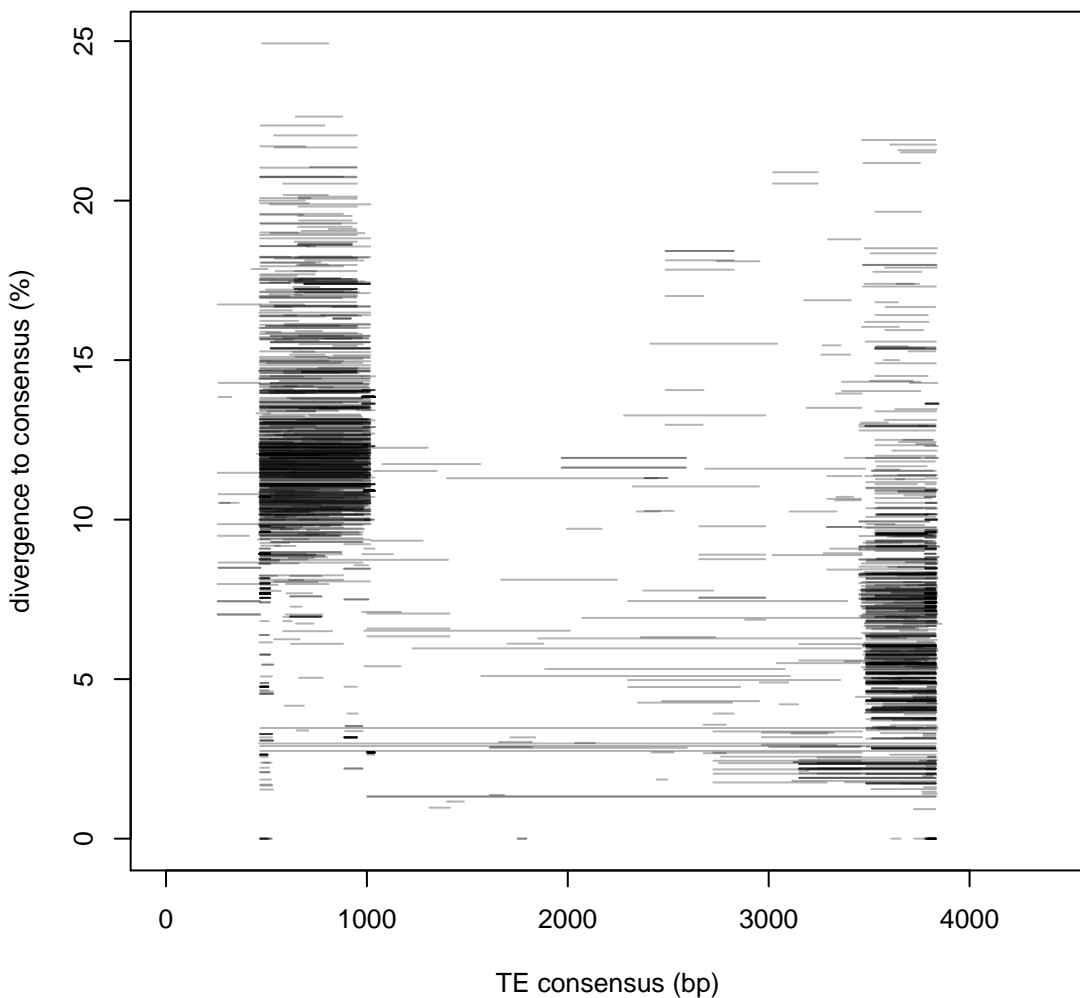
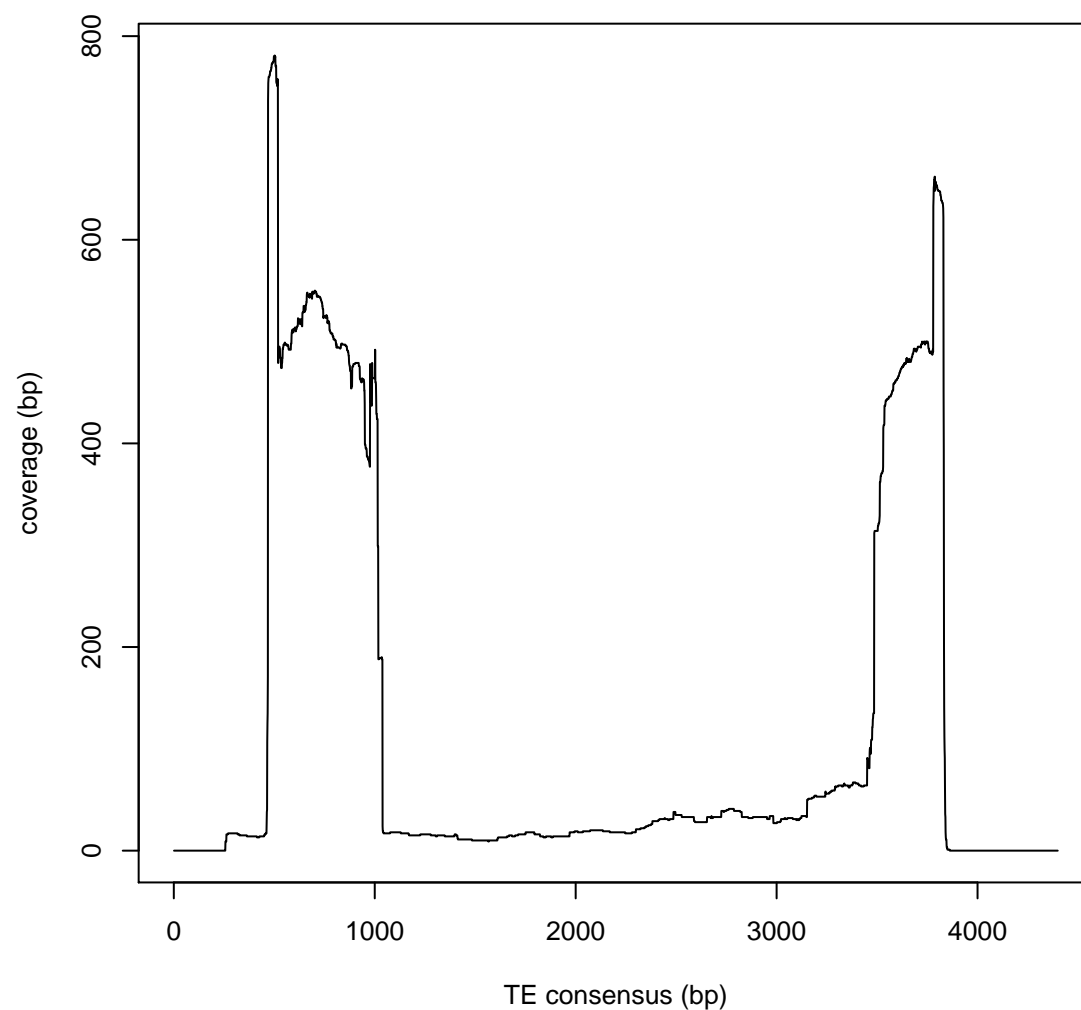


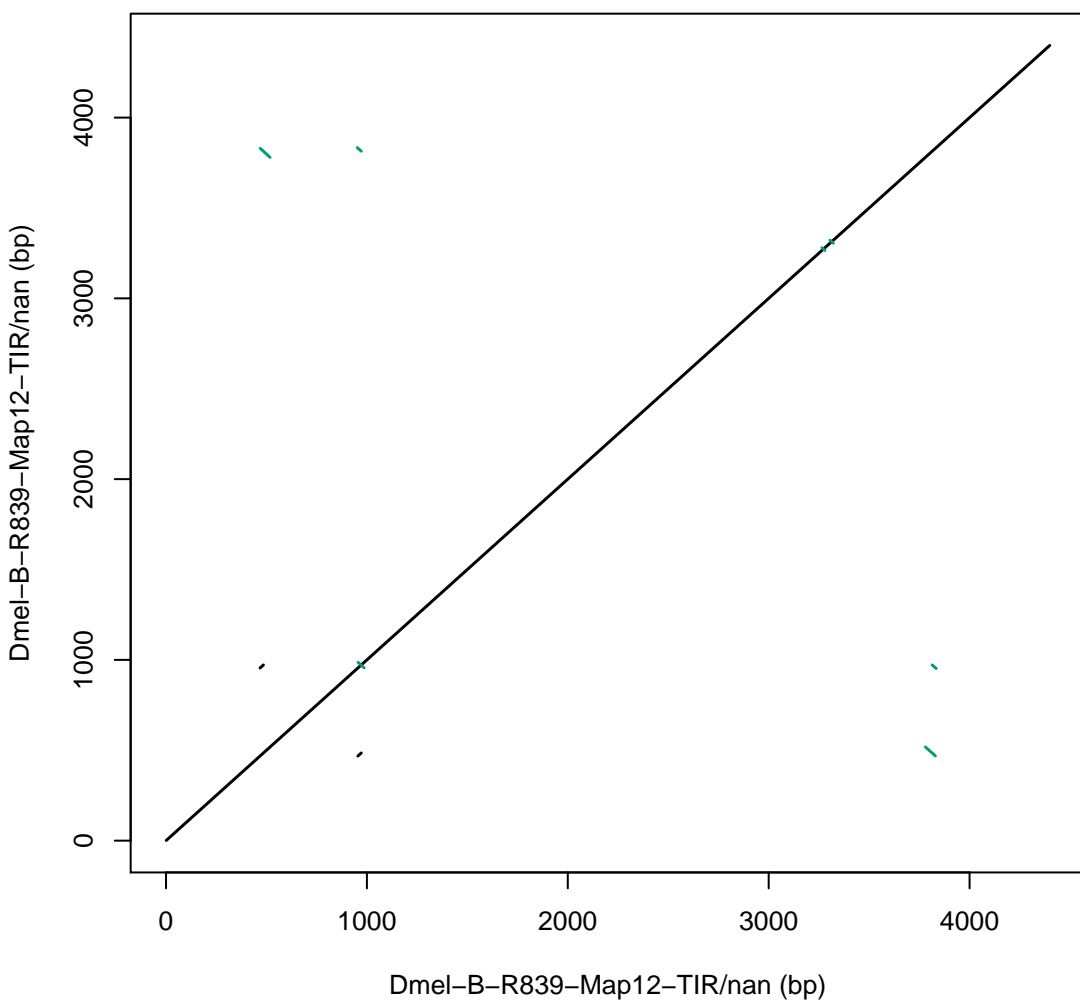
TE: Dmel-B-R839-Map12-TIR/nan  
consensus size: 4399bp; fragments: 2058; full length: 0 (>=3959.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

