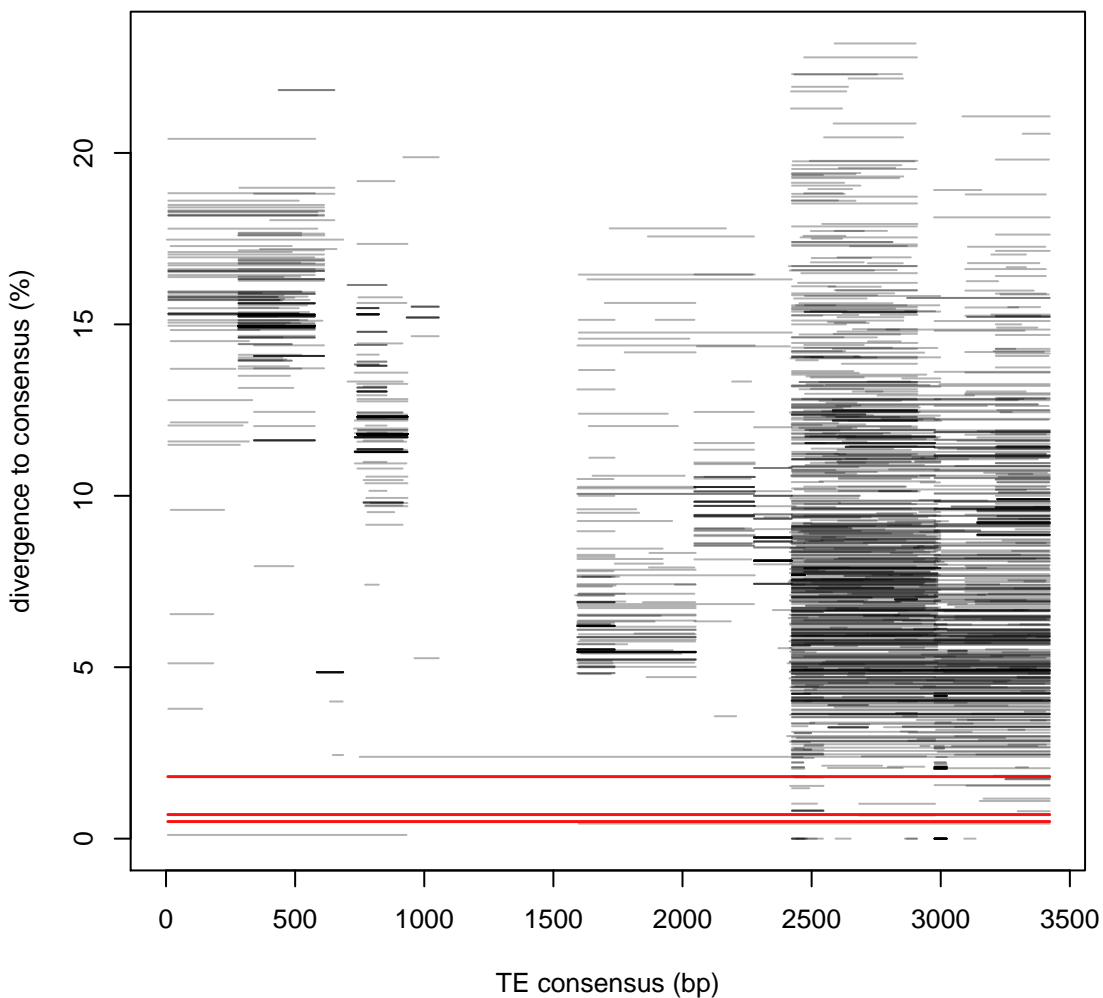
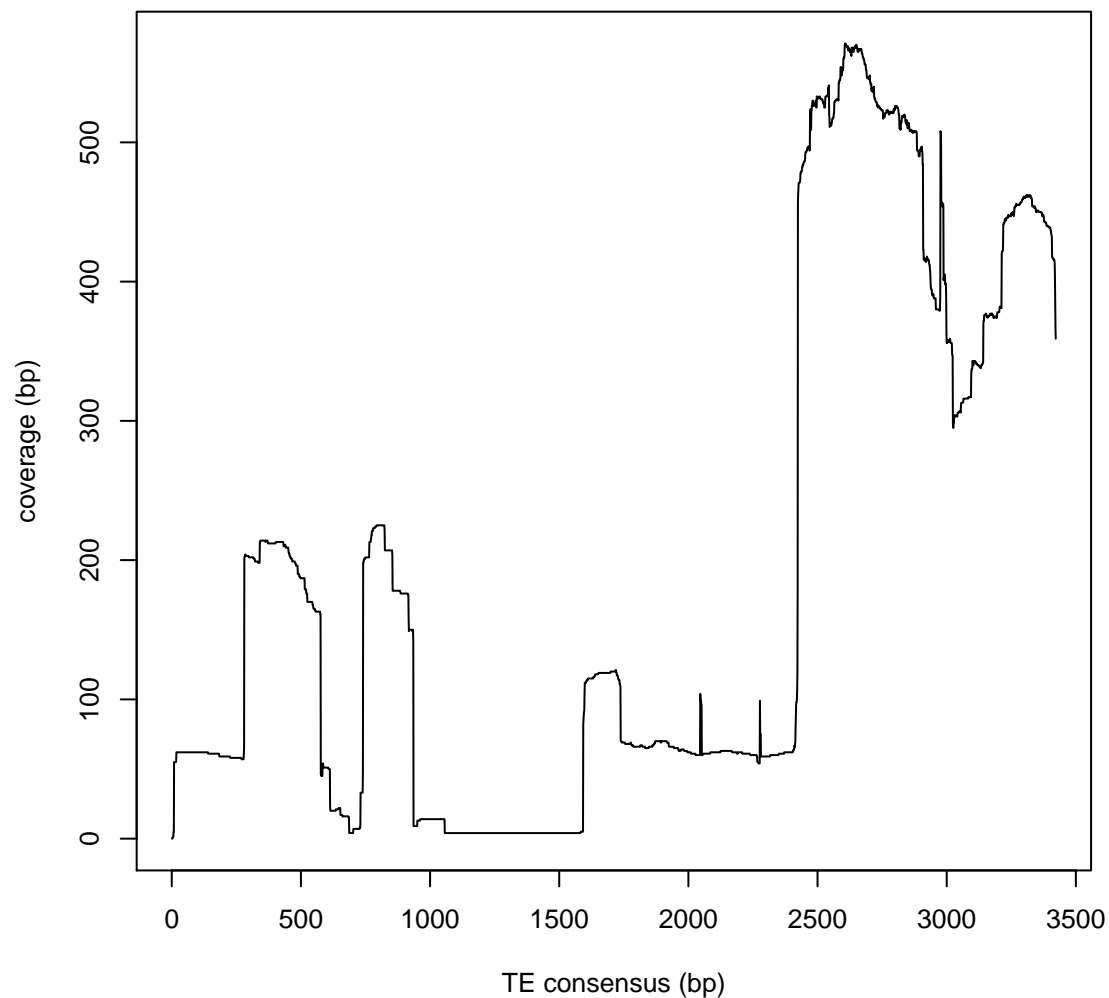


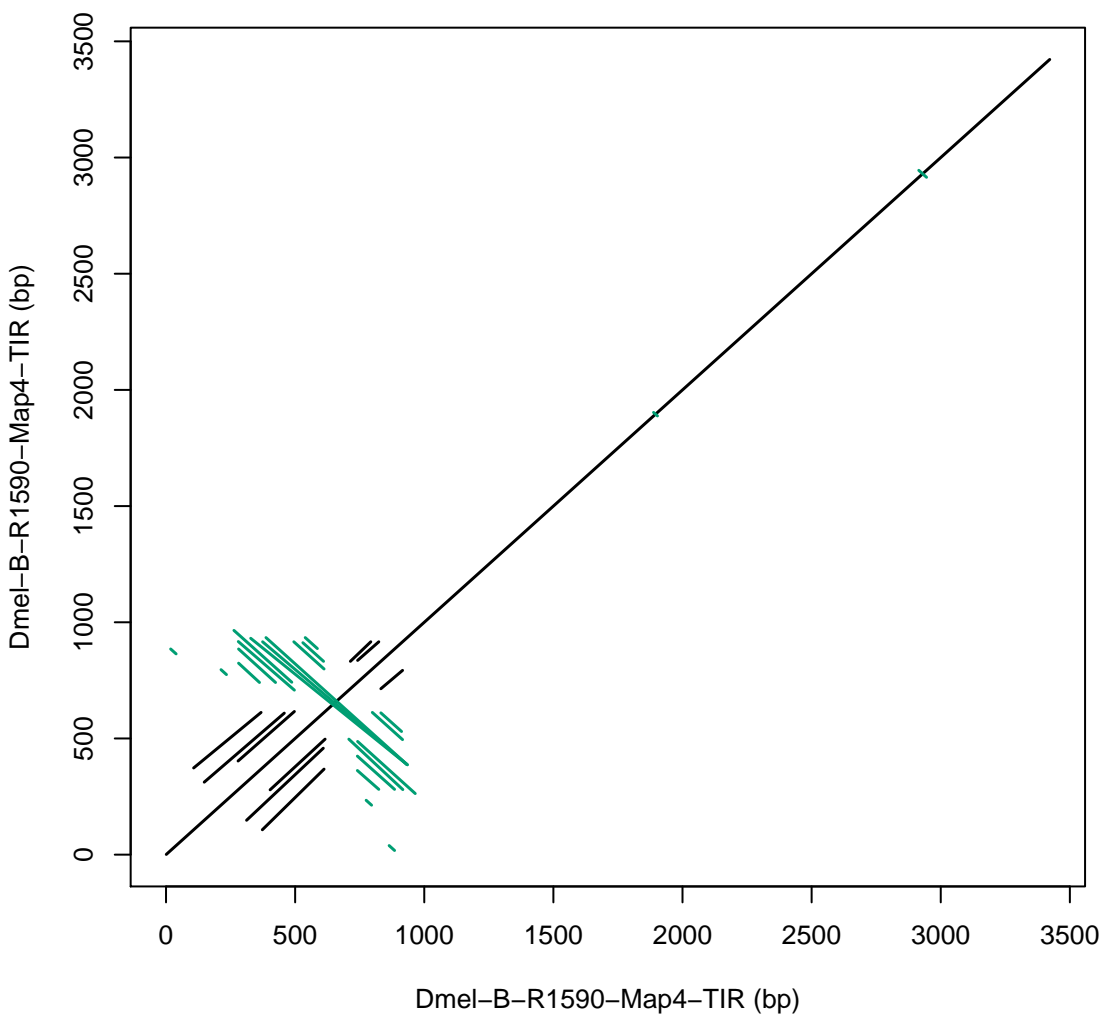
TE: Dmel-B-R1590-Map4-TIR
consensus size: 3422bp; fragments: 1874; full length: 3 (>=3079.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

