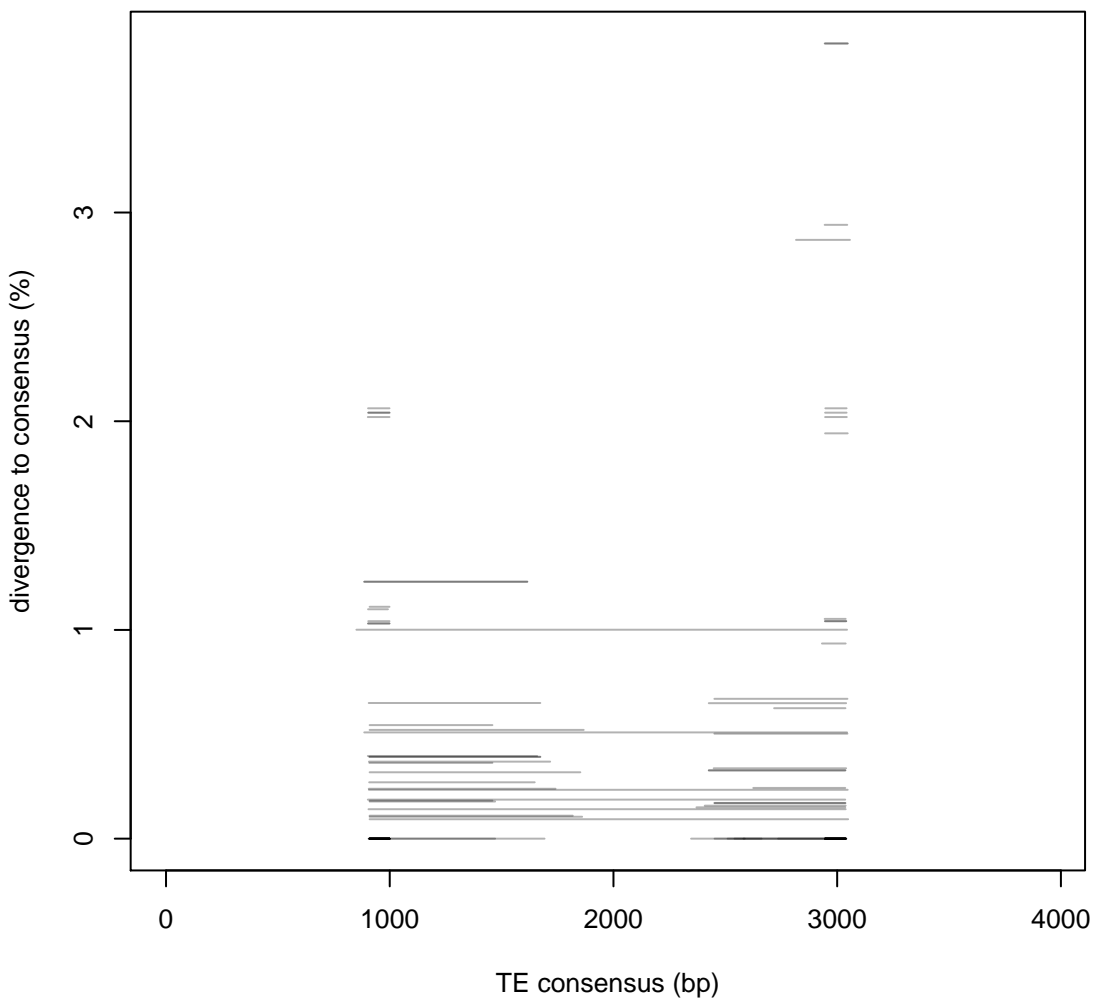
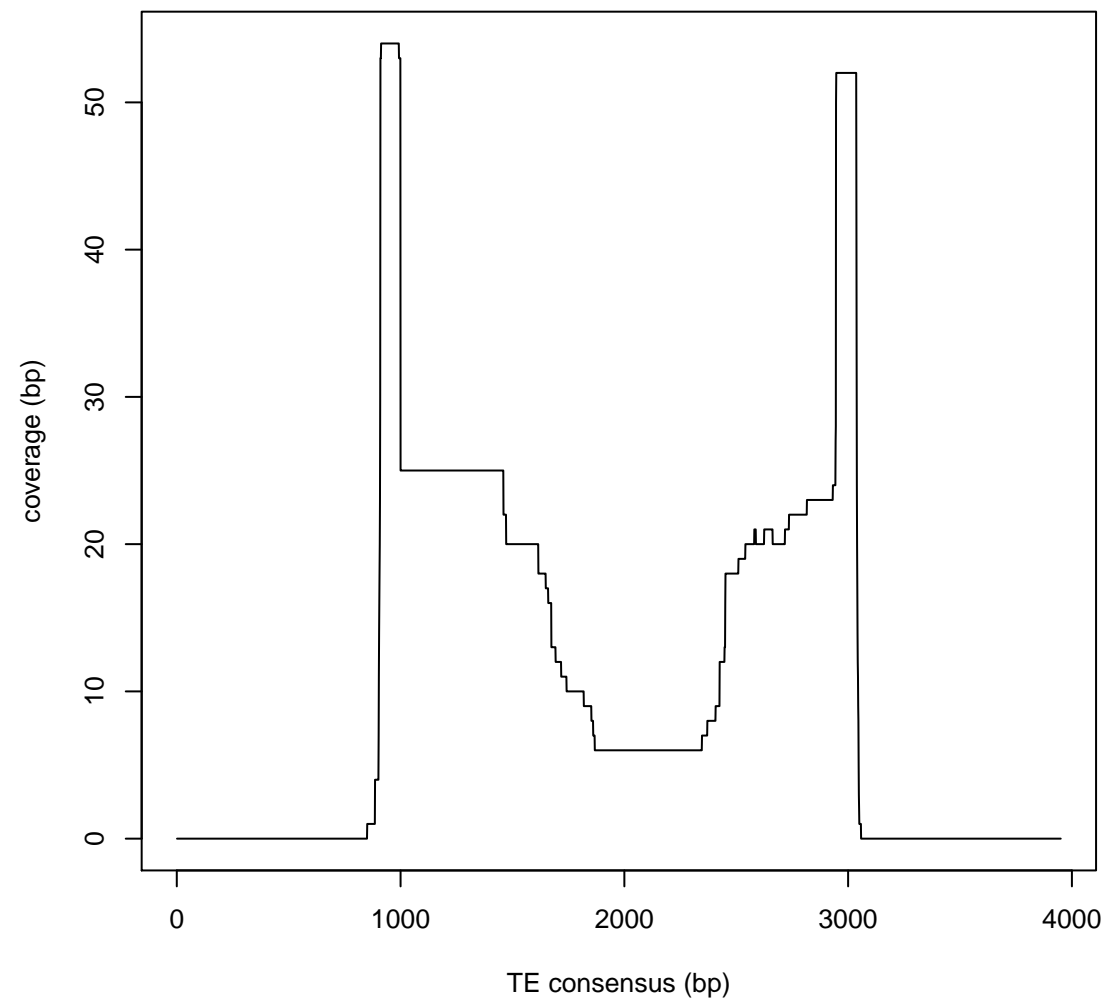


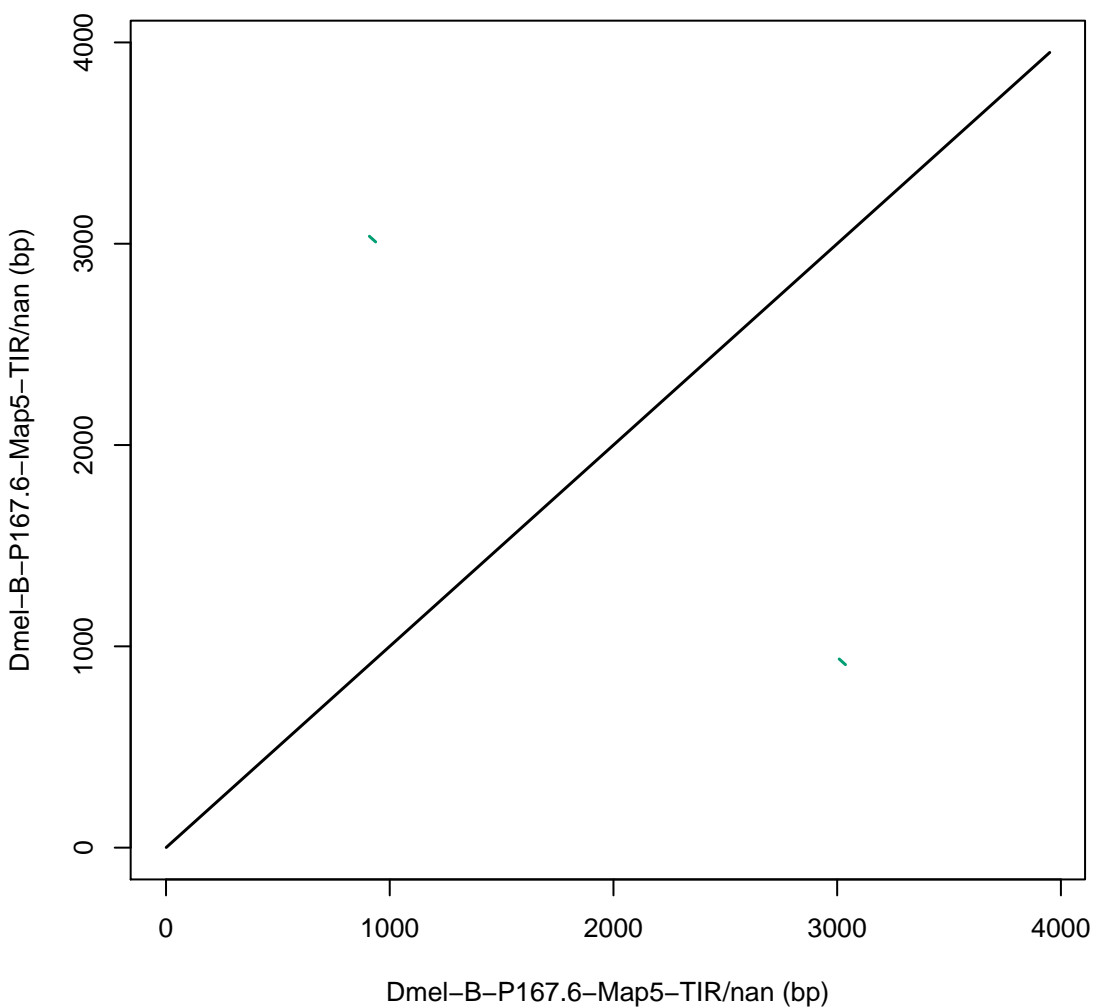
TE: Dmel-B-P167.6-Map5-TIR/nan
consensus size: 3950bp; fragments: 102; full length: 0 (>=3555bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

