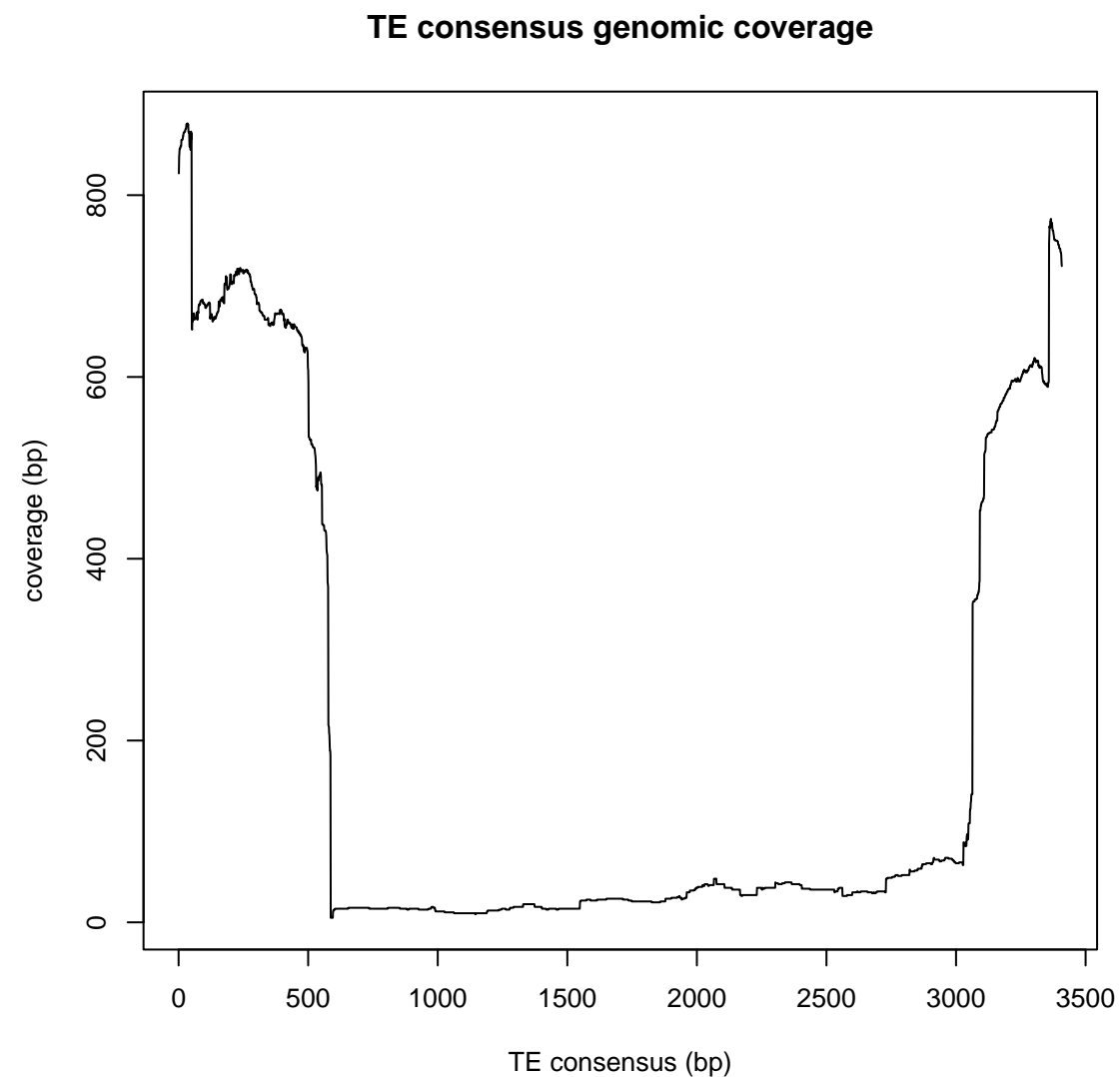
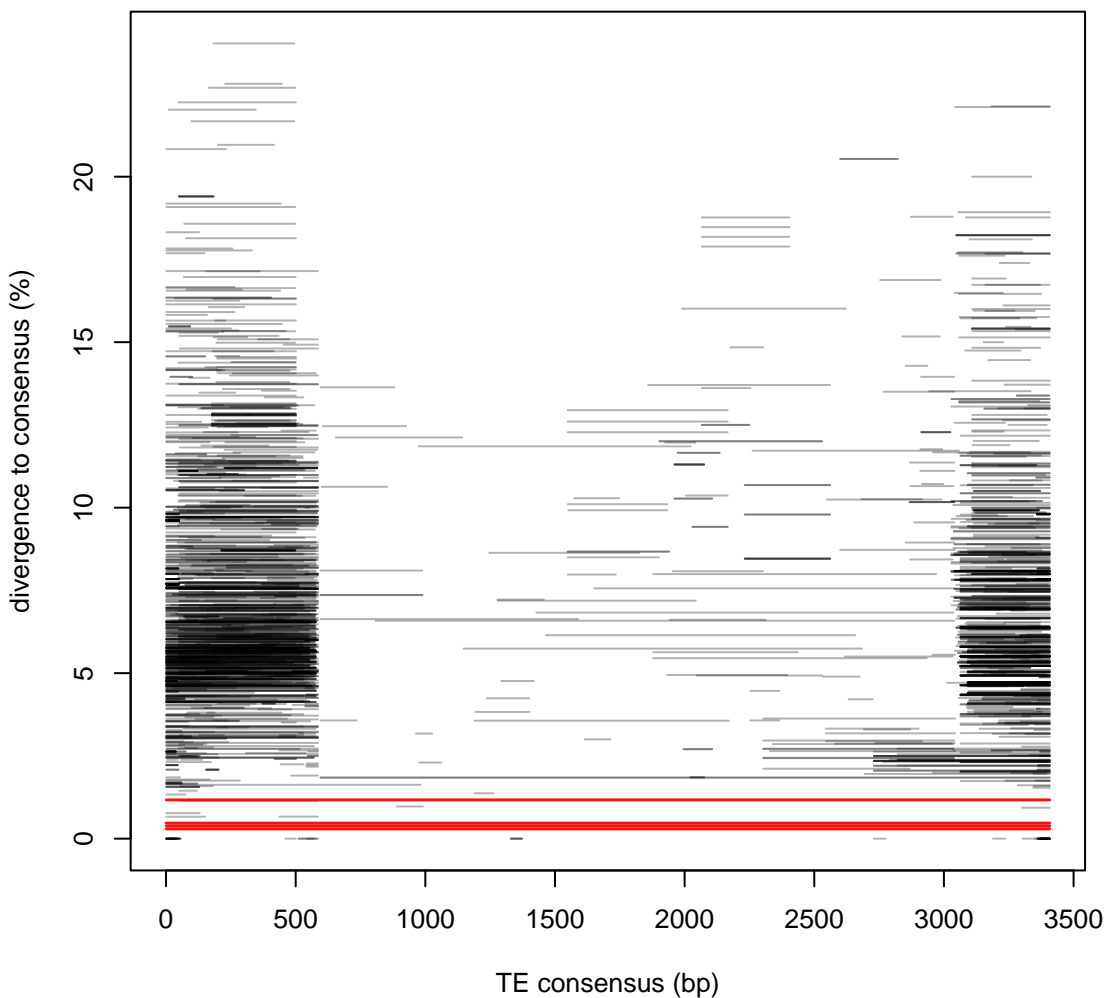
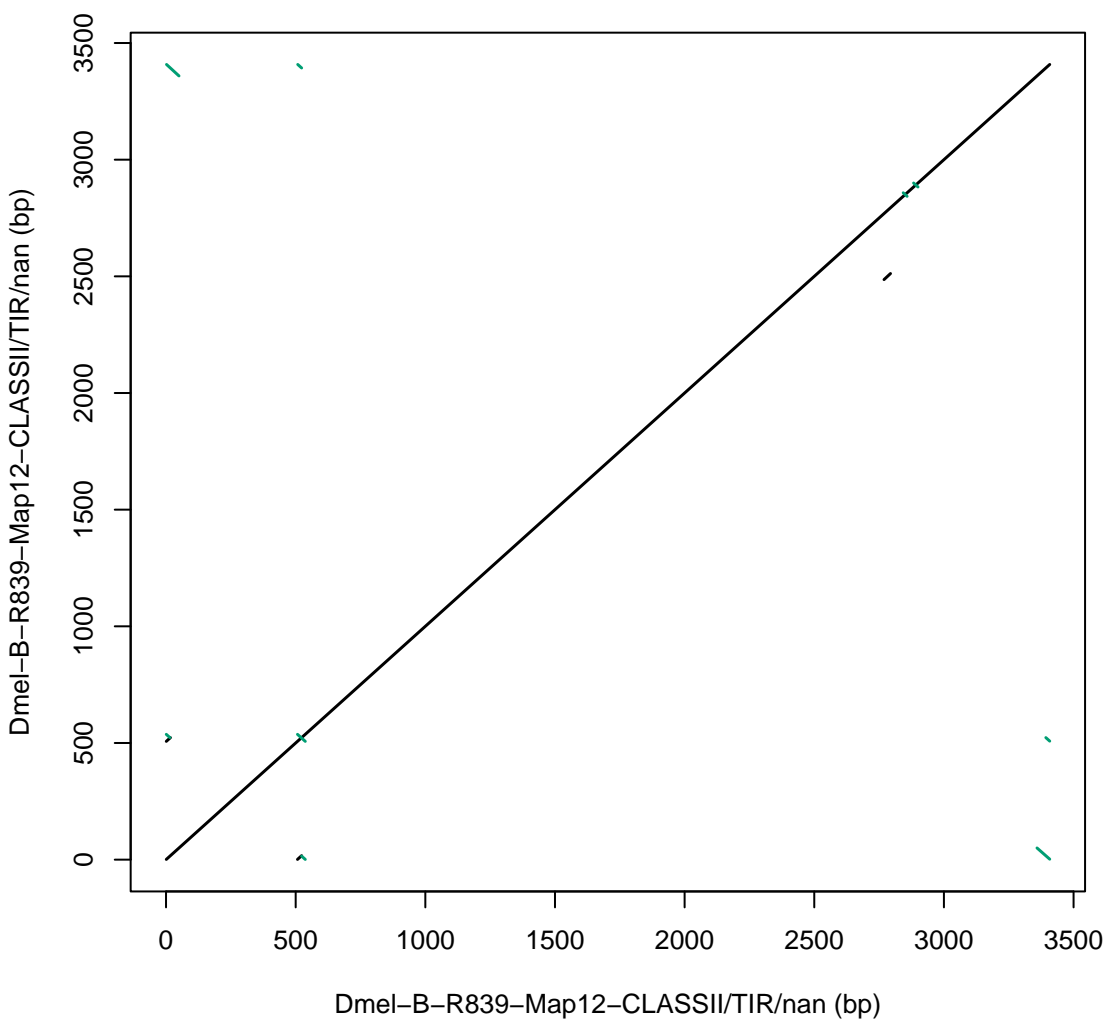


TE: Dmel-B-R839-Map12-CLASSII/TIR/nan
consensus size: 3408bp; fragments: 2390; full length: 4 (>=3067.2bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

