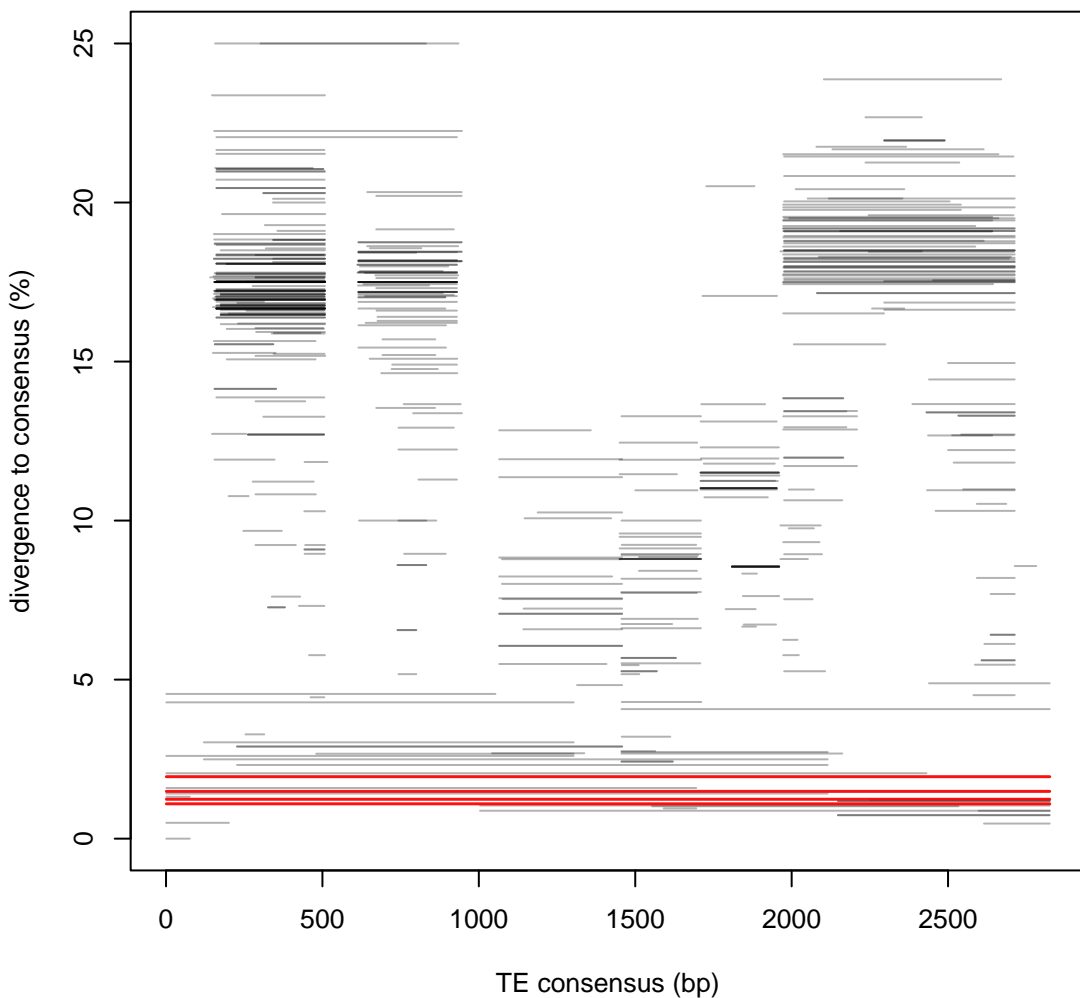
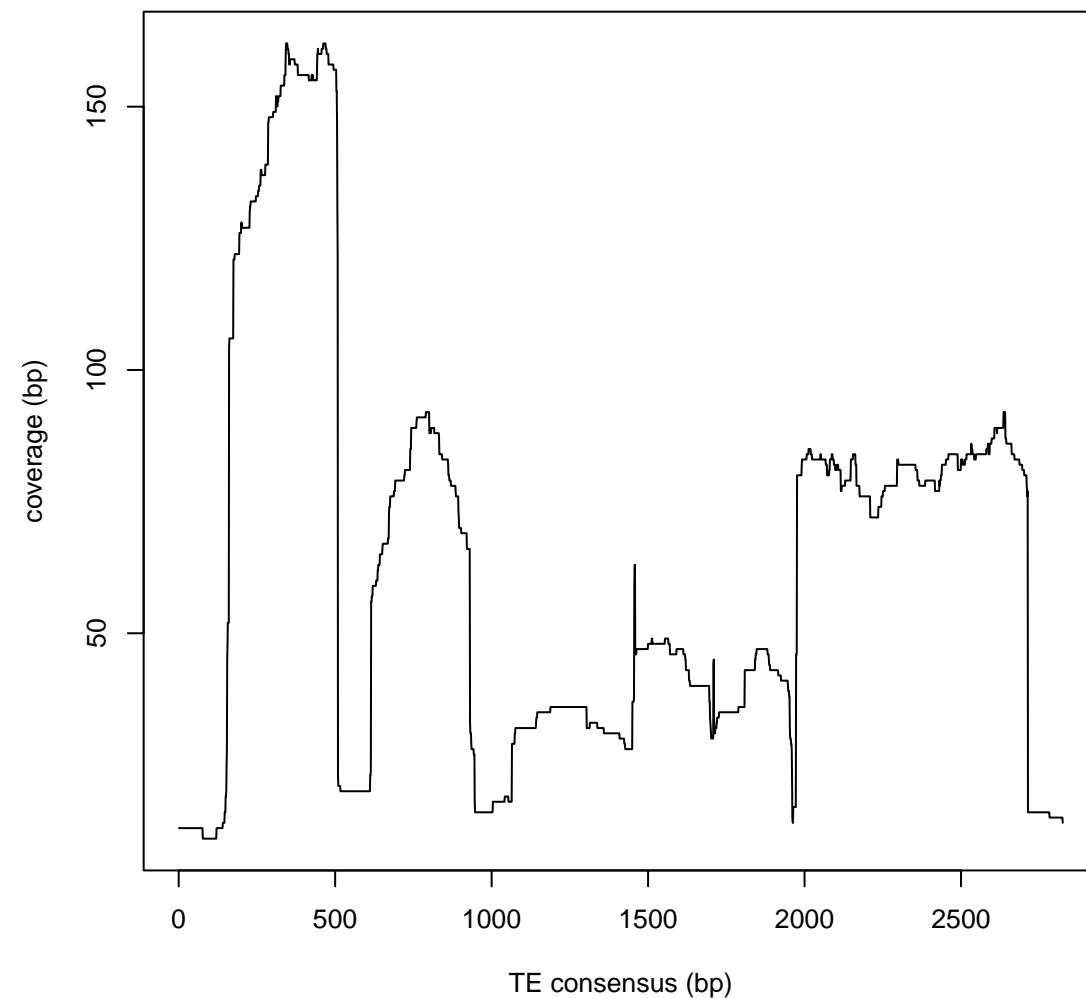


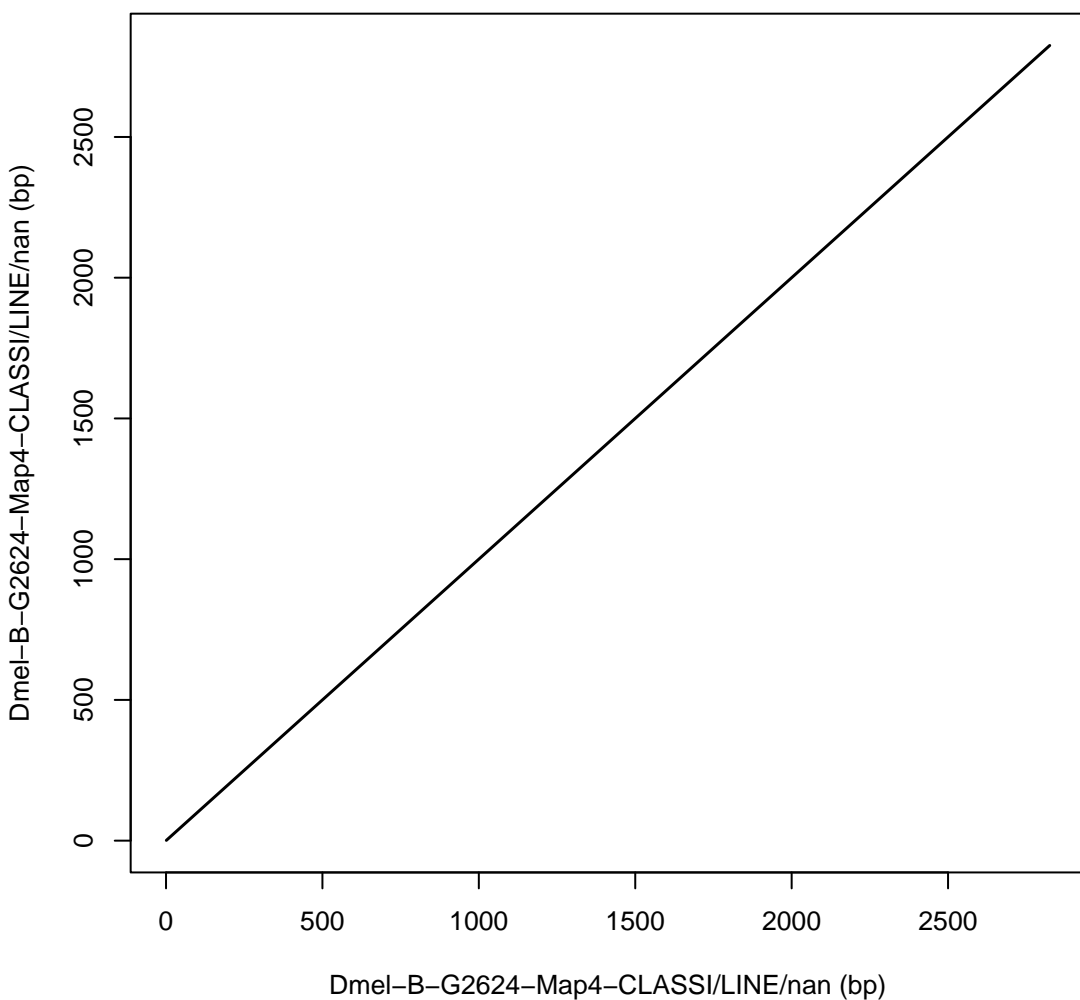
TE: Dmel-B-G2624-Map4-CLASSI/LINE/nan  
consensus size: 2825bp; fragments: 477; full length: 4 ( $\geq 2542.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

