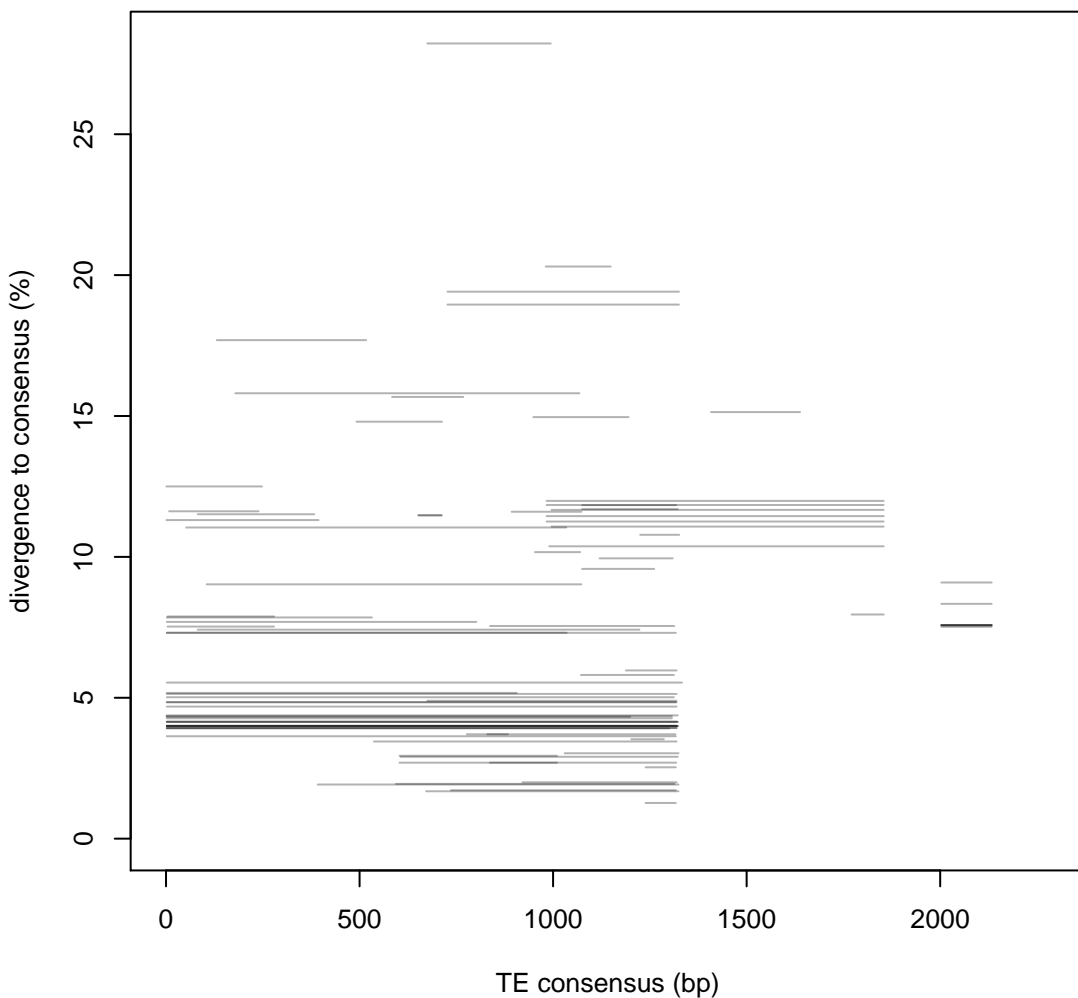
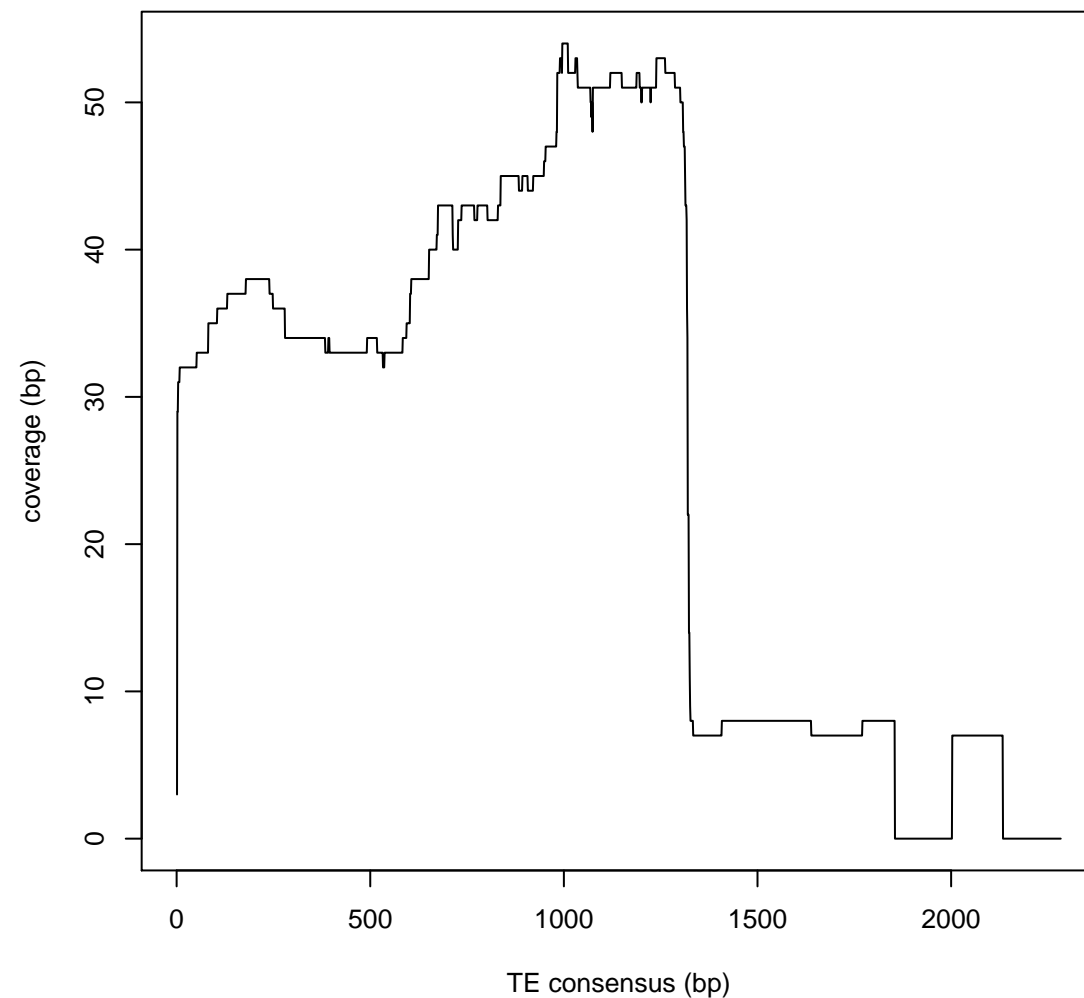


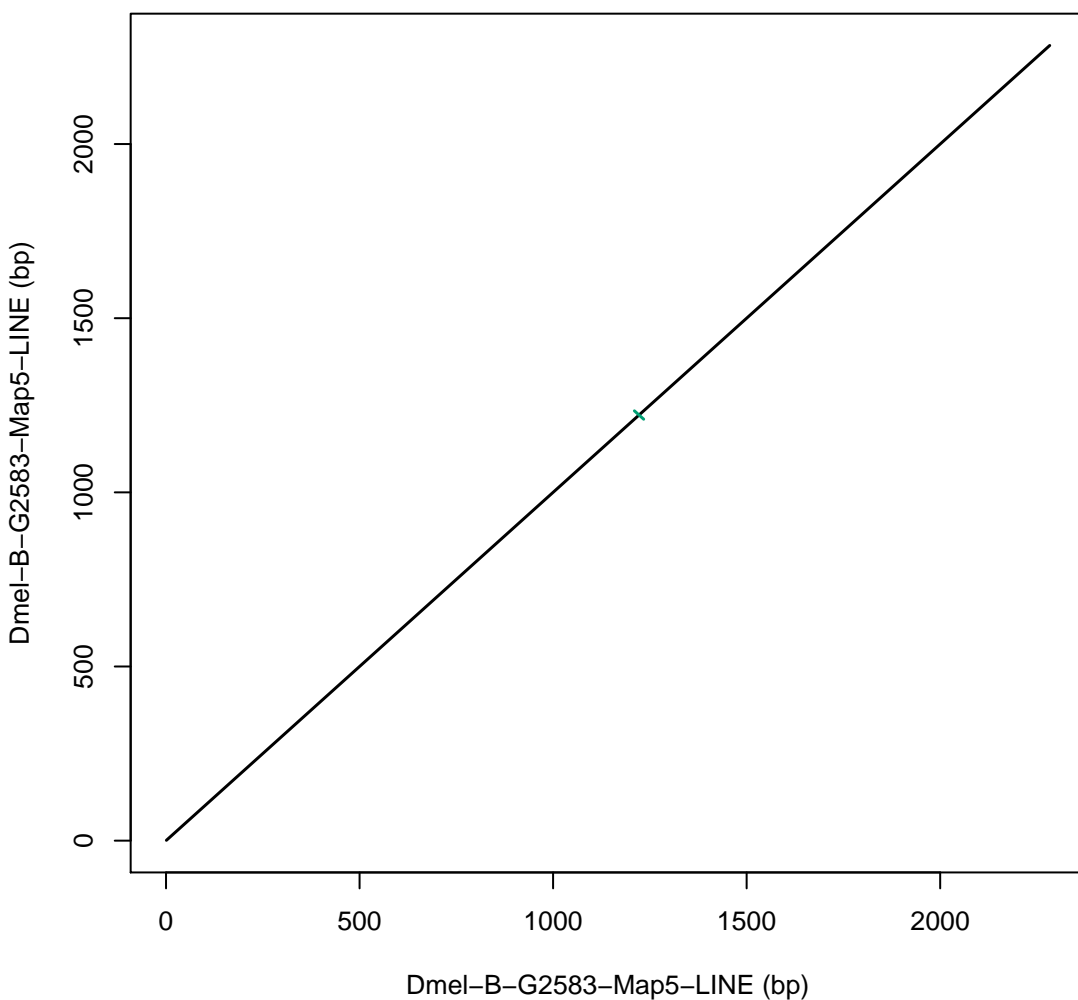
TE: Dmel-B-G2583-Map5-LINE
consensus size: 2283bp; fragments: 90; full length: 0 (>=2054.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

