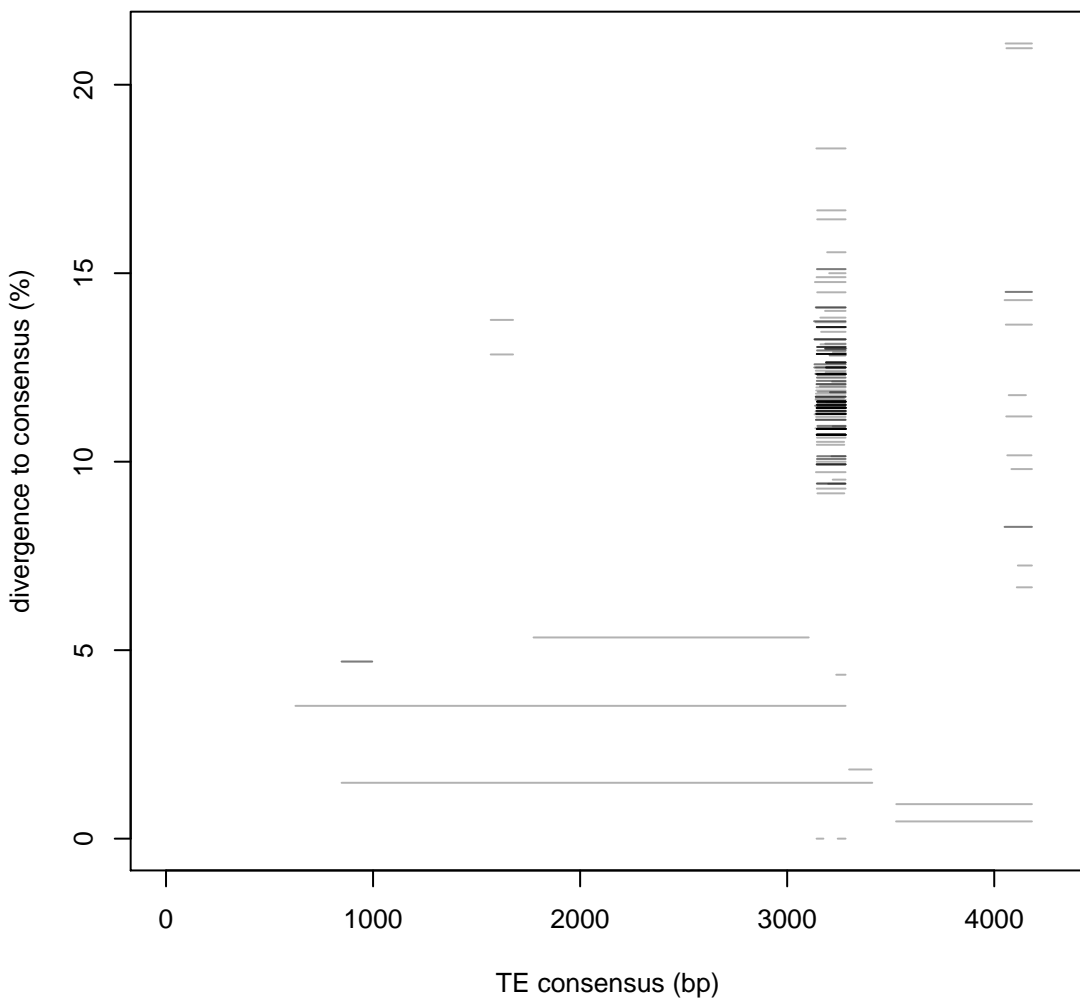
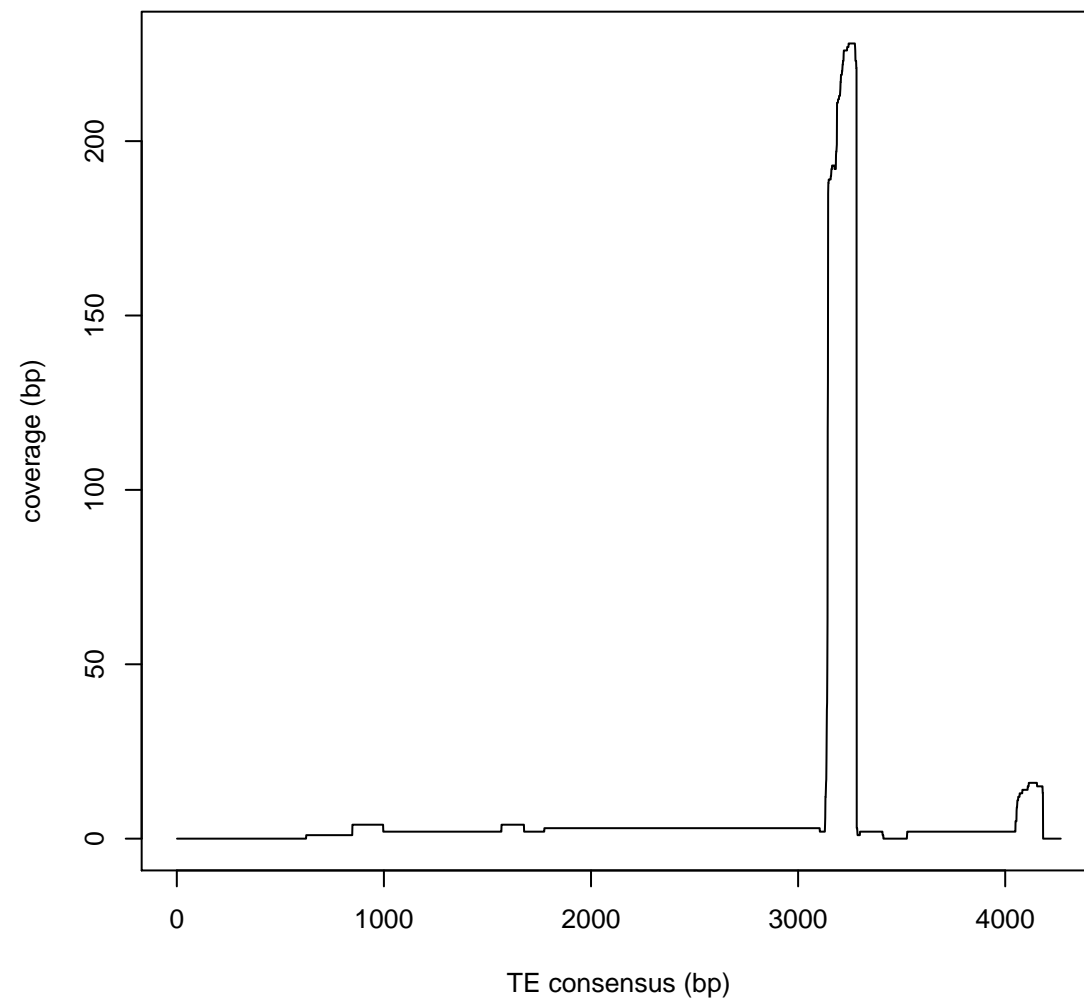


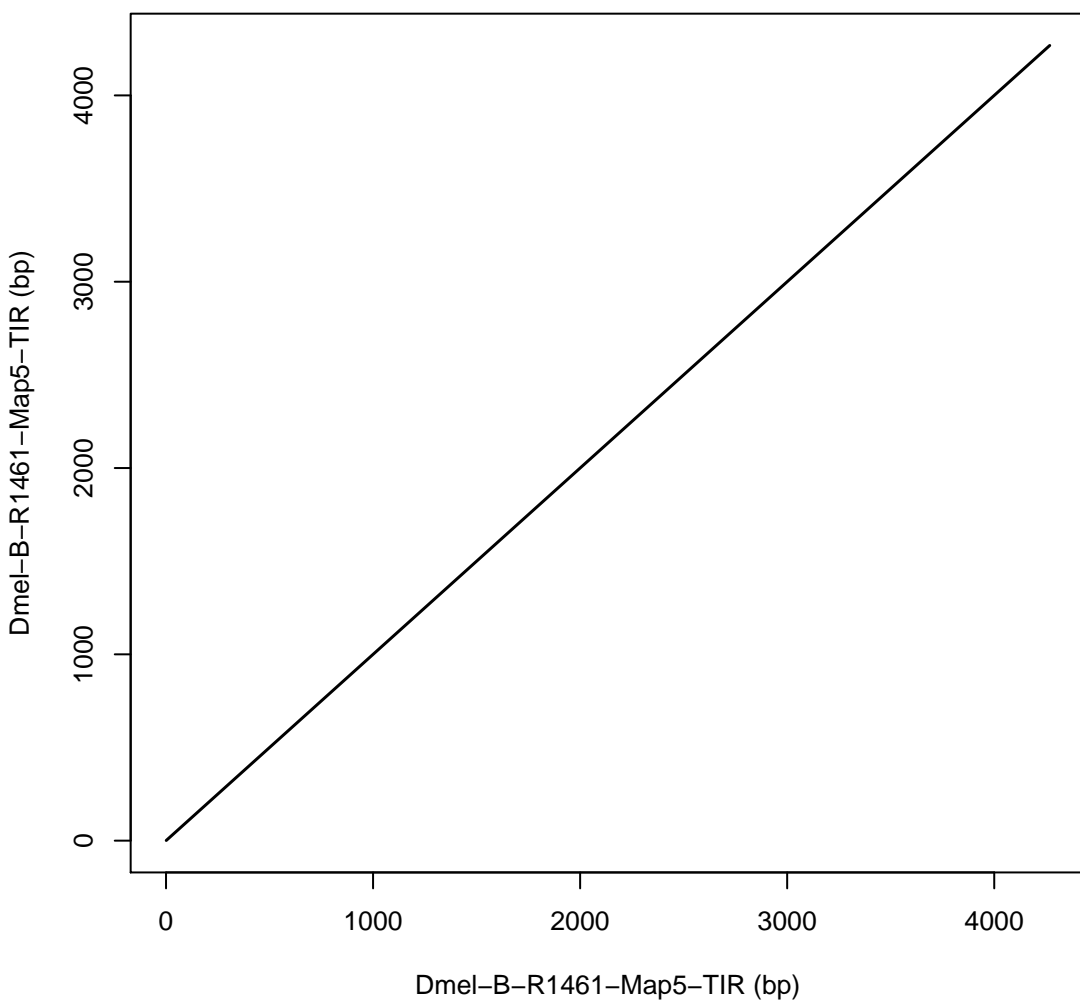
TE: Dmel-B-R1461-Map5-TIR
consensus size: 4268bp; fragments: 251; full length: 0 (>=3841.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

