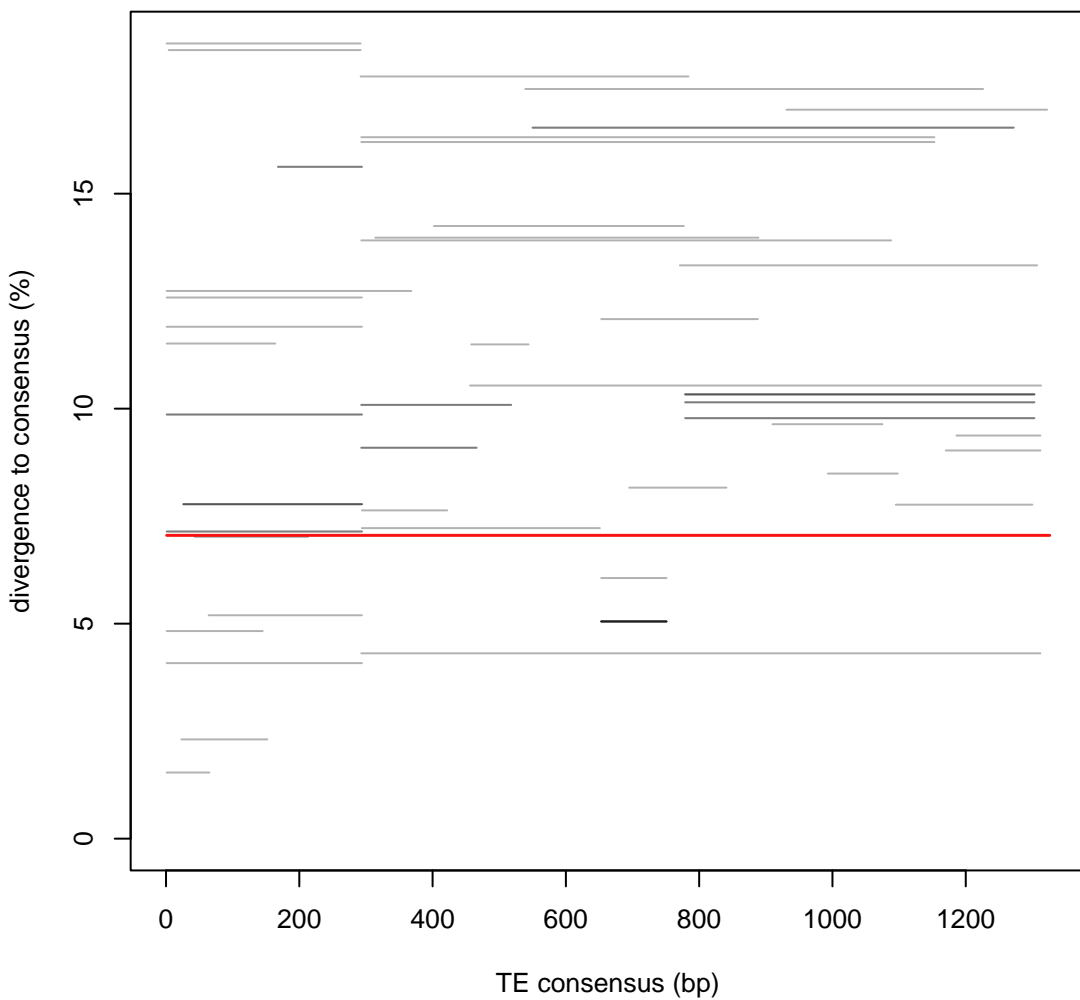
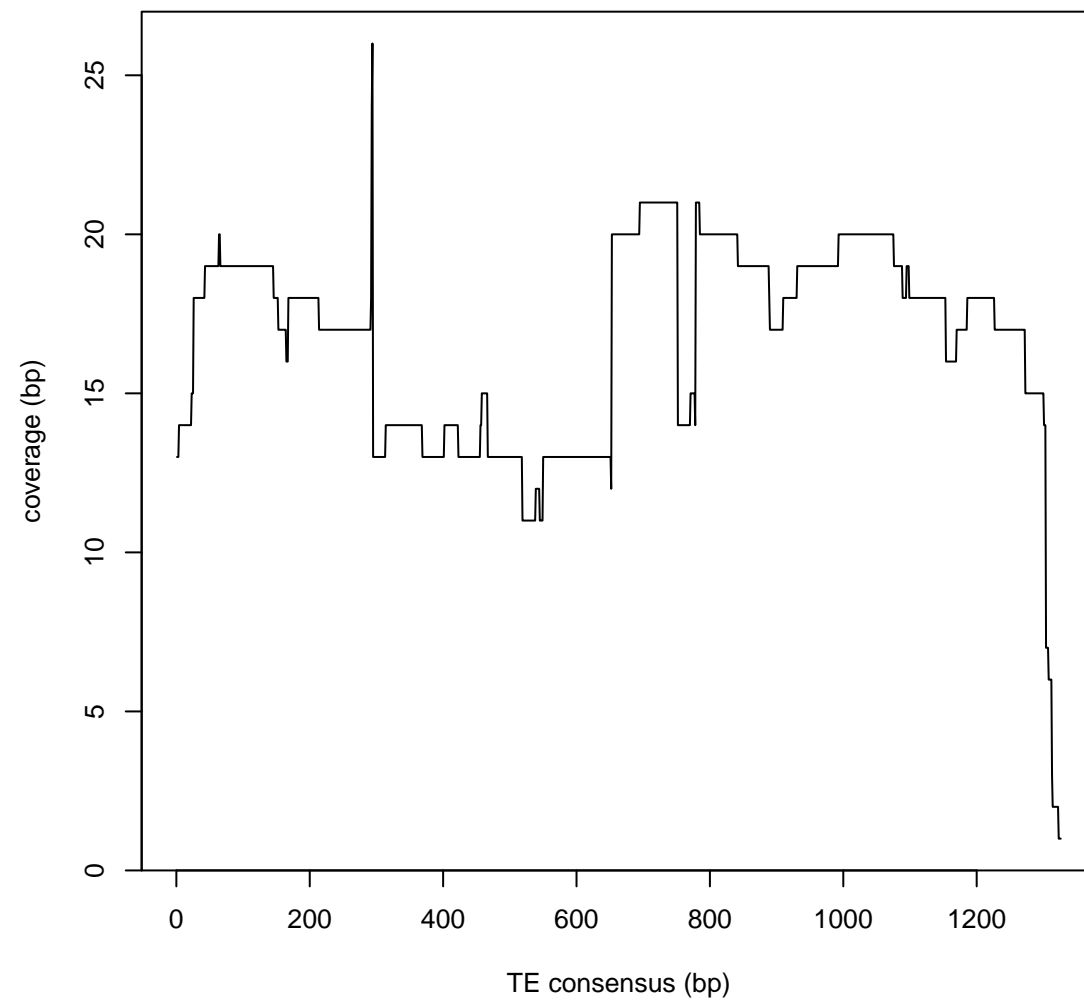


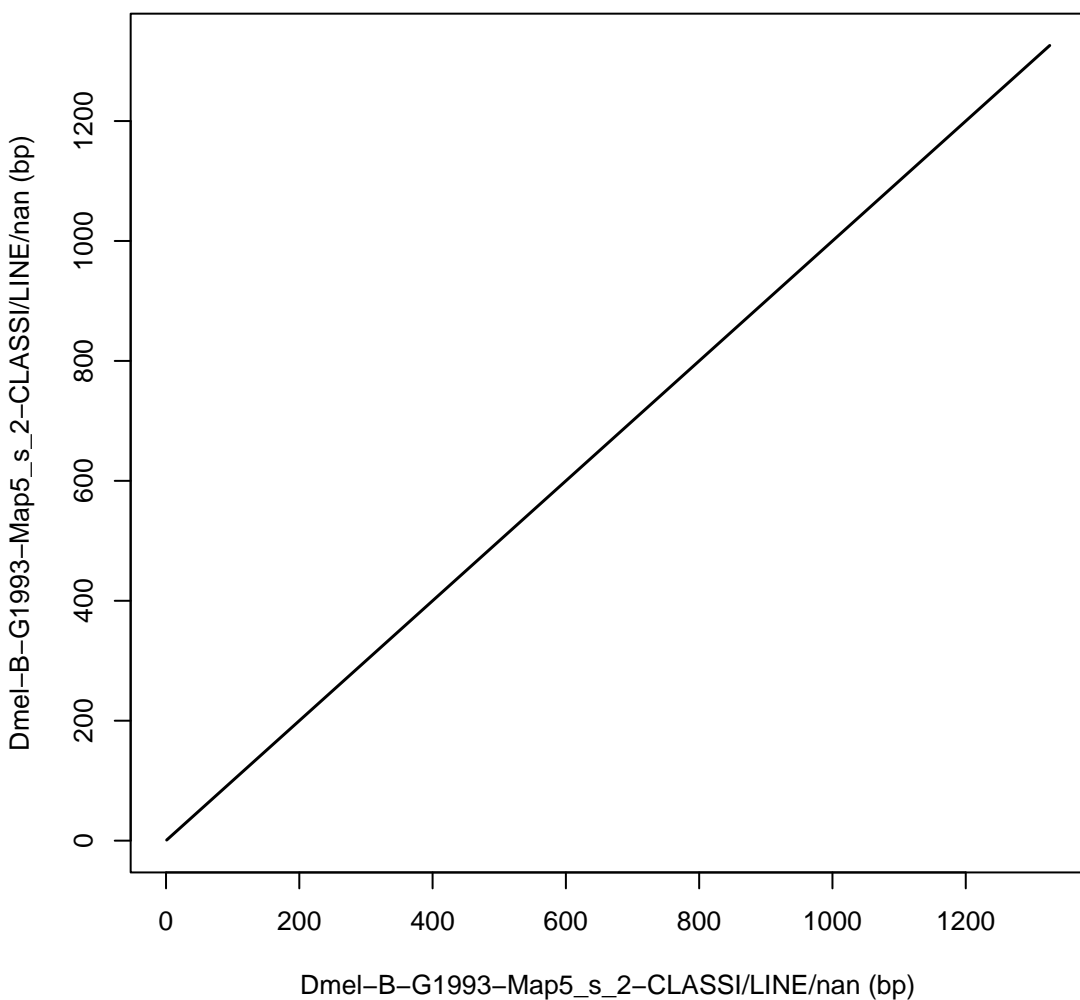
TE: Dmel-B-G1993-Map5\_s\_2-CLASSI/LINE/nan  
consensus size: 1326bp; fragments: 63; full length: 1 (>=1193.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

