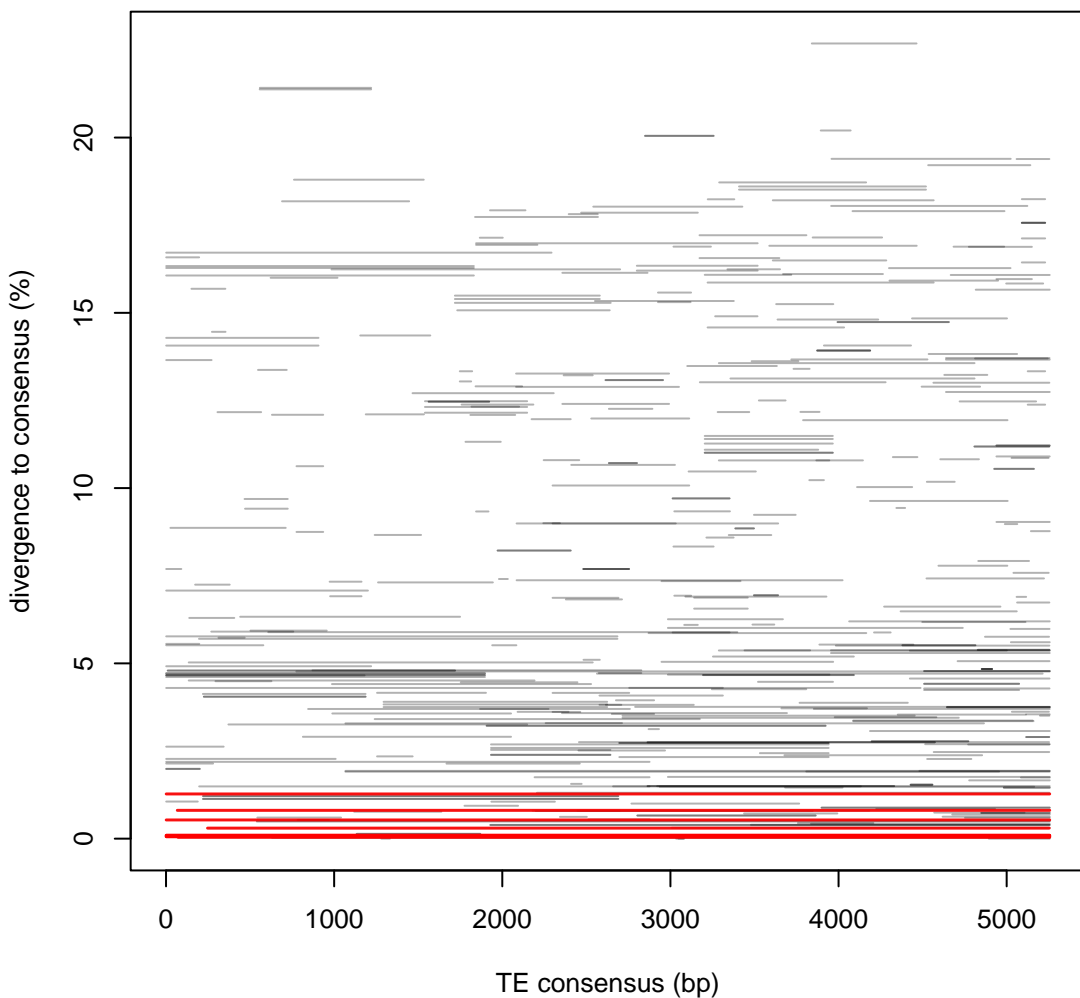
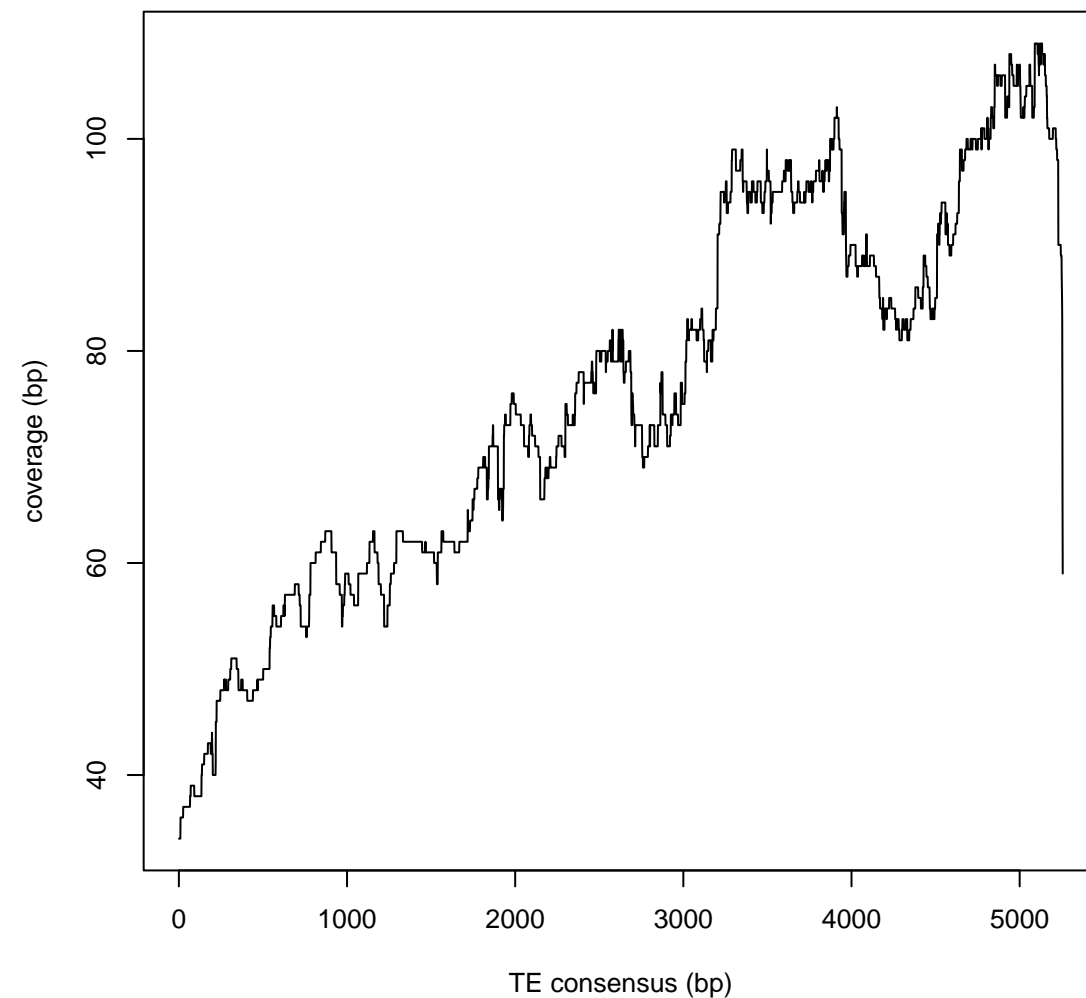


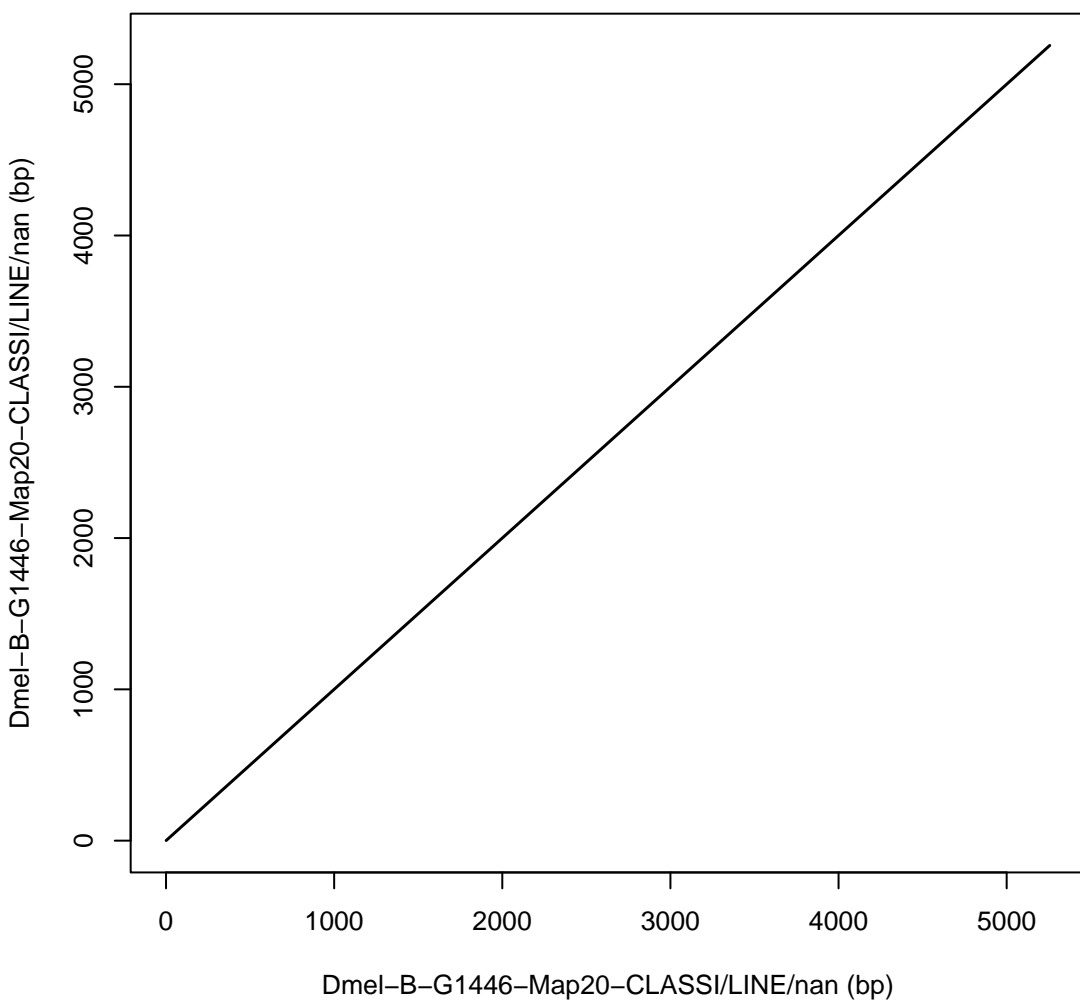
TE: Dmel-B-G1446-Map20-CLASSI/LINE/nan  
consensus size: 5256bp; fragments: 443; full length: 9 (>=4730.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

