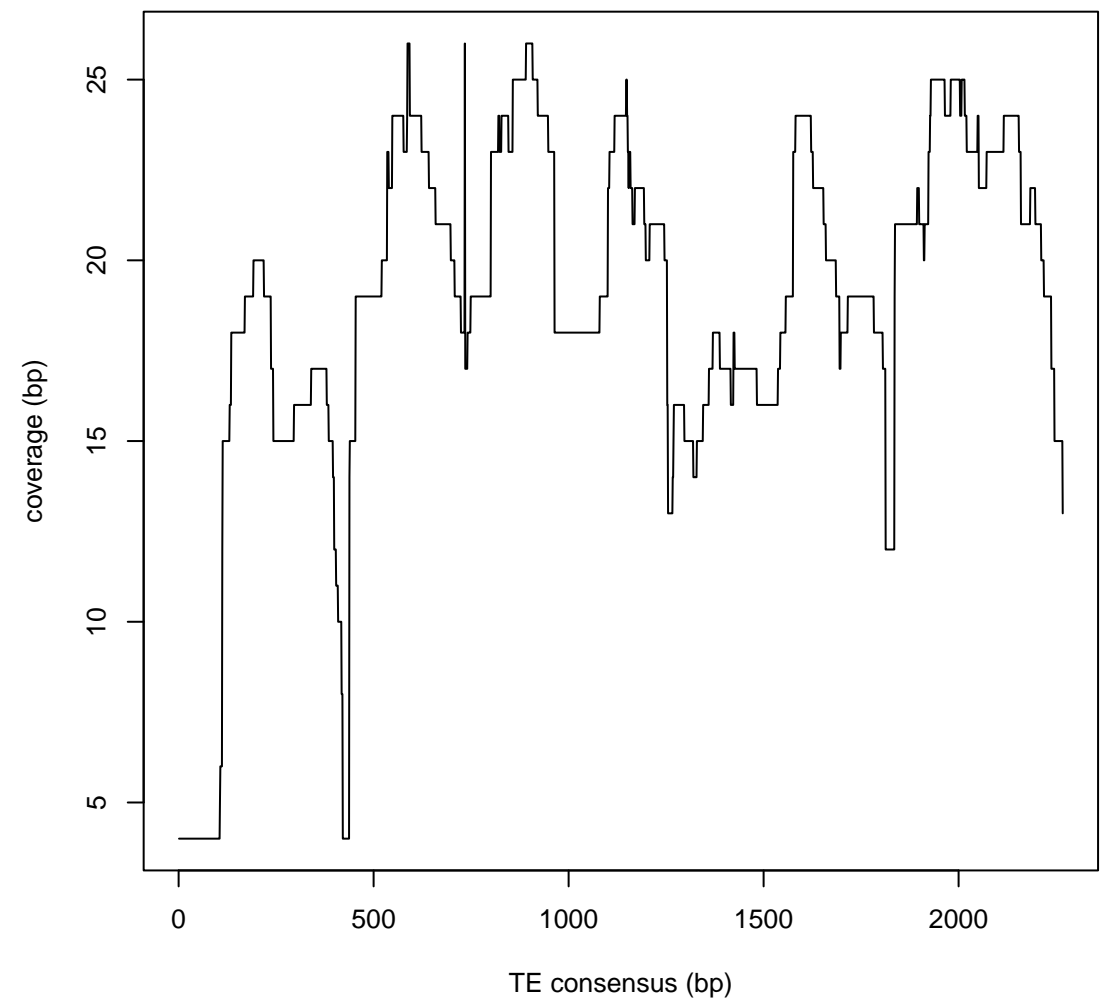


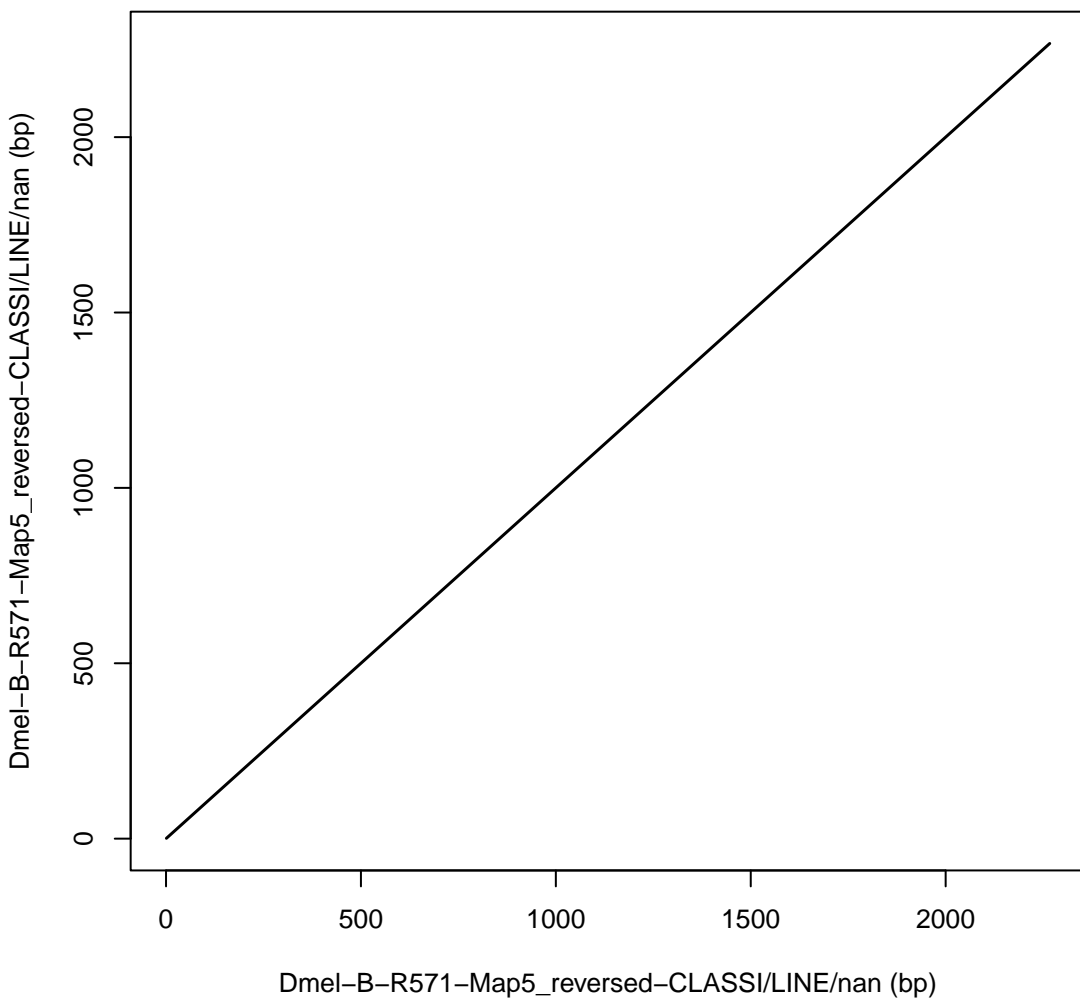
TE: Dmel-B-R571-Map5\_reversed-CLASSI/LINE/nan  
consensus size: 2267bp; fragments: 120; full length: 1 (>=2040.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

