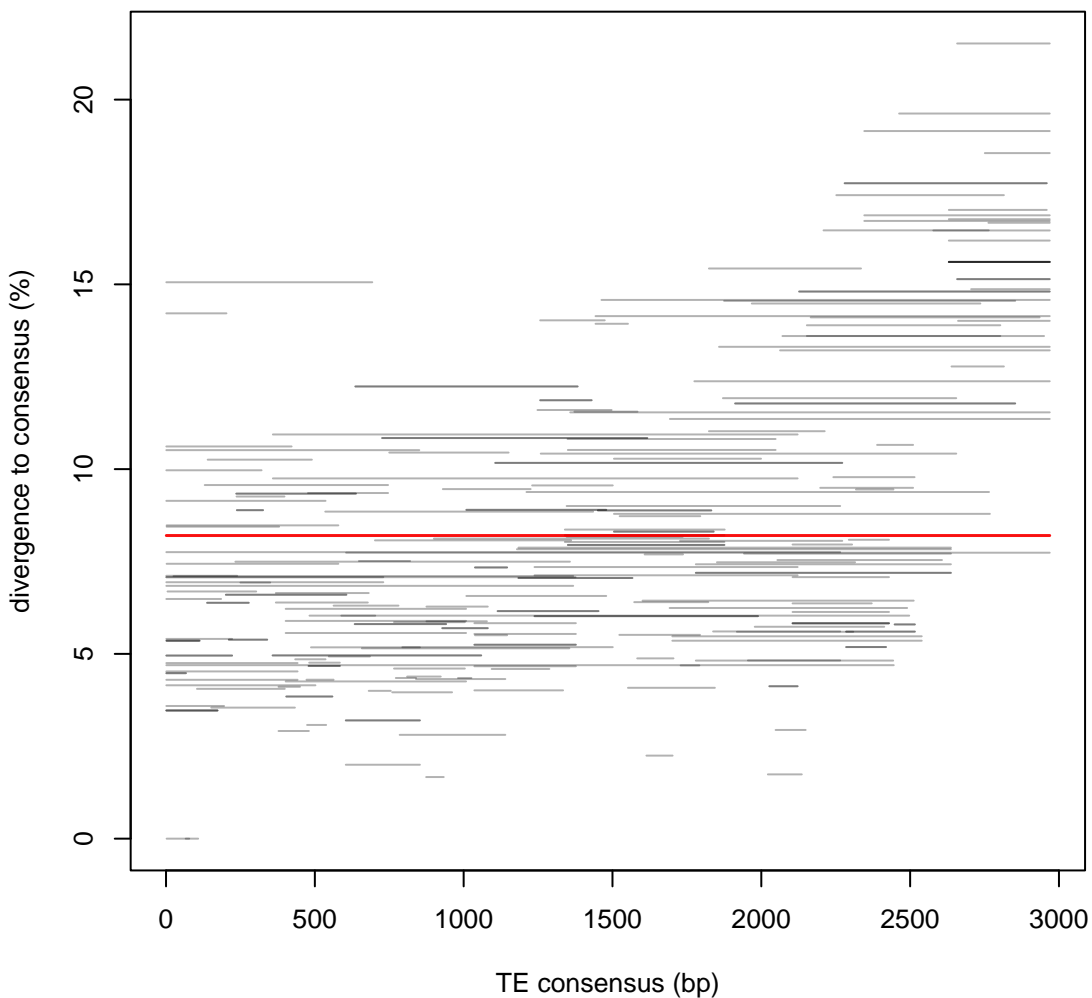
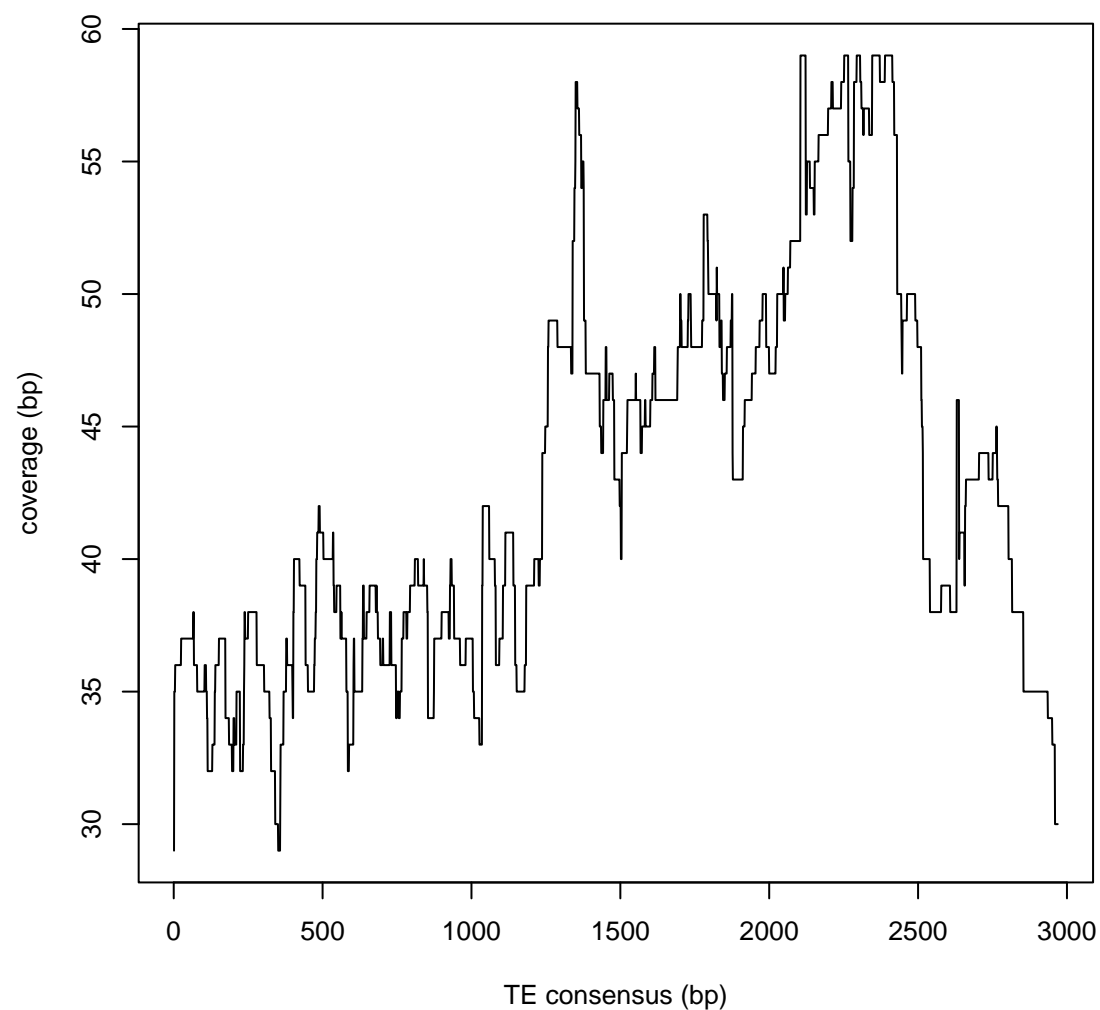


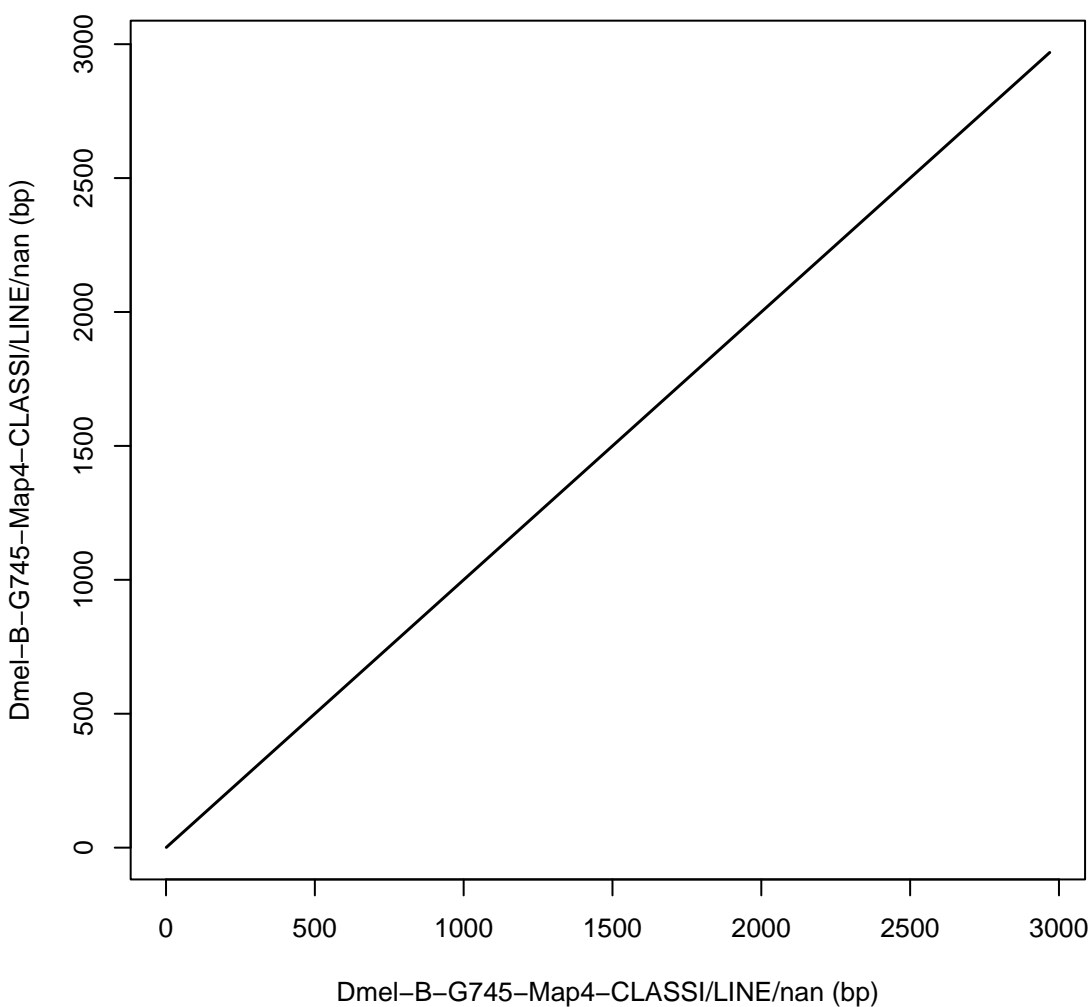
TE: Dmel-B-G745-Map4-CLASSI/LINE/nan  
consensus size: 2969bp; fragments: 255; full length: 1 (>=2672.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

