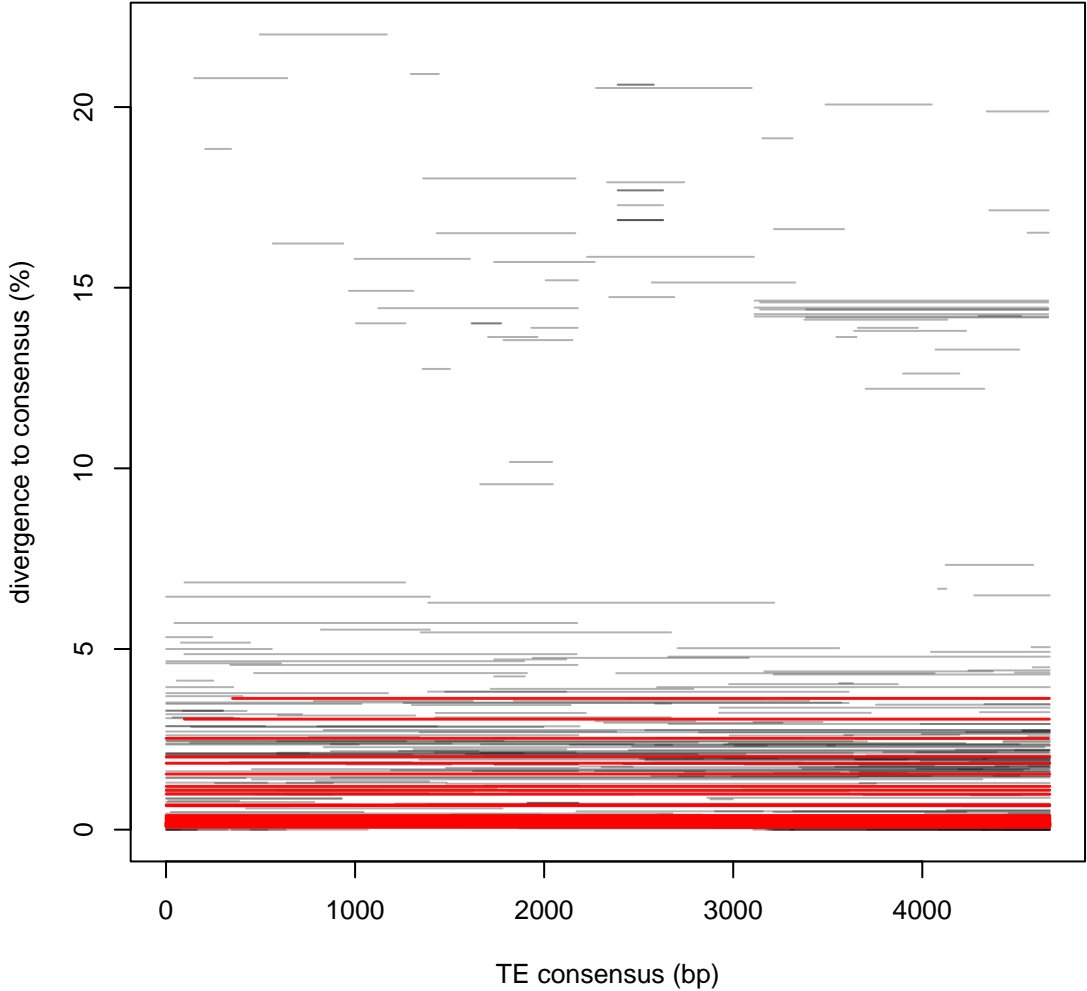
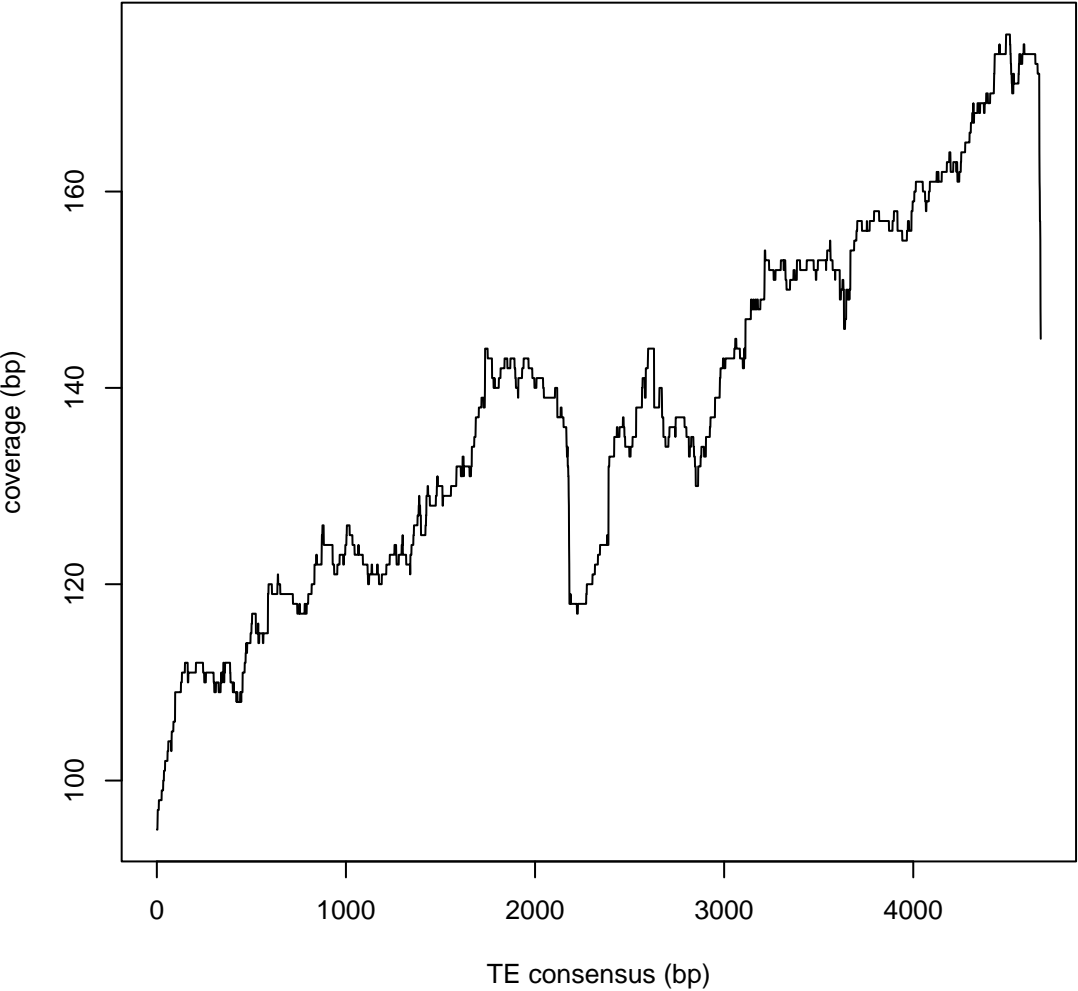


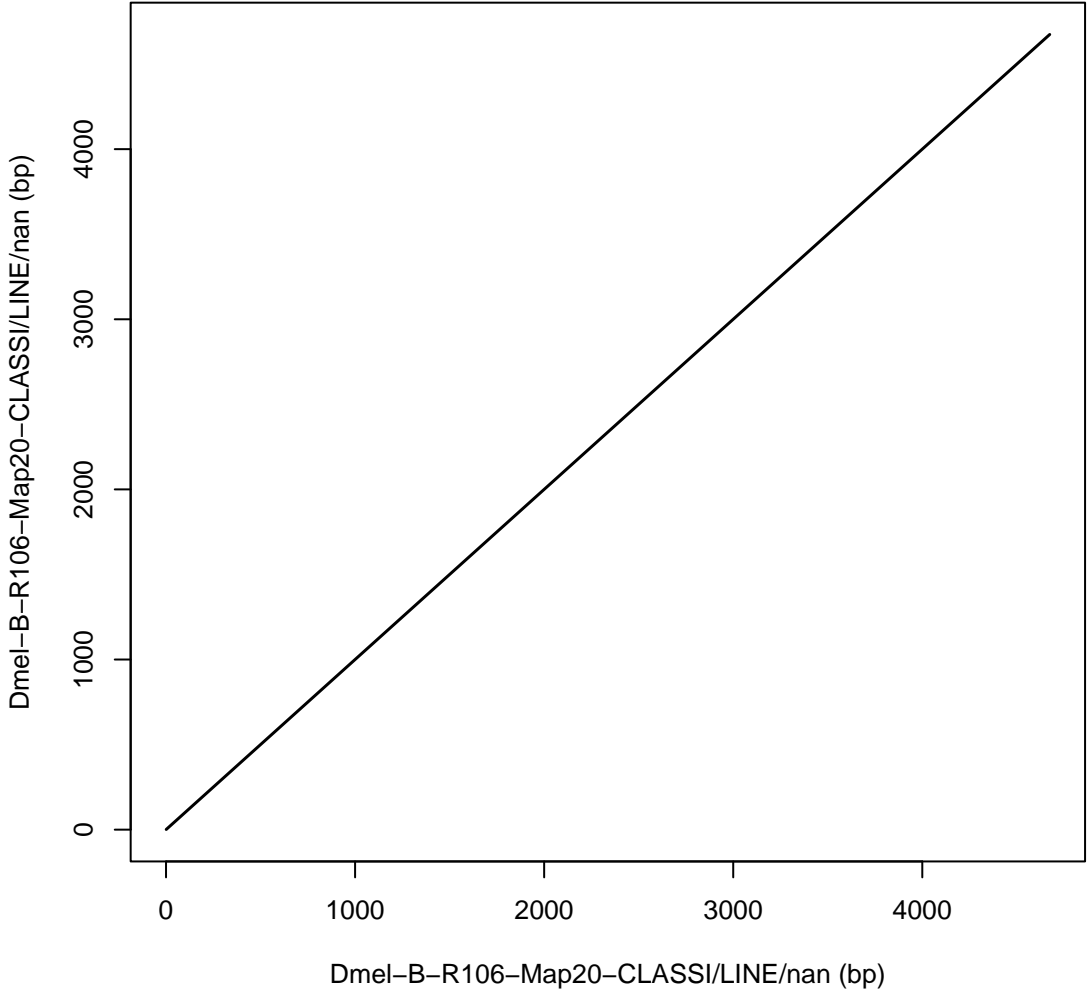
TE: Dmel-B-R106-Map20-CLASSI/LINE/nan  
consensus size: 4674bp; fragments: 392; full length: 51 (>=4206.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

