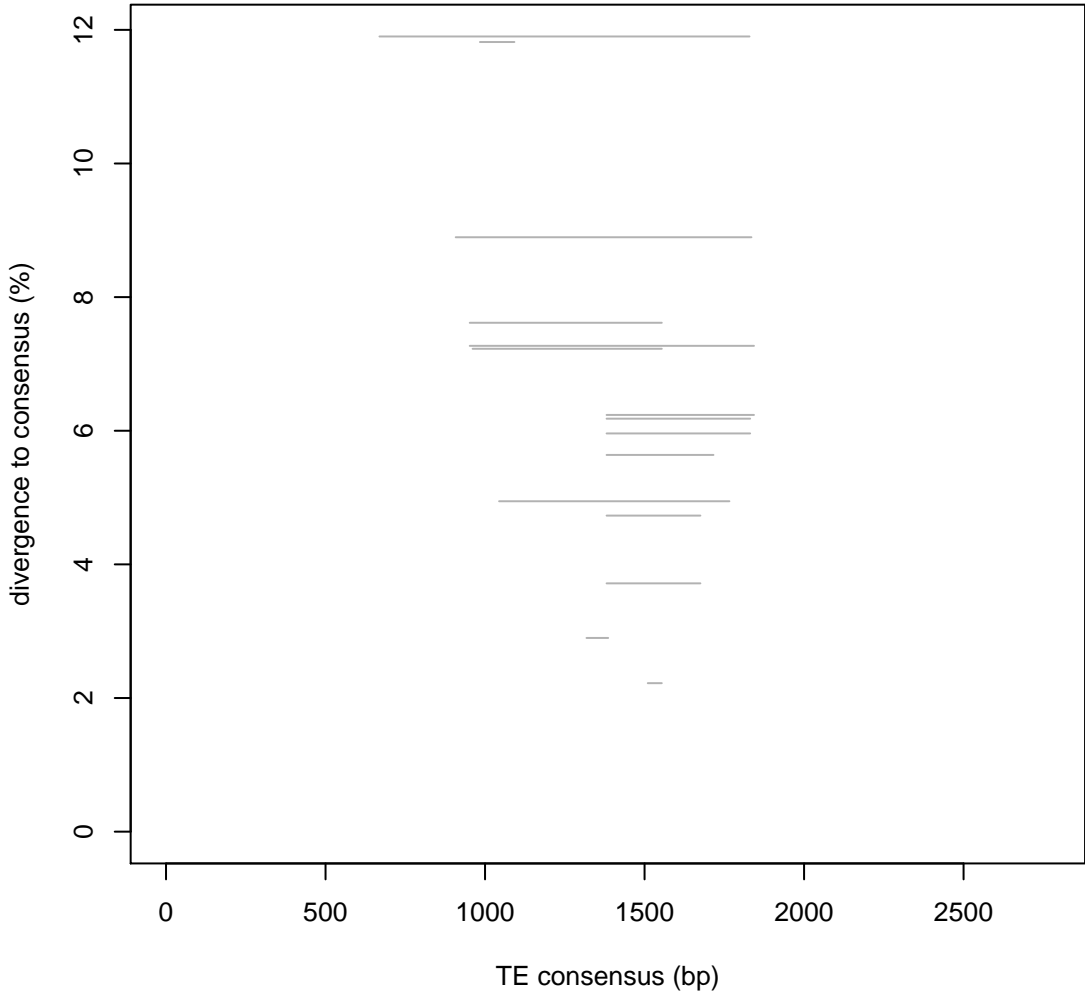
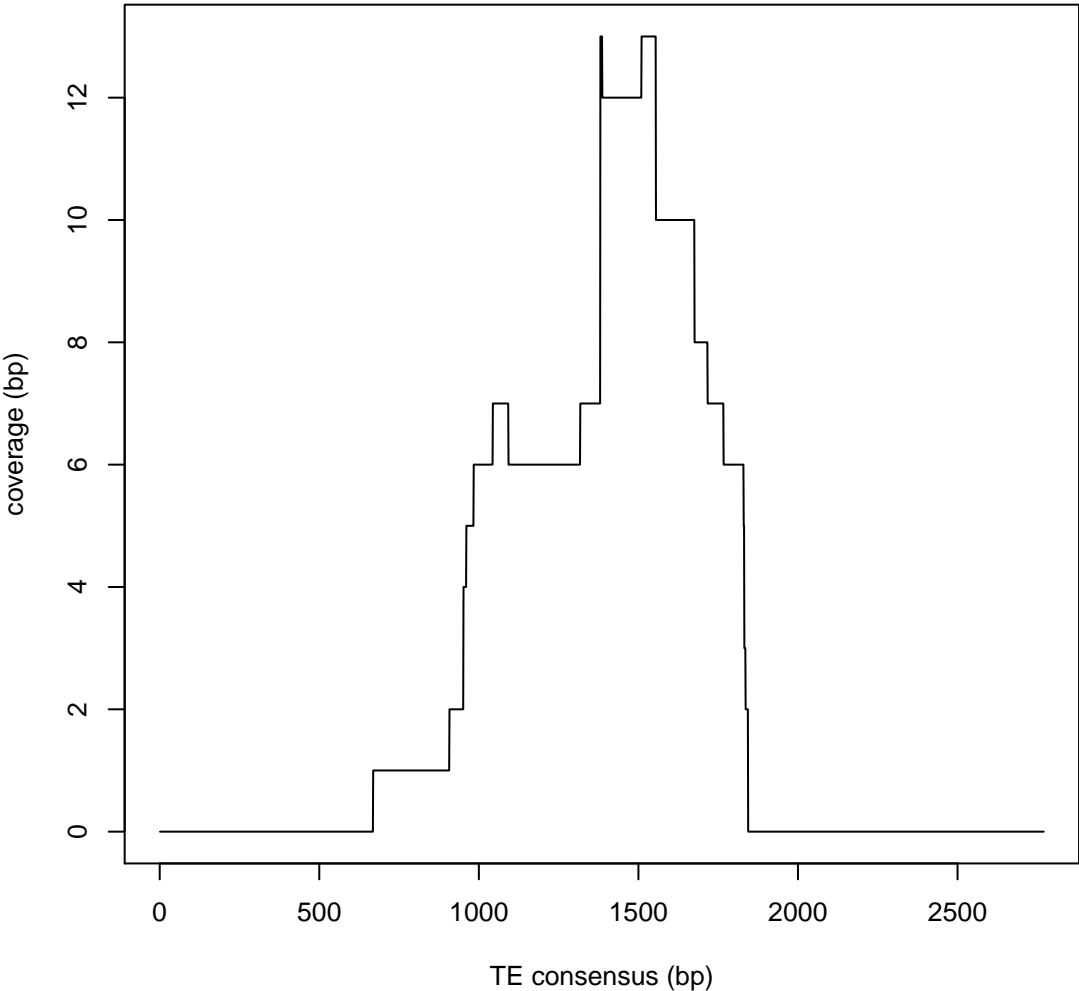


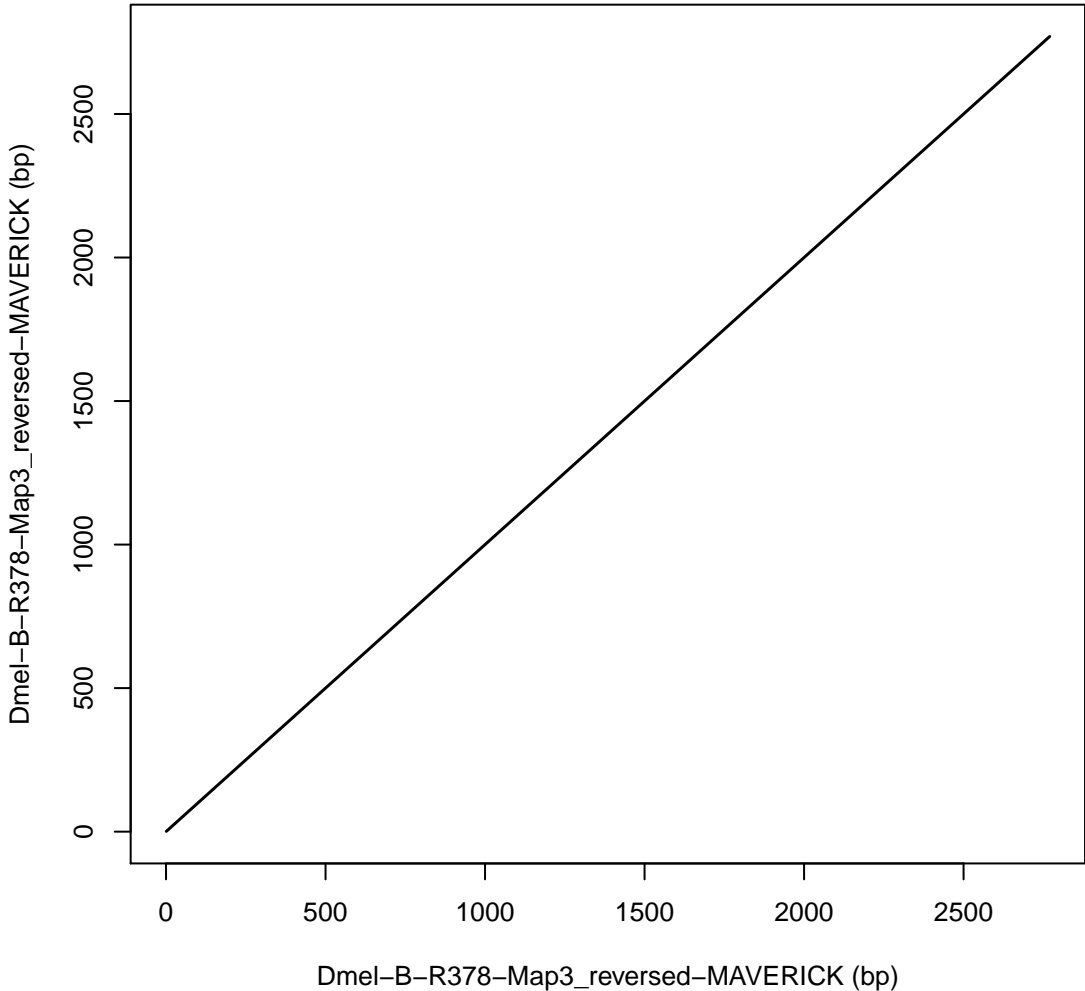
TE: Dmel-B-R378-Map3\_reversed-MAVERICK  
consensus size: 2770bp; fragments: 15; full length: 0 (>=2493bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

