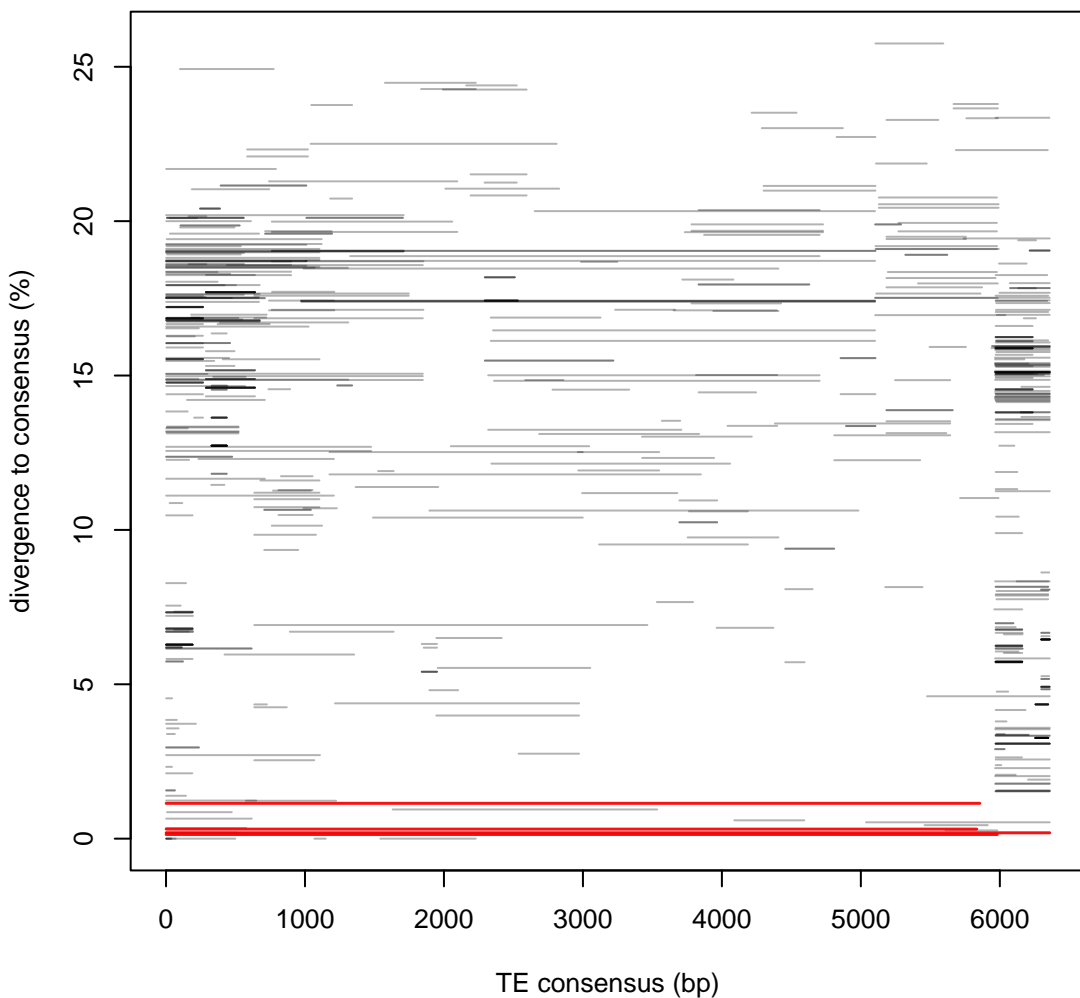
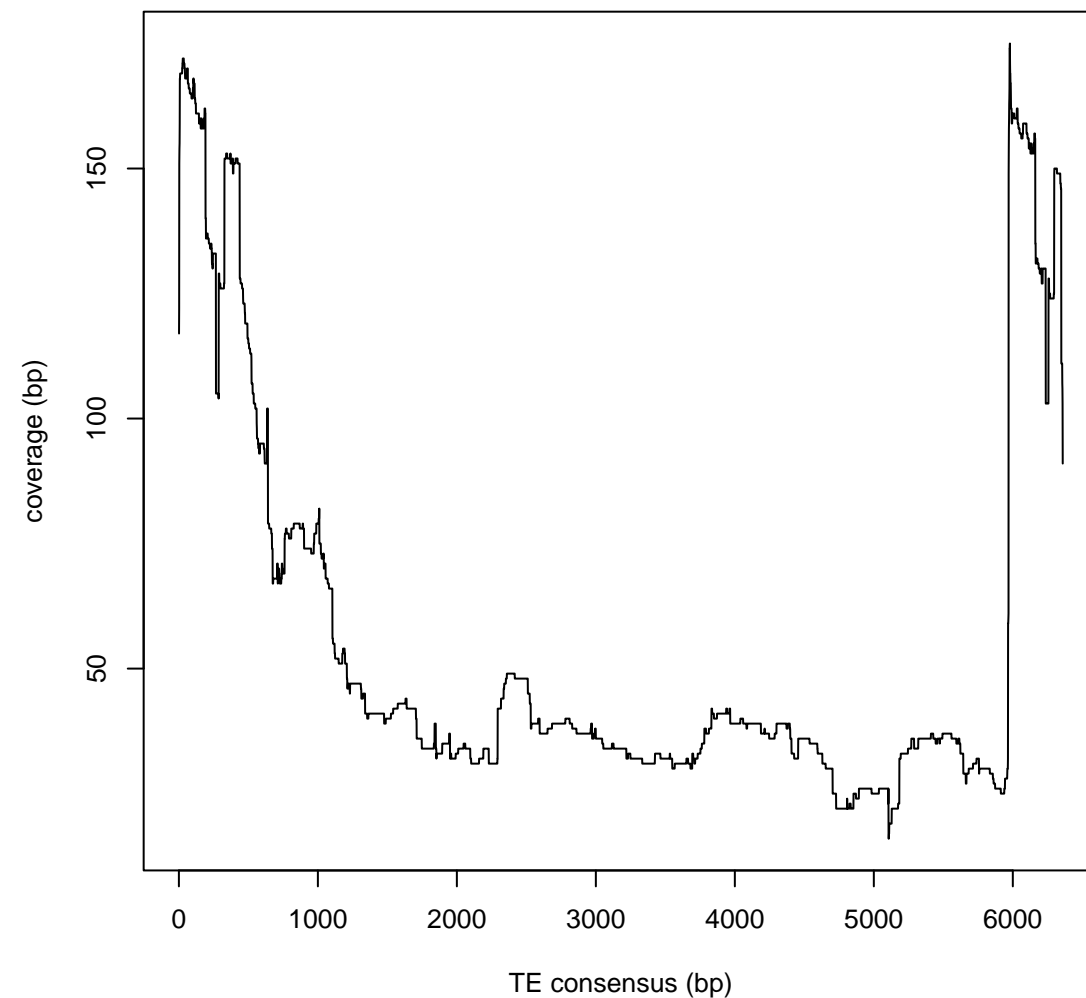


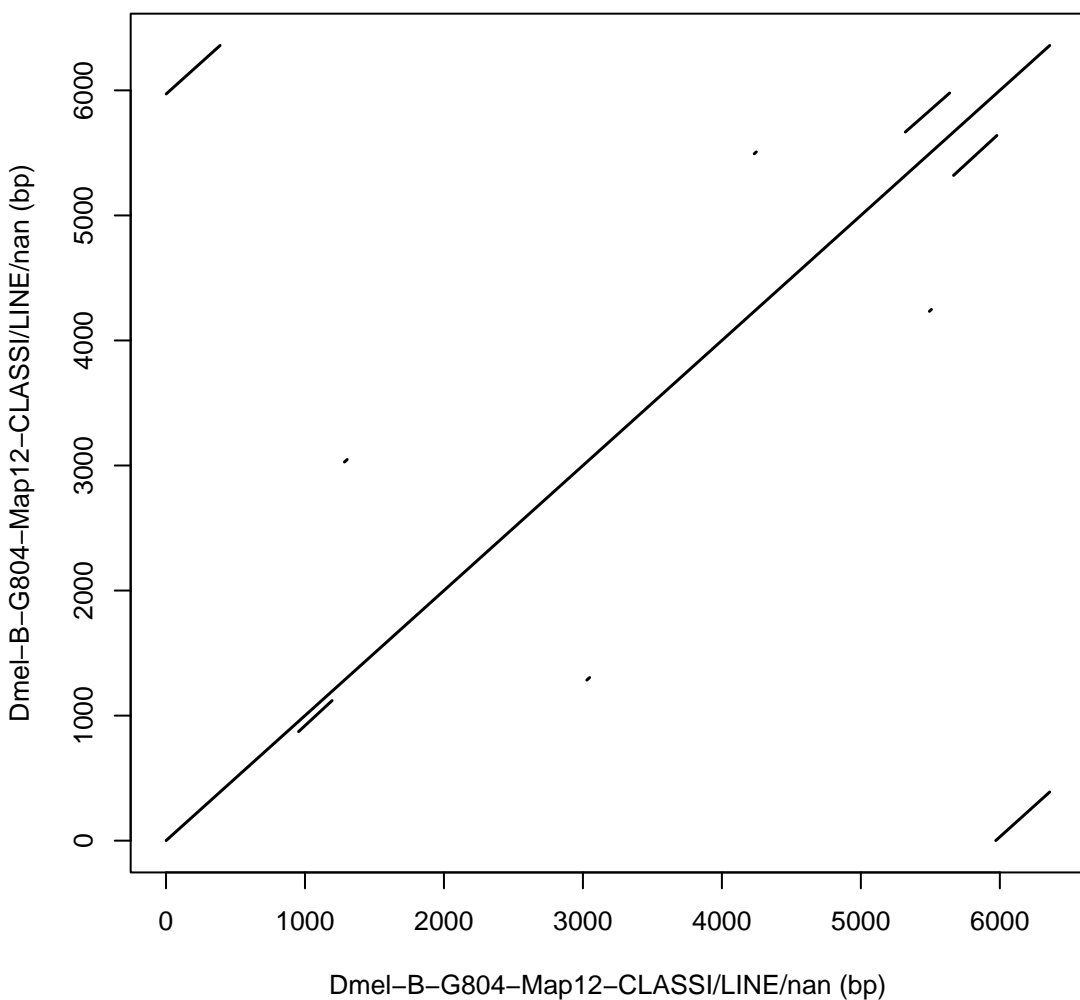
TE: Dmel-B-G804-Map12-CLASSI/LINE/nan
consensus size: 6359bp; fragments: 664; full length: 4 (≥ 5723.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

