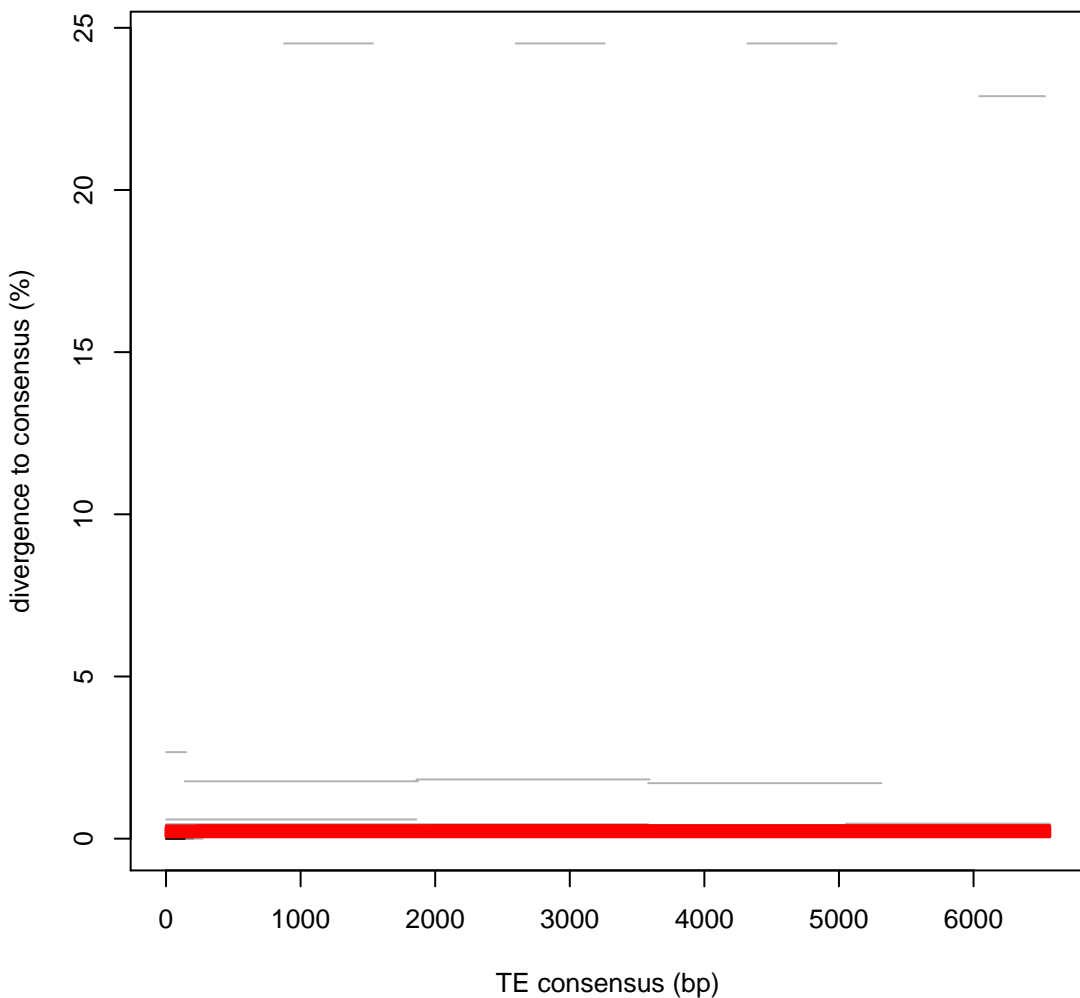
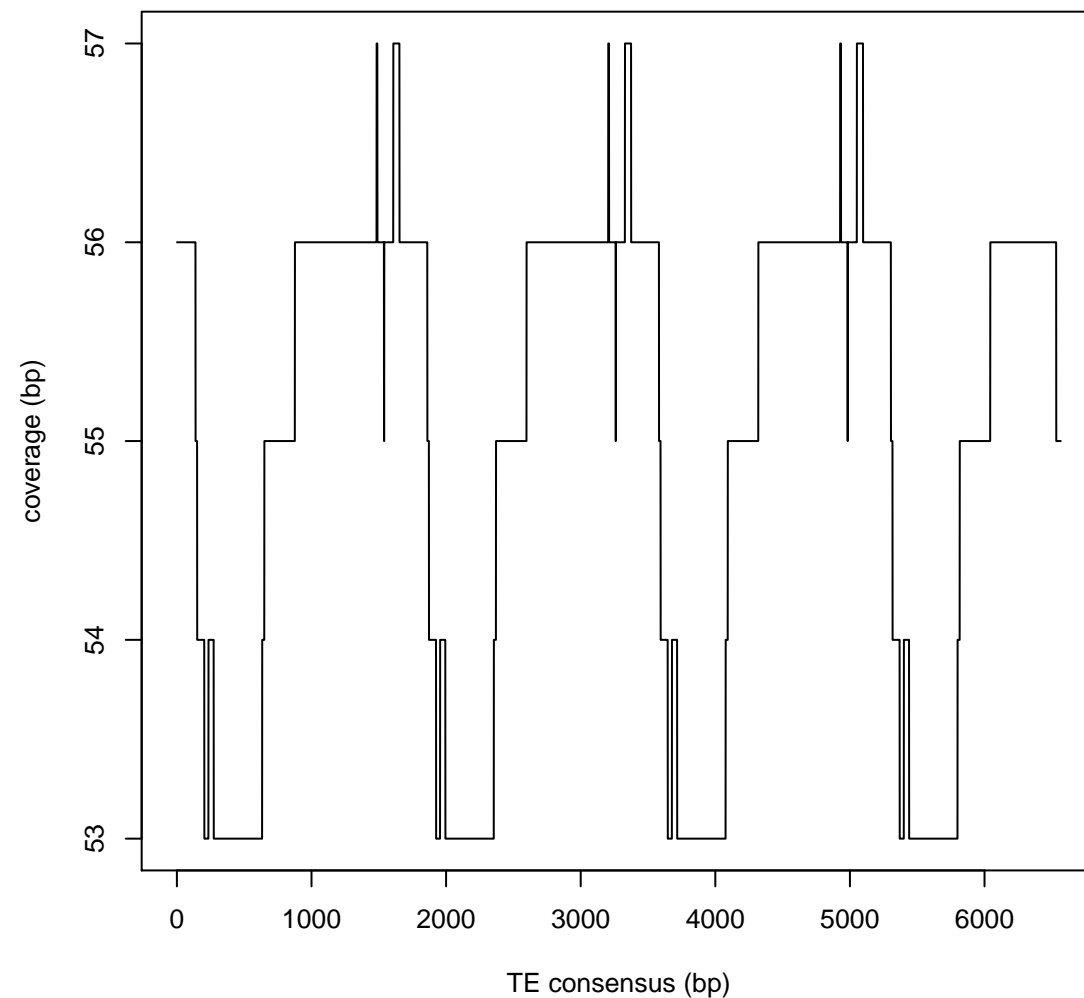


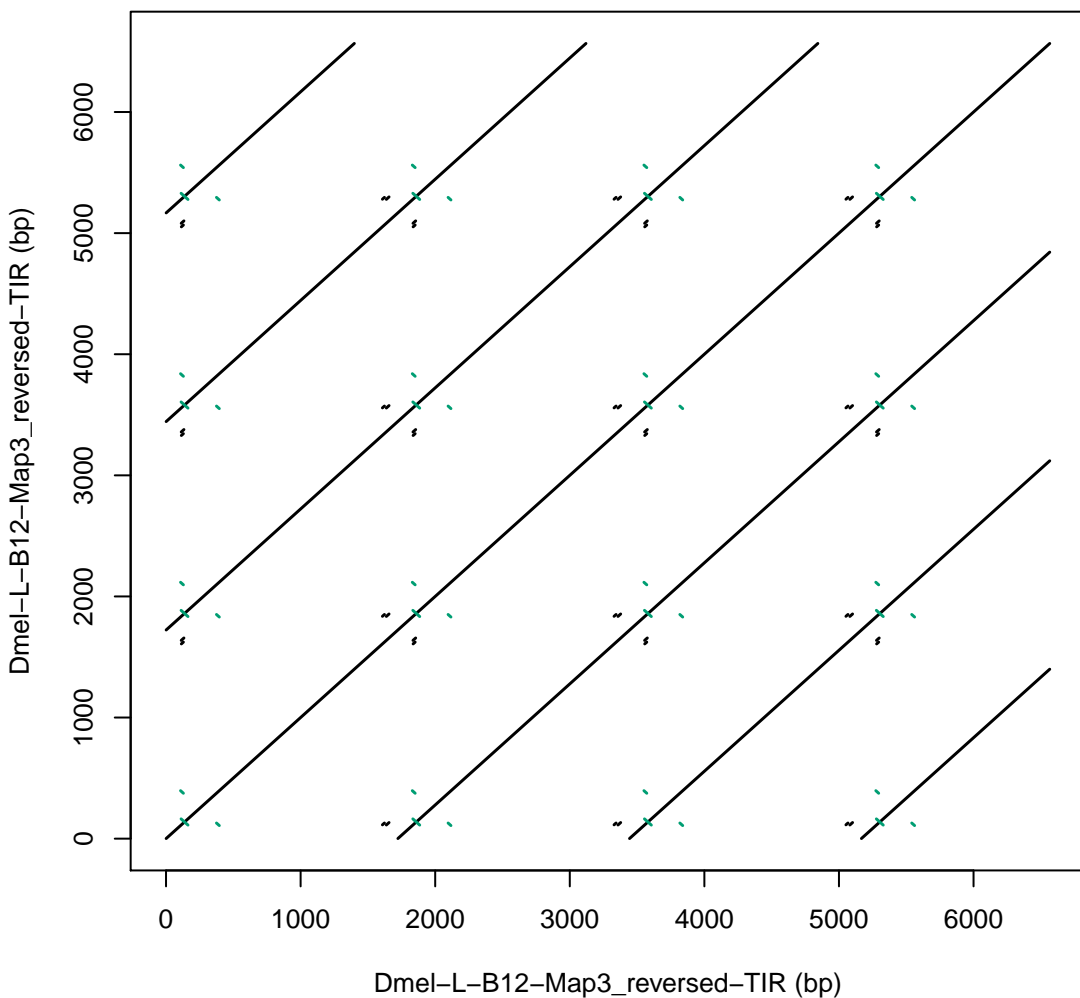
TE: Dmel-L-B12-Map3_reversed-TIR
consensus size: 6566bp; fragments: 105; full length: 34 (>=5909.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

