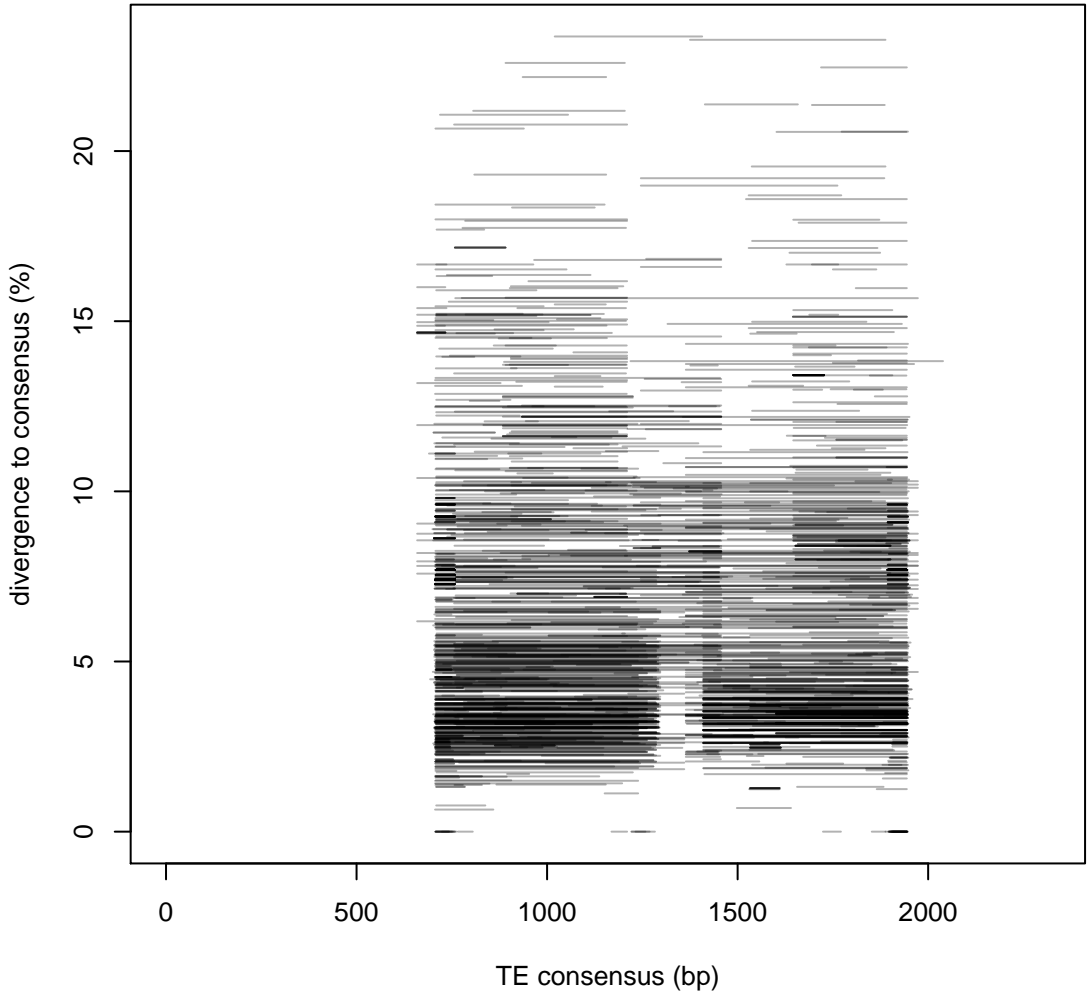
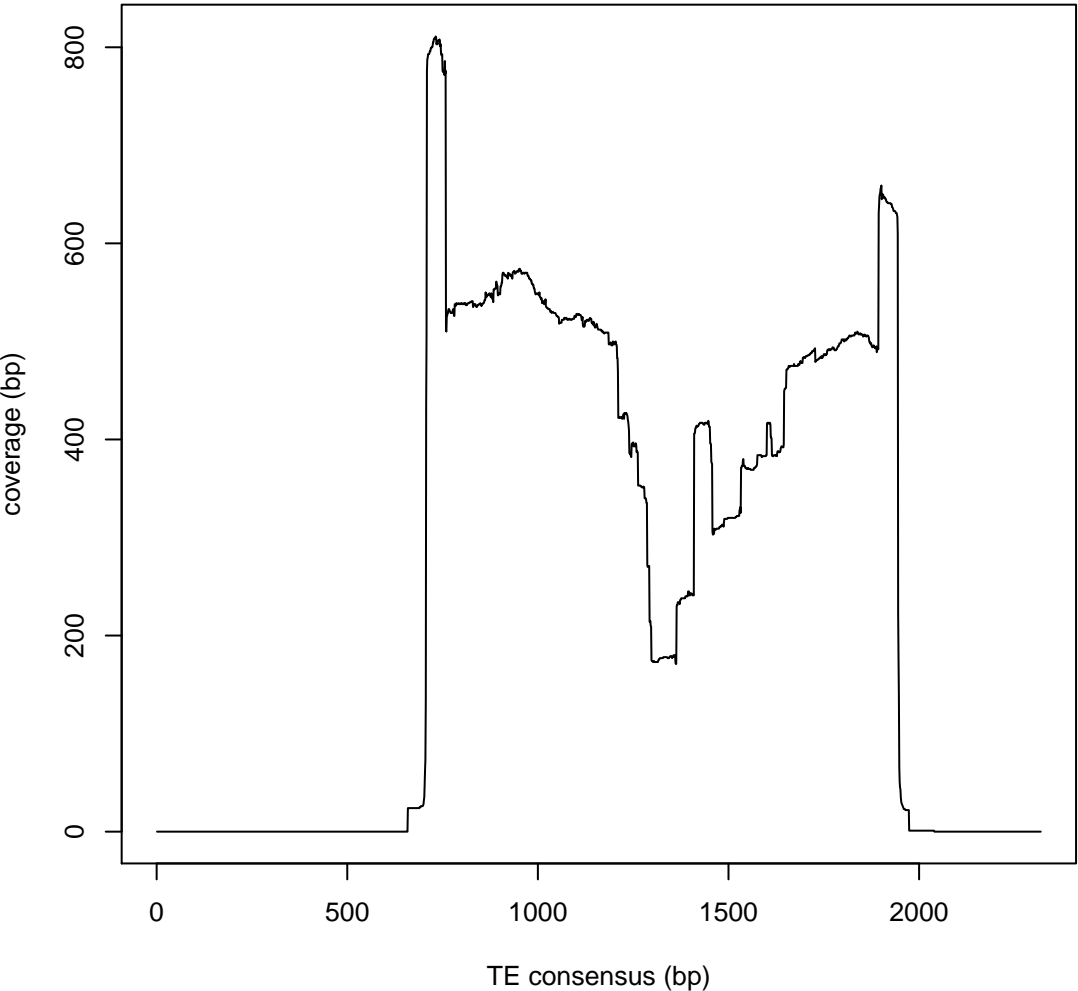


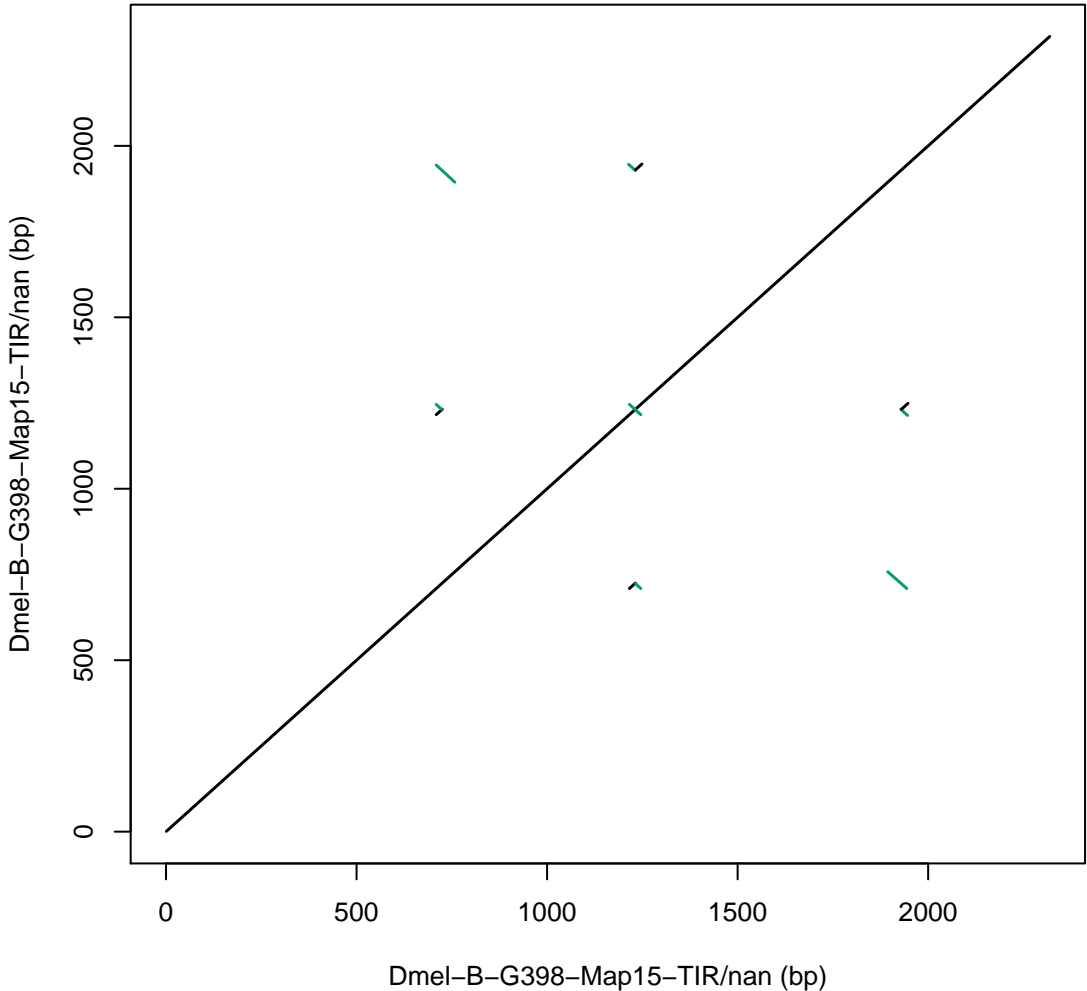
TE: Dmel-B-G398-Map15-TIR/nan
consensus size: 2319bp; fragments: 2003; full length: 0 (>=2087.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

