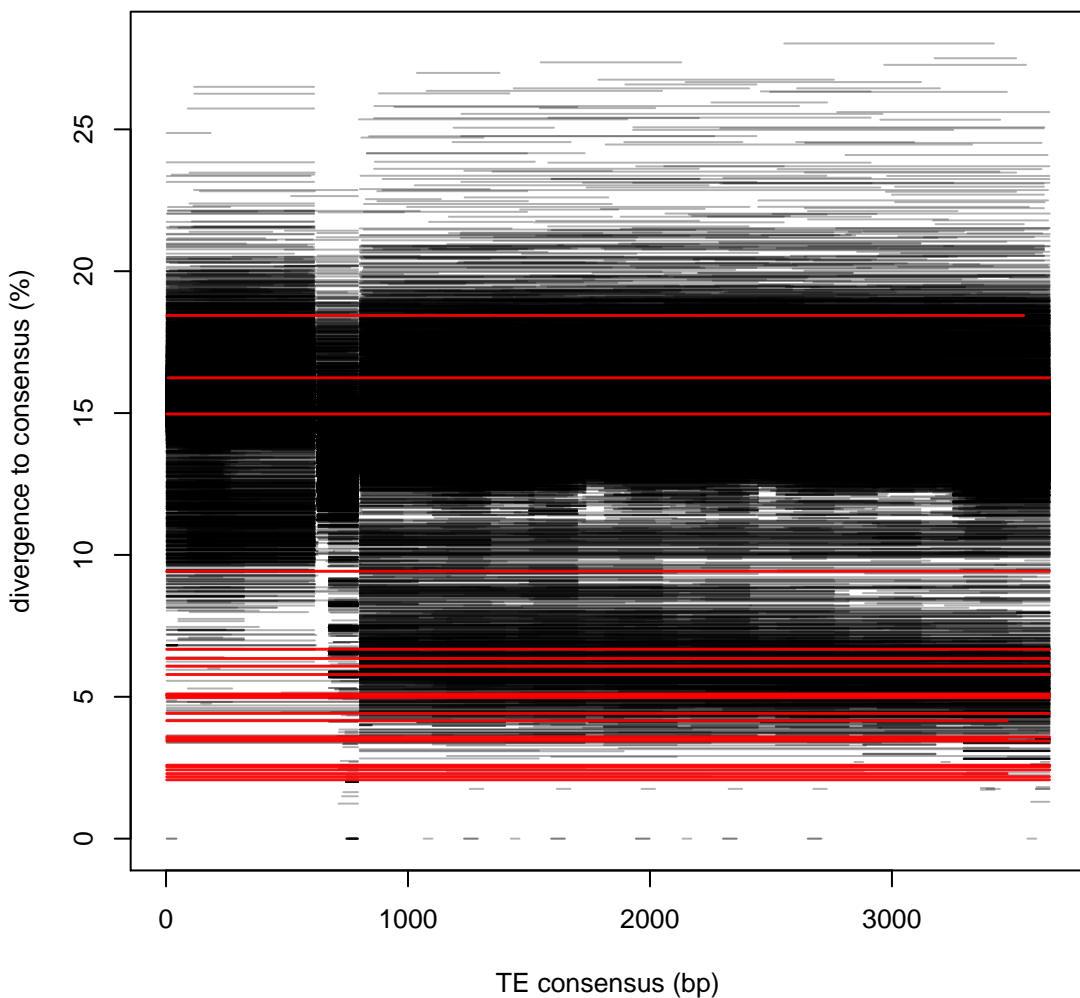
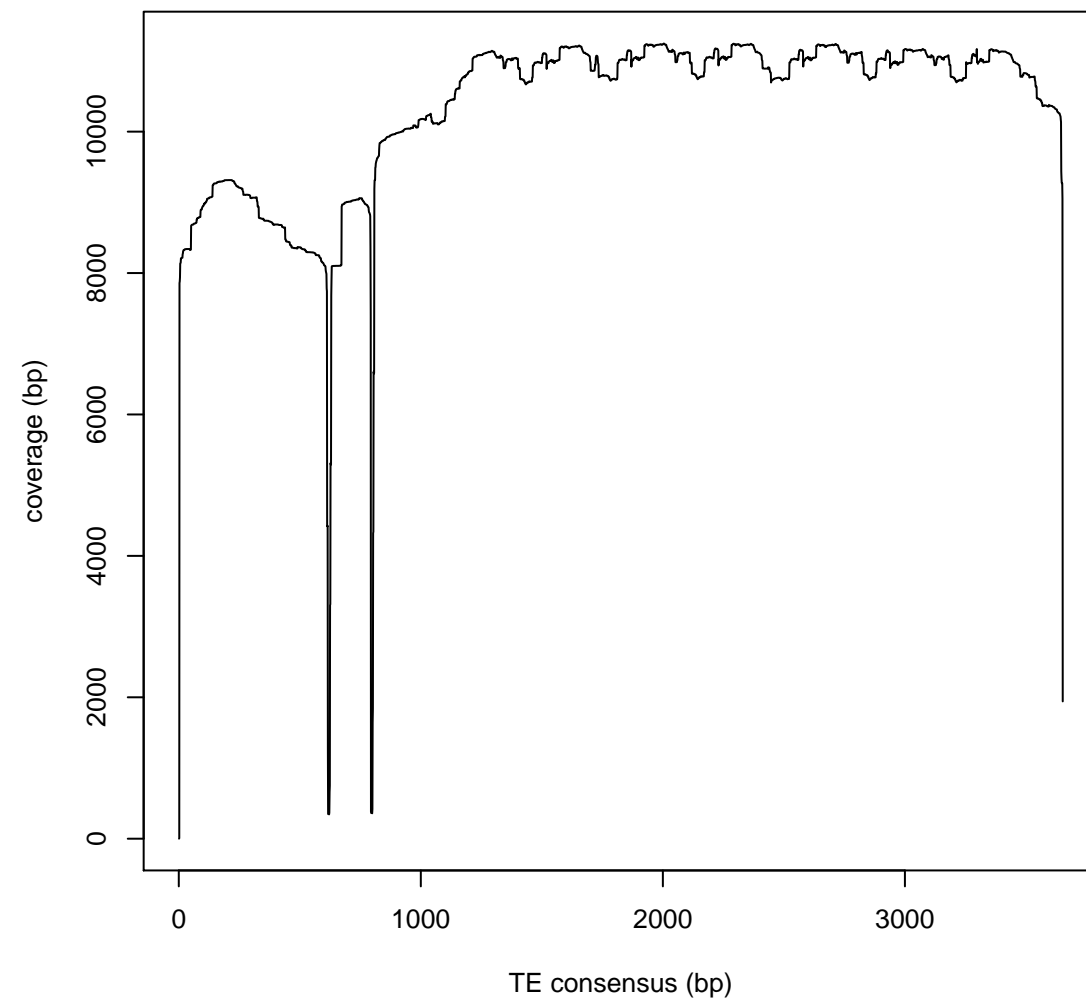


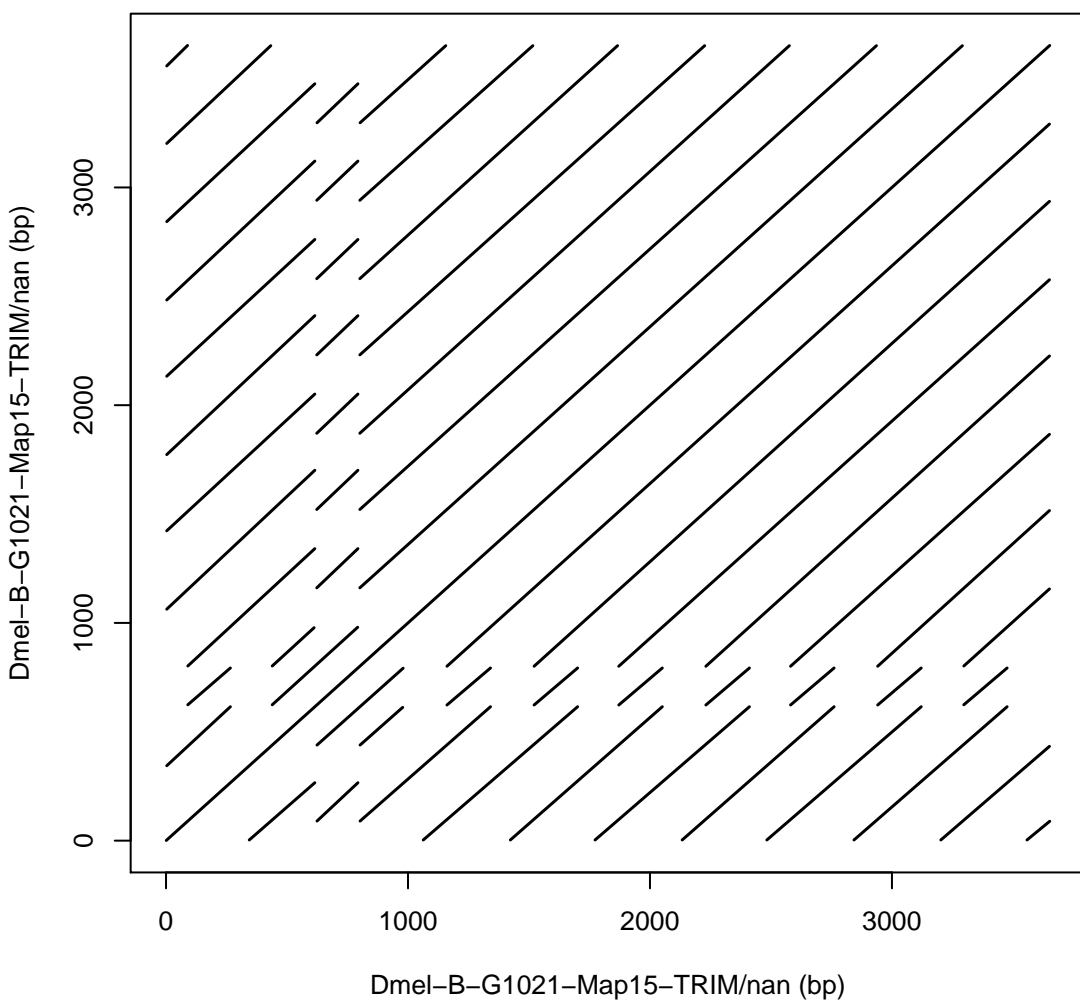
TE: Dmel-B-G1021-Map15-TRIM/nan  
consensus size: 3652bp; fragments: 37946; full length: 23 (>=3286.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

