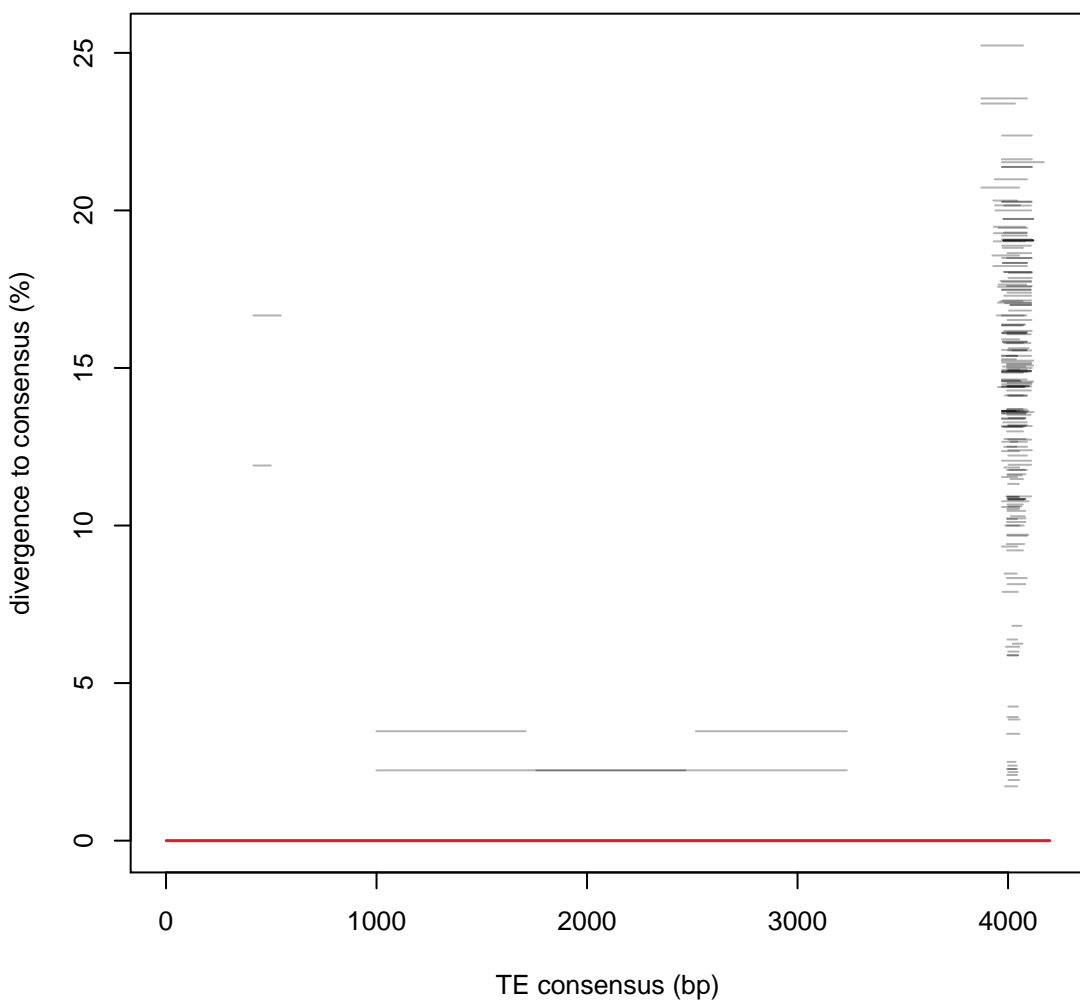
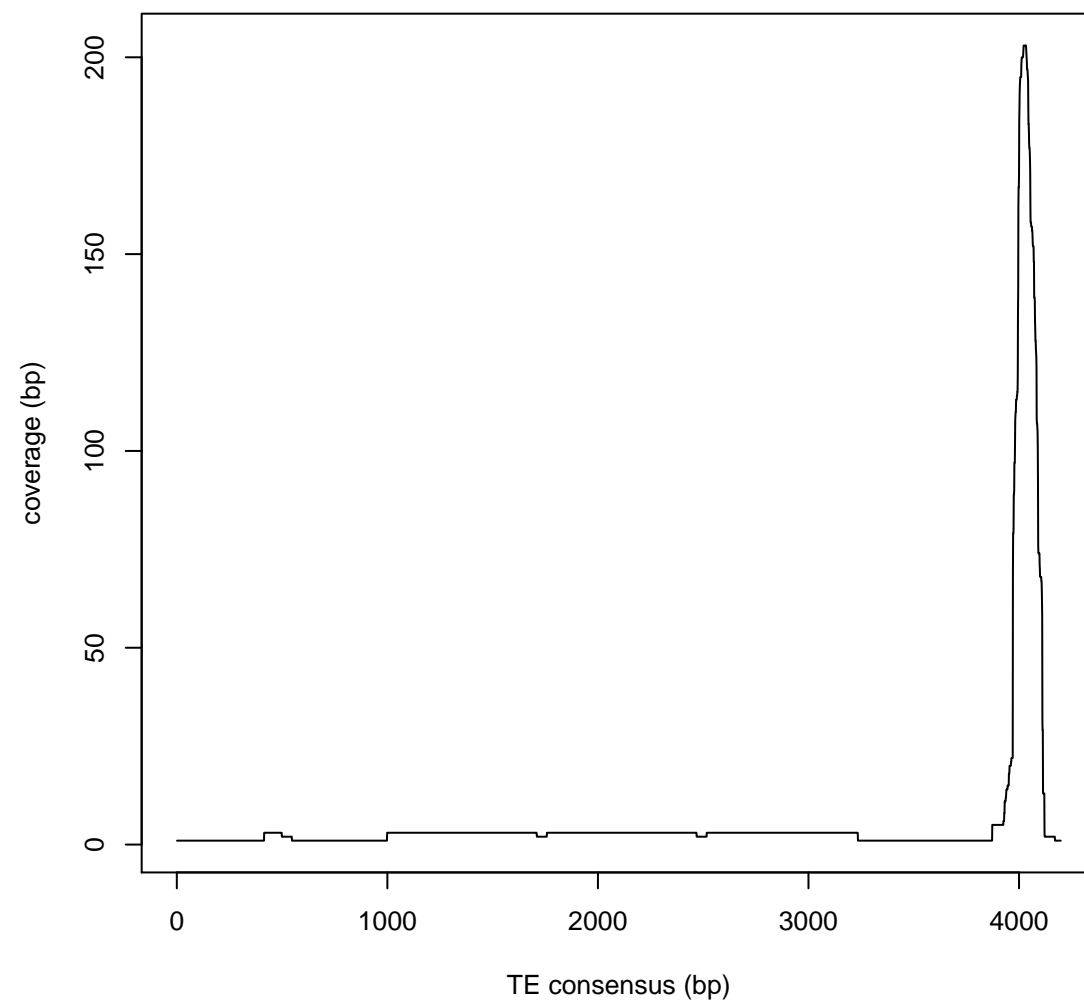


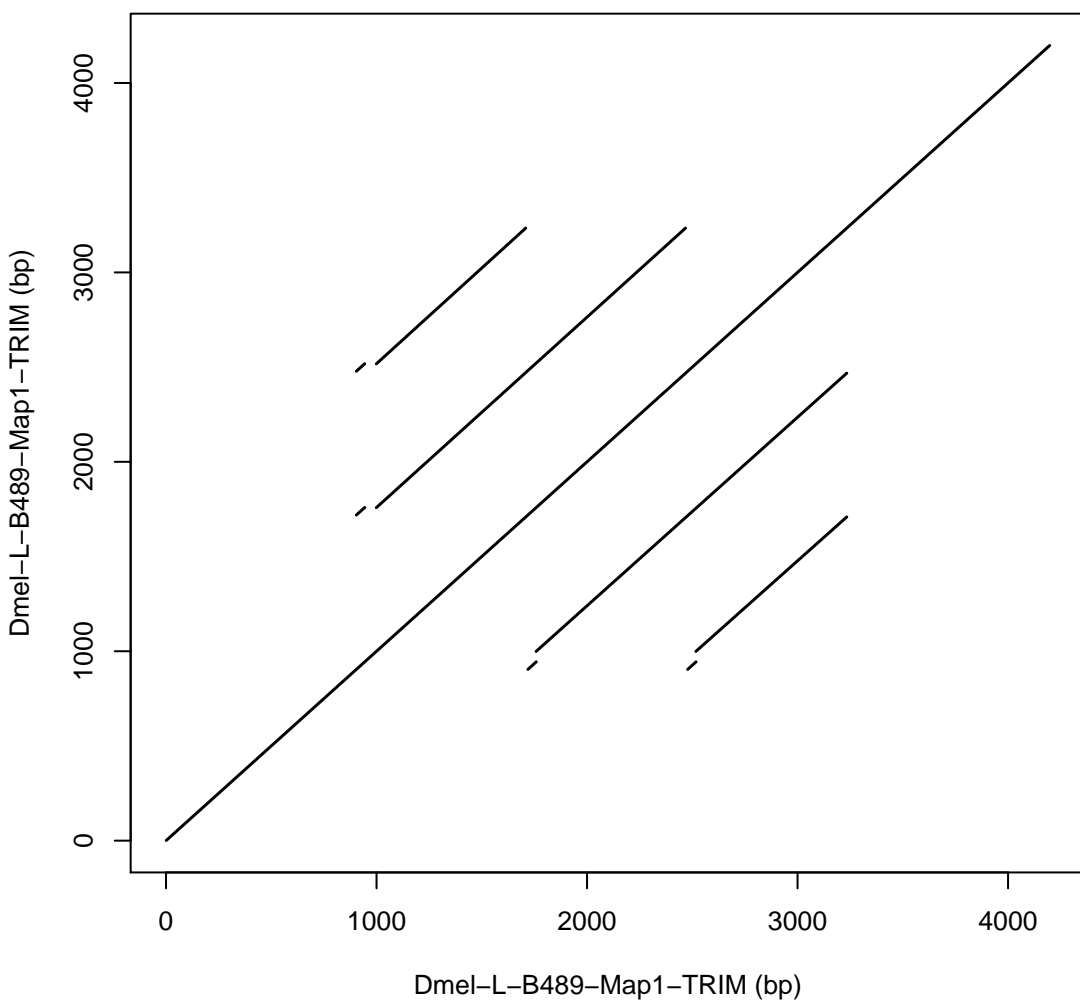
TE: Dmel-L-B489-Map1-TRIM  
consensus size: 4198bp; fragments: 209; full length: 1 (>=3778.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

