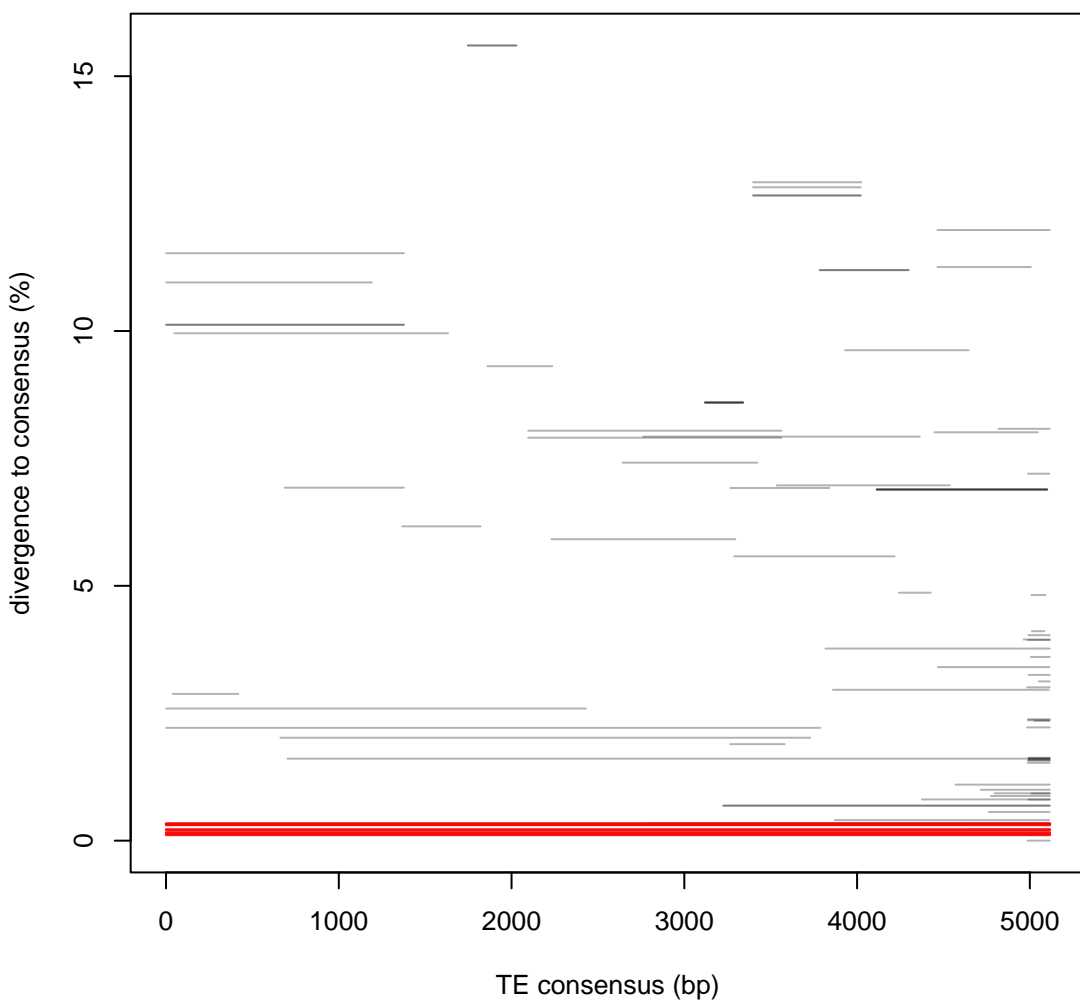
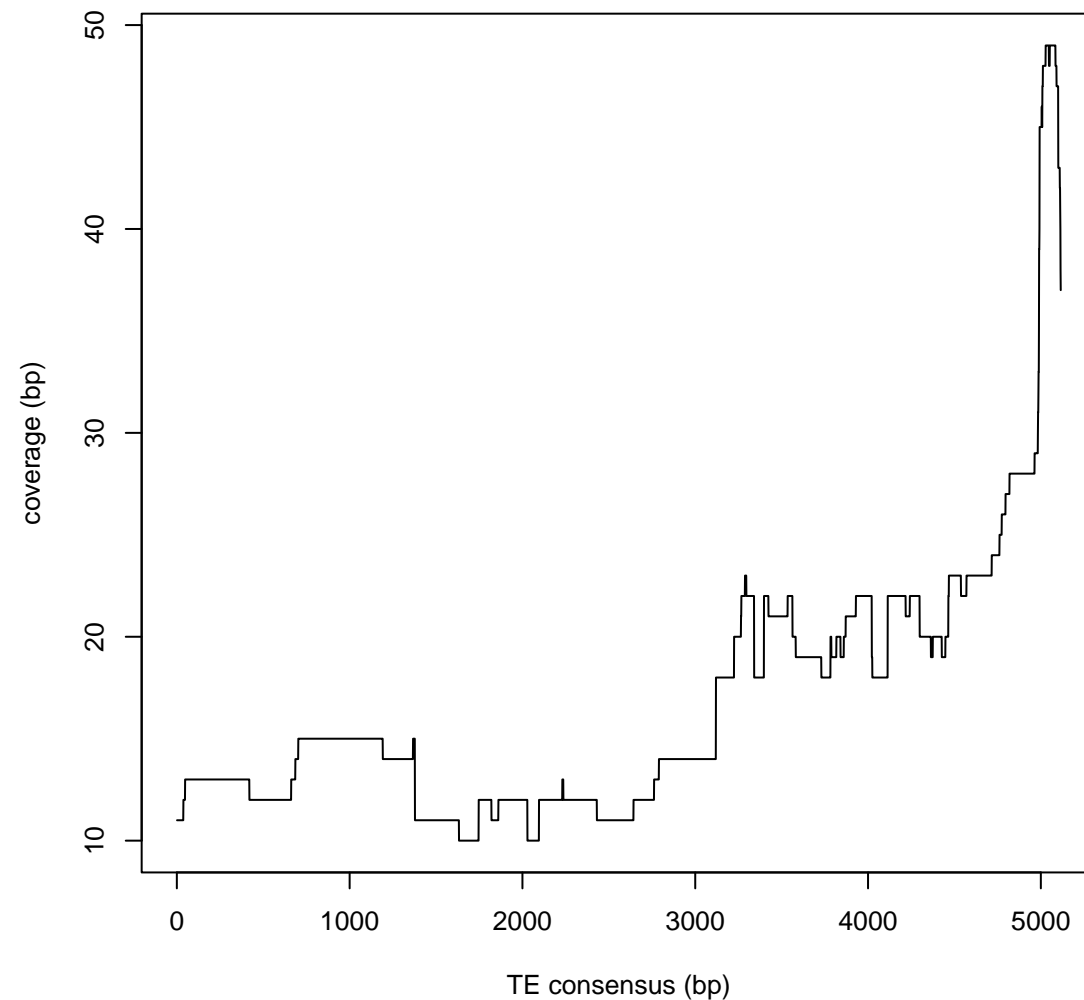


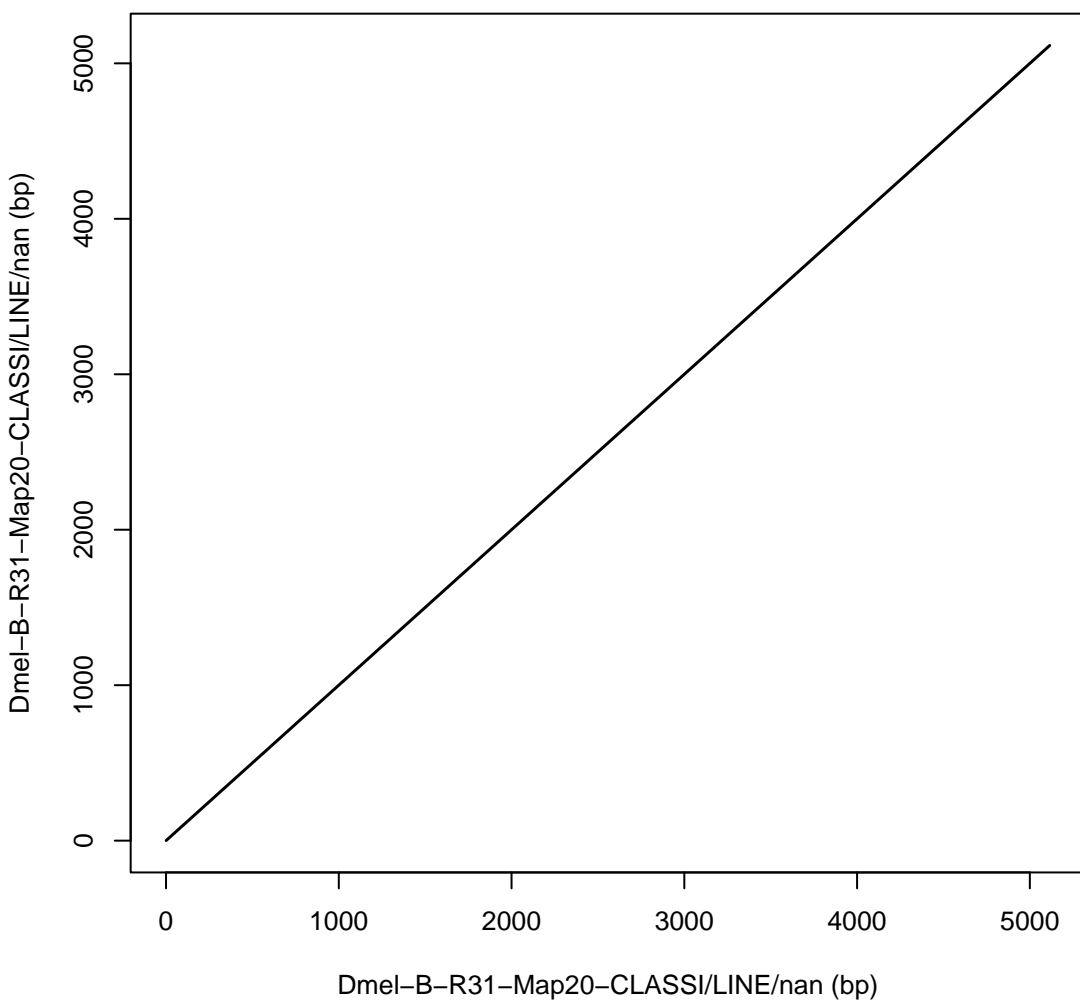
TE: Dmel-B-R31-Map20-CLASSI/LINE/nan  
consensus size: 5115bp; fragments: 86; full length: 5 (>=4603.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

