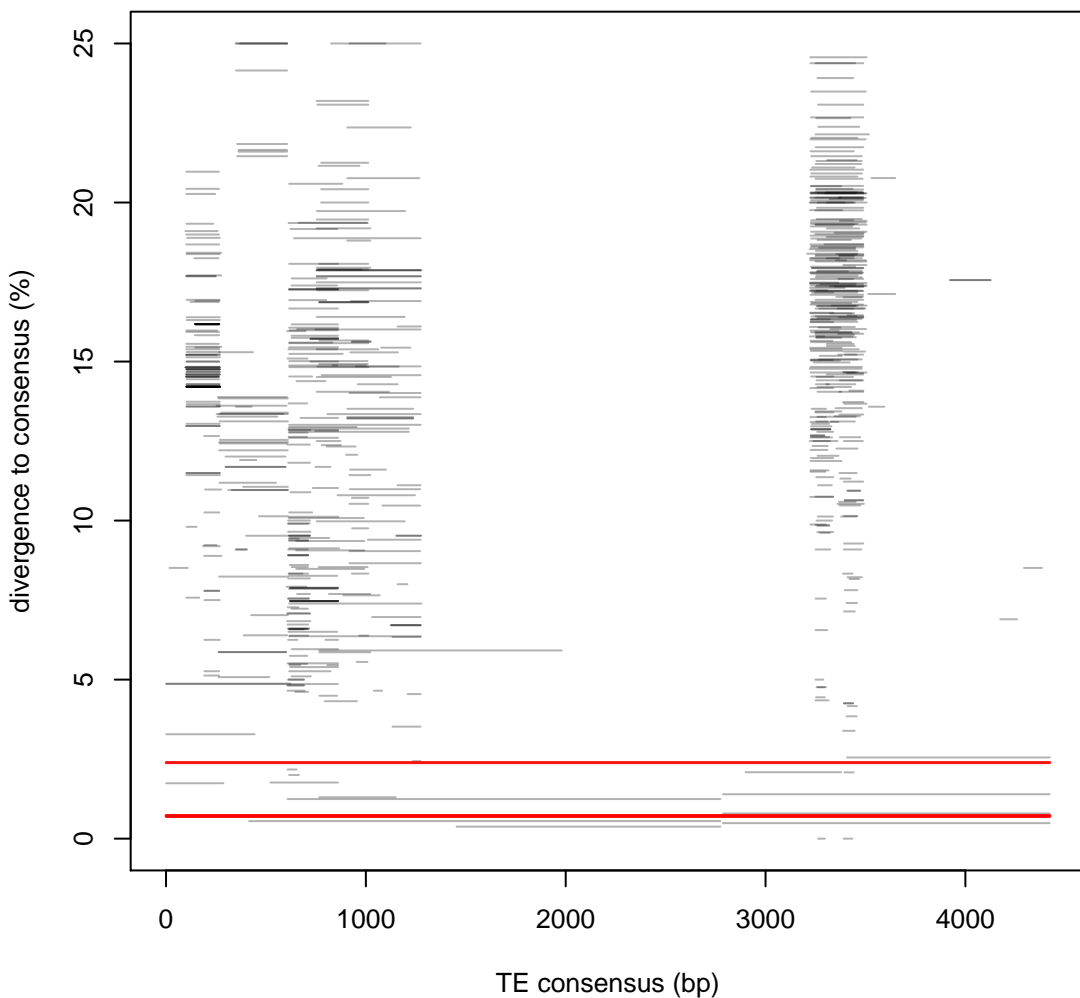
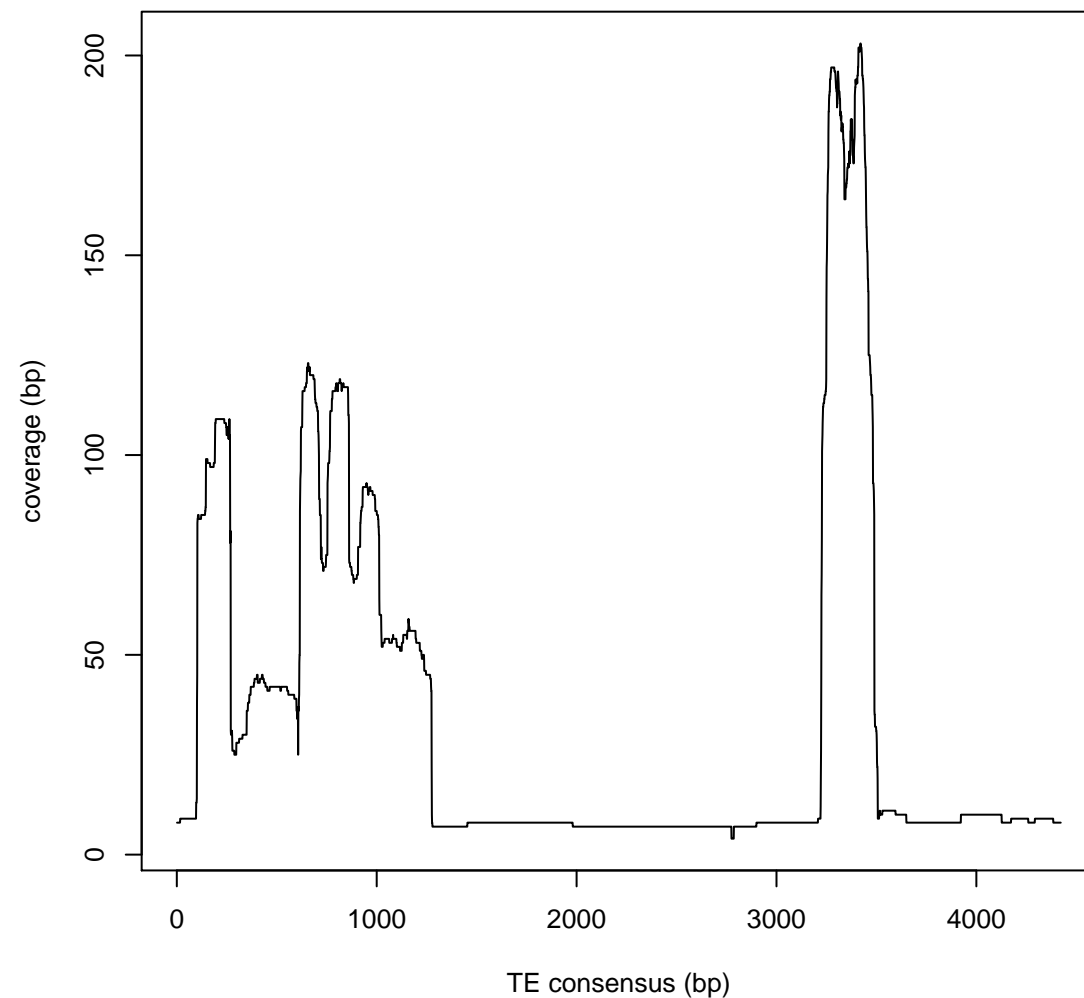


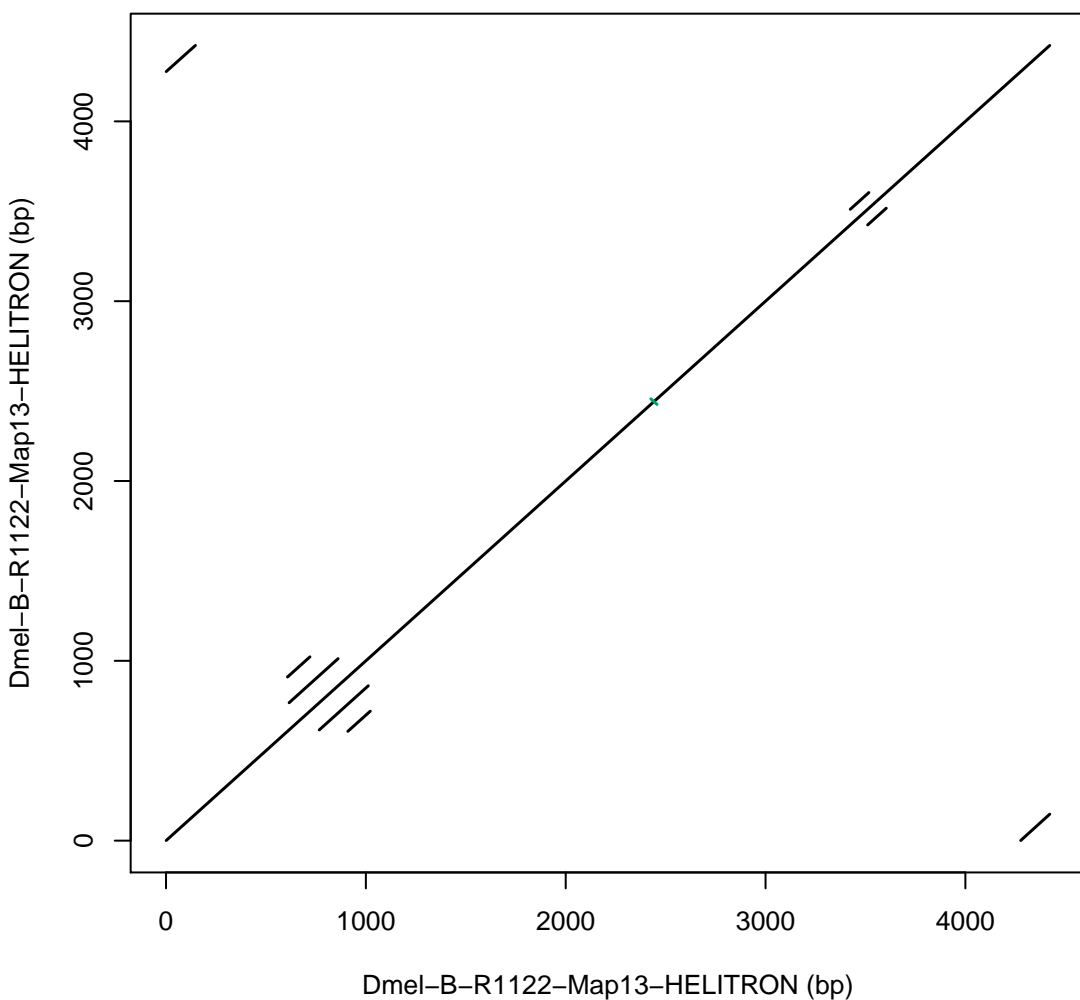
TE: Dmel-B-R1122-Map13-HELITRON  
consensus size: 4422bp; fragments: 659; full length: 4 (>=3979.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

