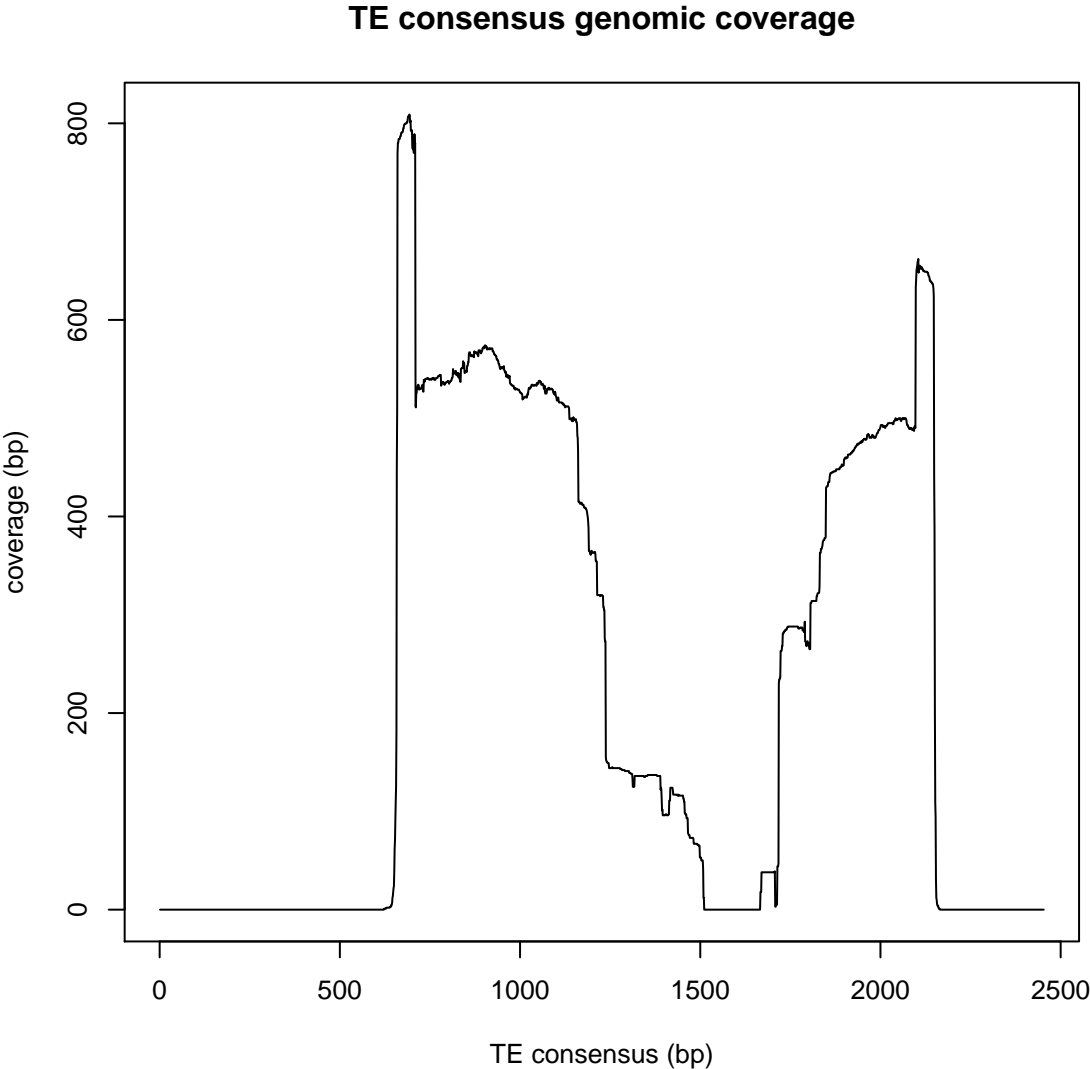
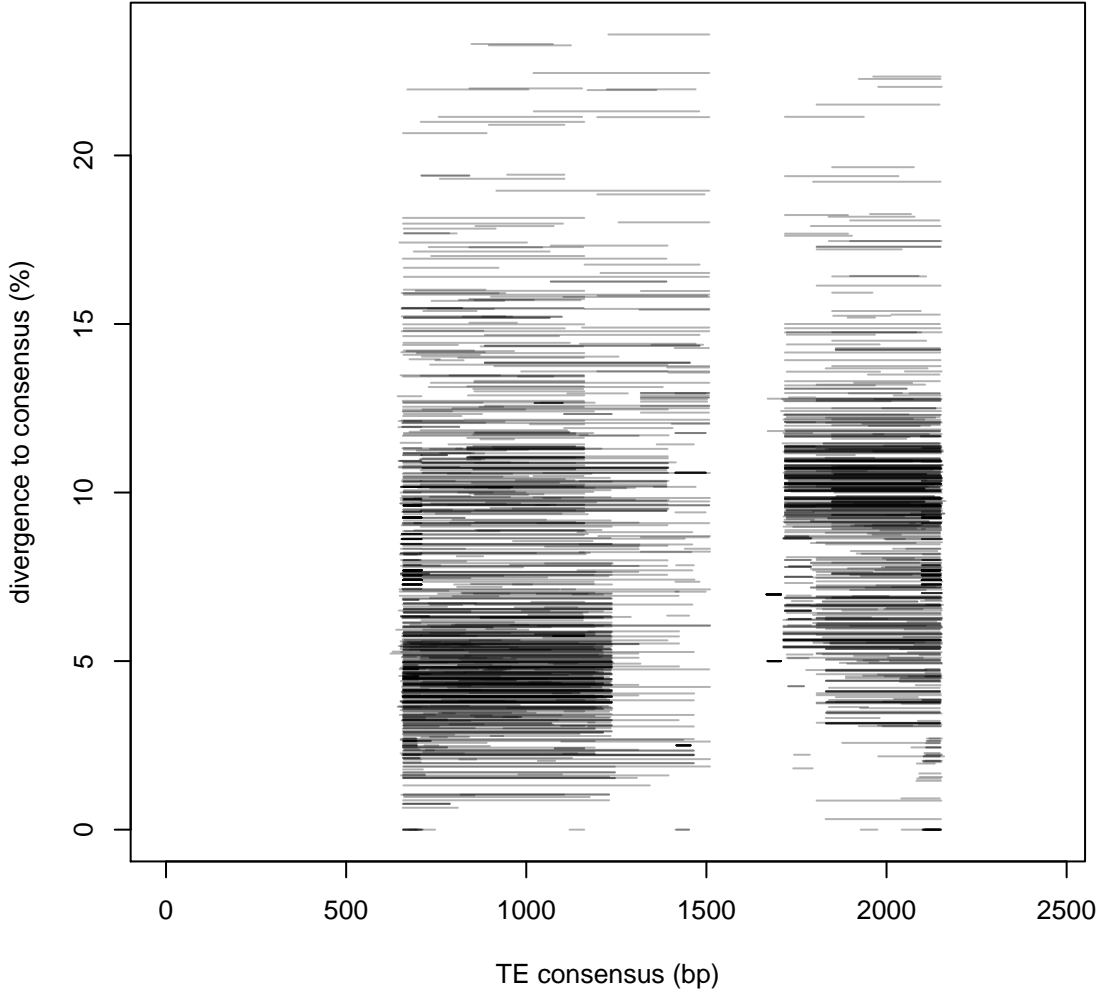
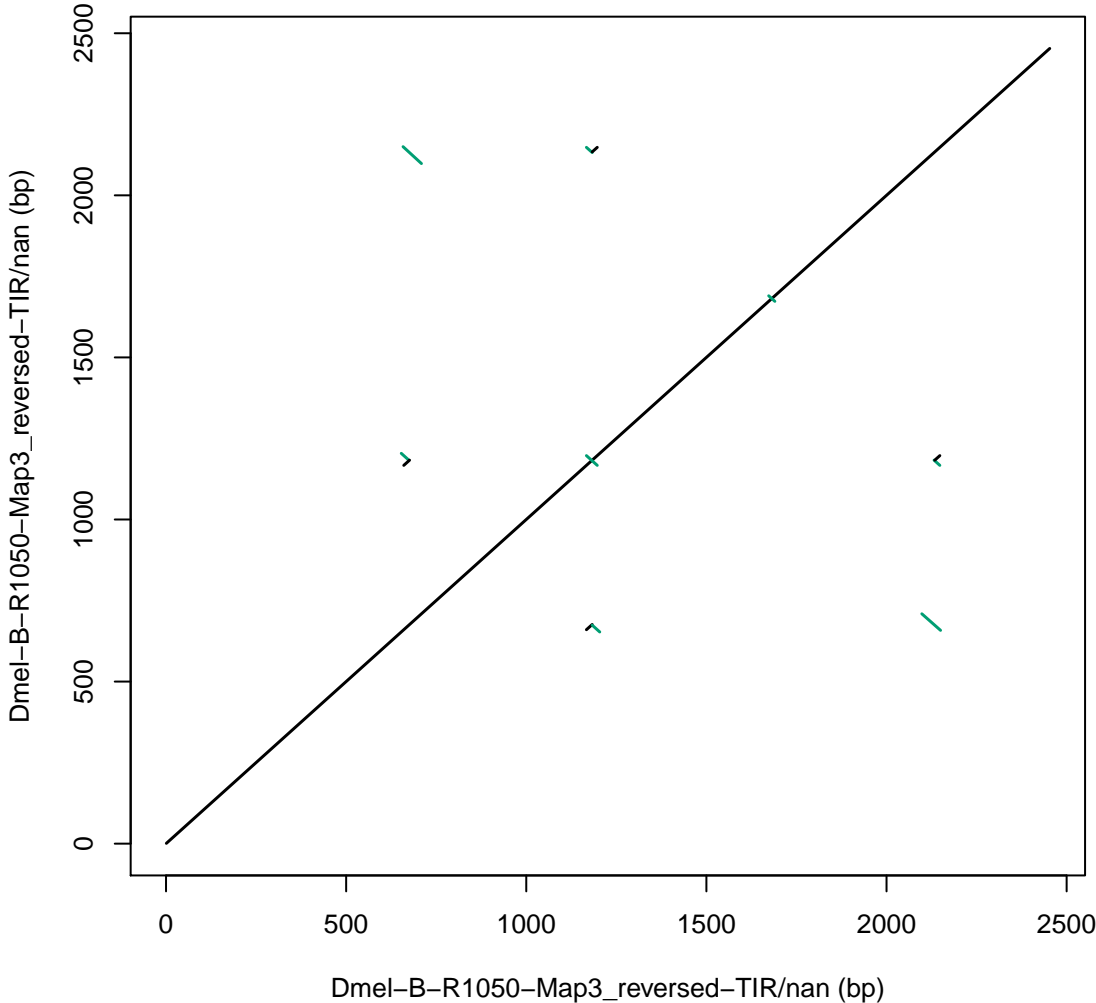


TE: Dmel-B-R1050-Map3_reversed-TIR/nan
consensus size: 2453bp; fragments: 2025; full length: 0 (>=2207.7bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

