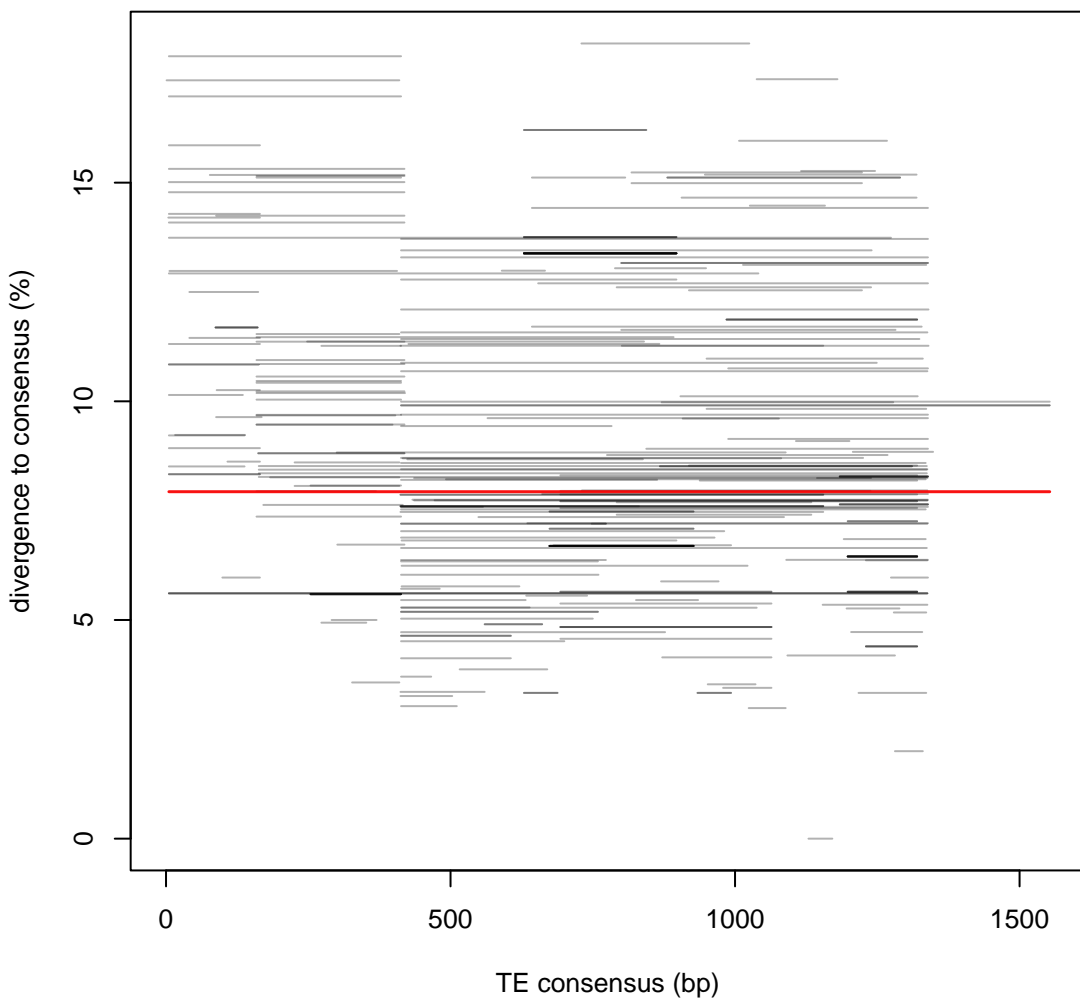
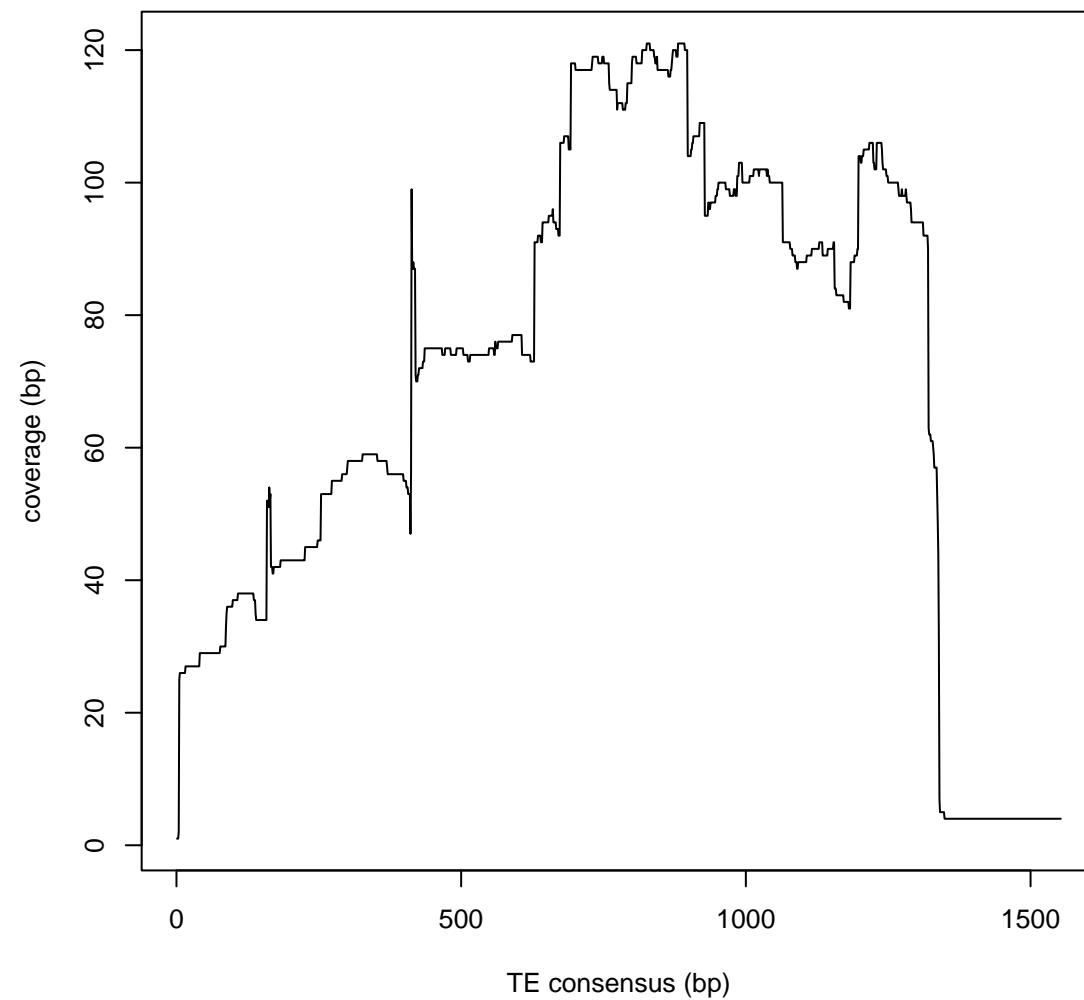


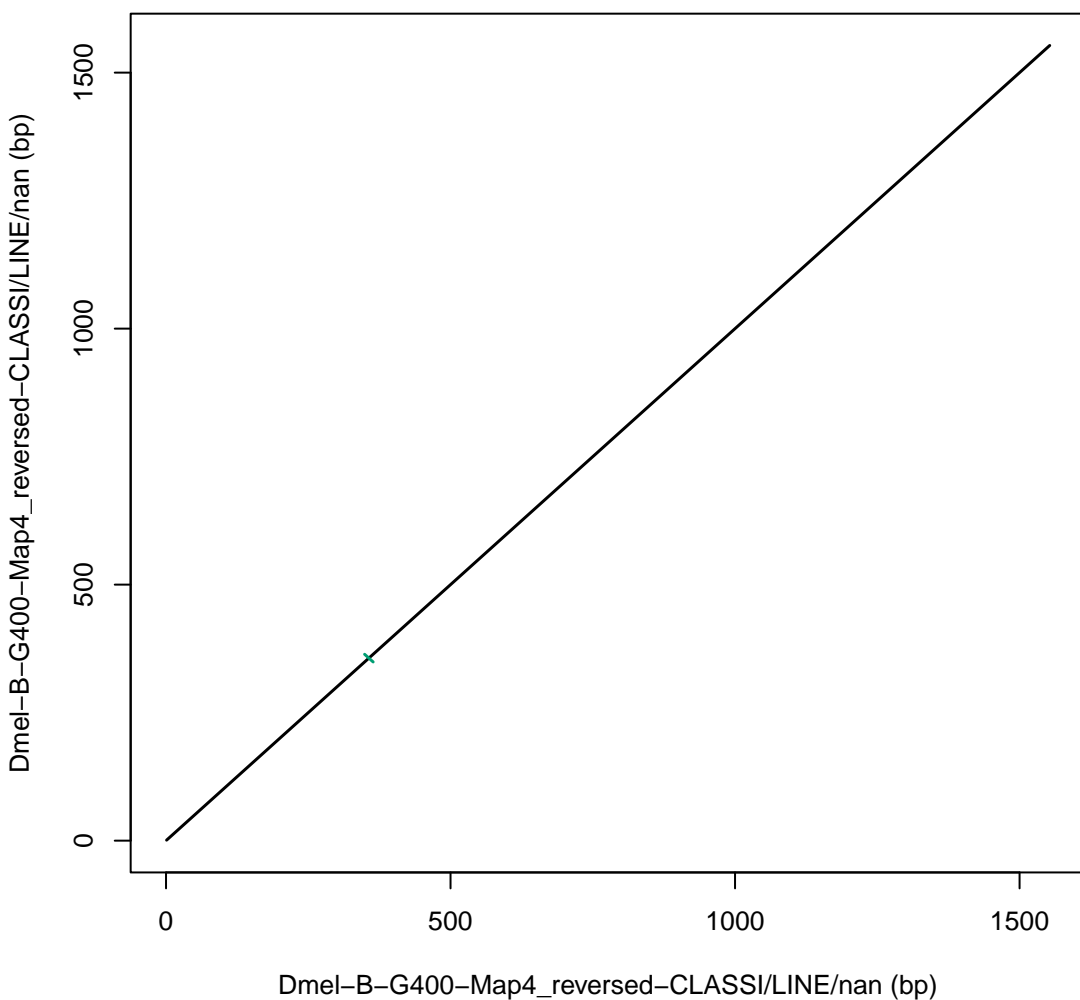
TE: Dmel-B-G400-Map4\_reversed-CLASSI/LINE/nan  
consensus size: 1553bp; fragments: 290; full length: 1 (>=1397.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

