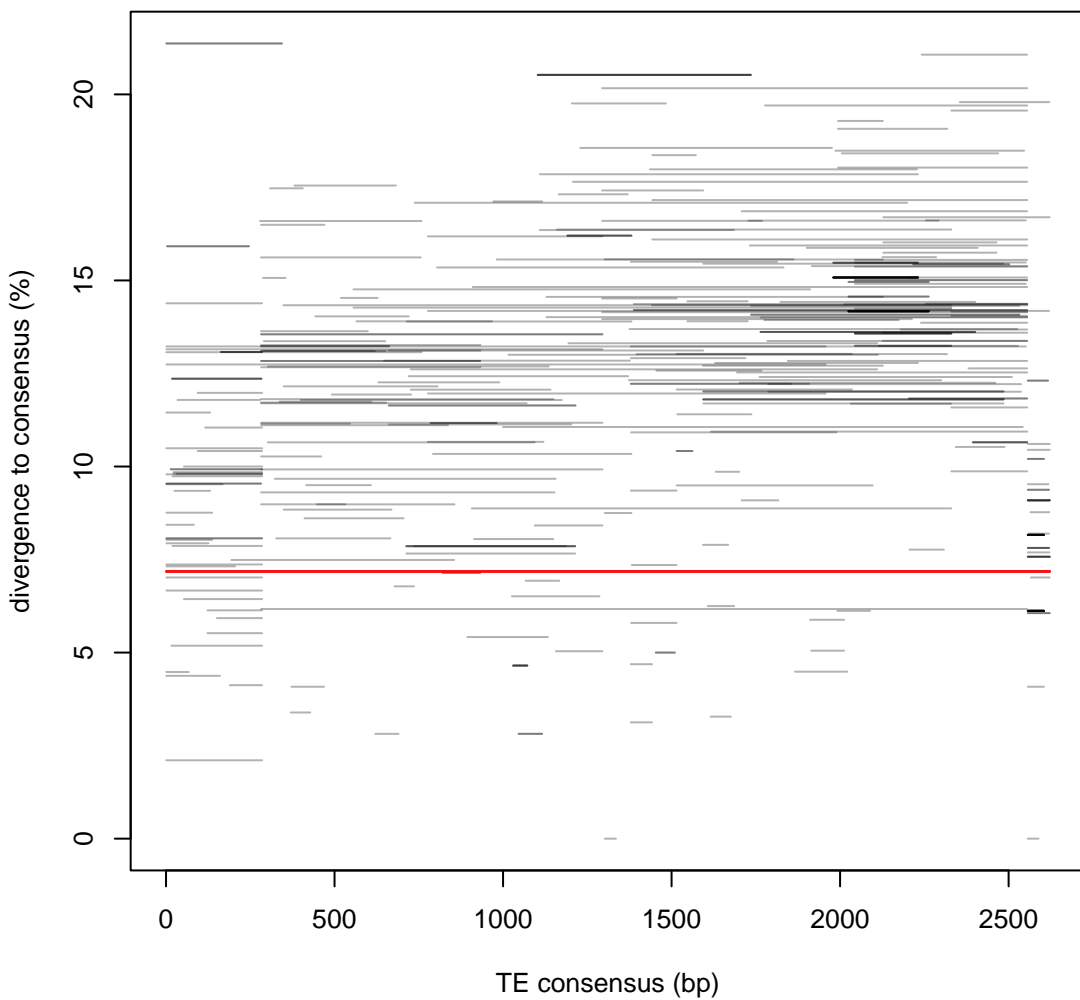
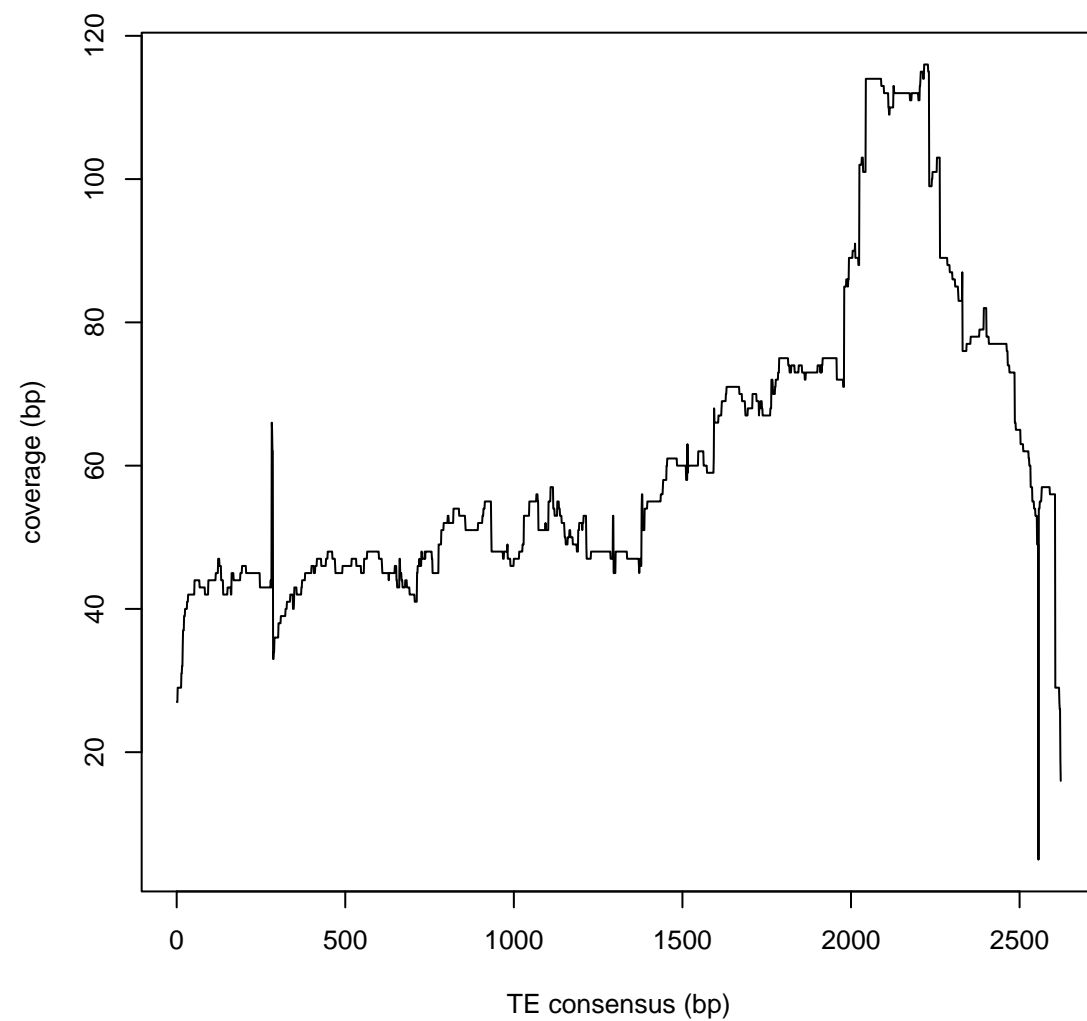


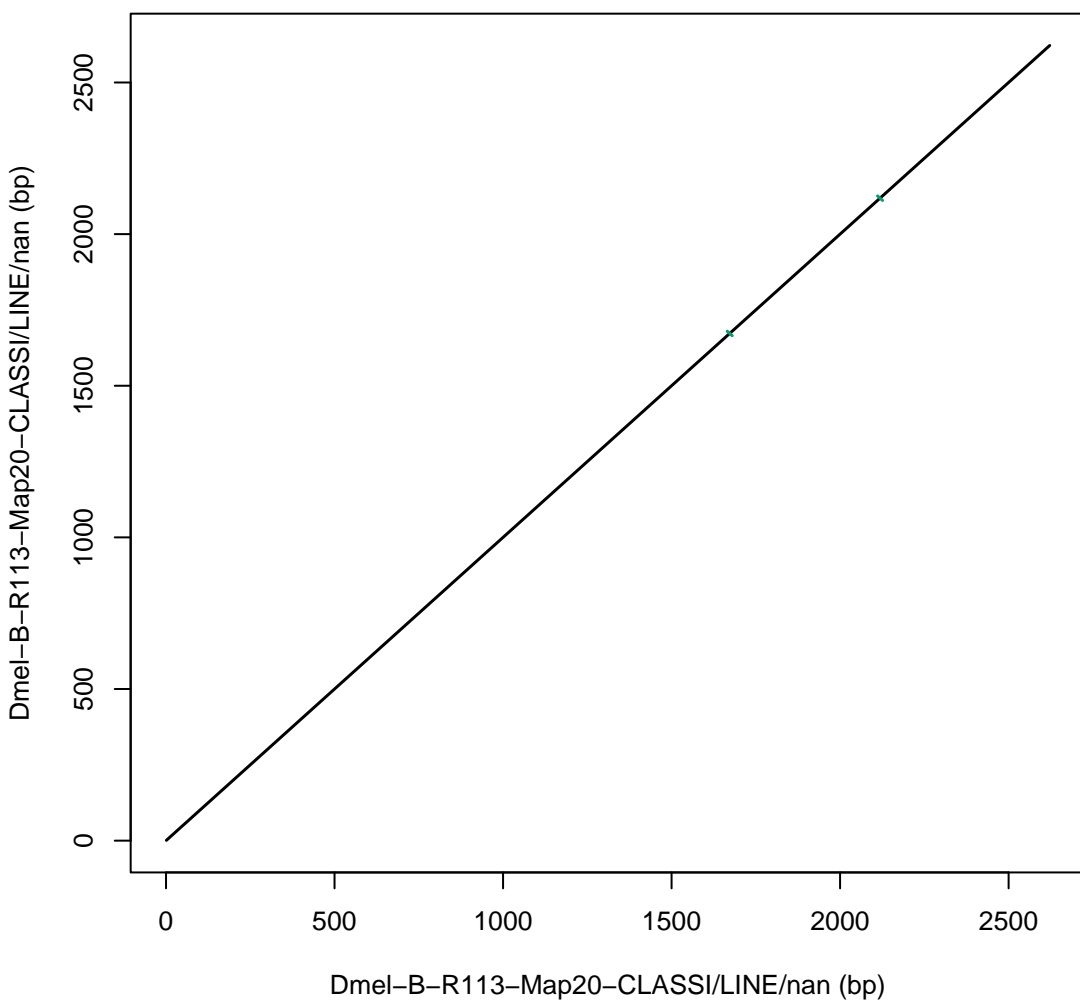
TE: Dmel-B-R113-Map20-CLASSI/LINE/nan  
consensus size: 2622bp; fragments: 381; full length: 1 ( $\geq 2359.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

