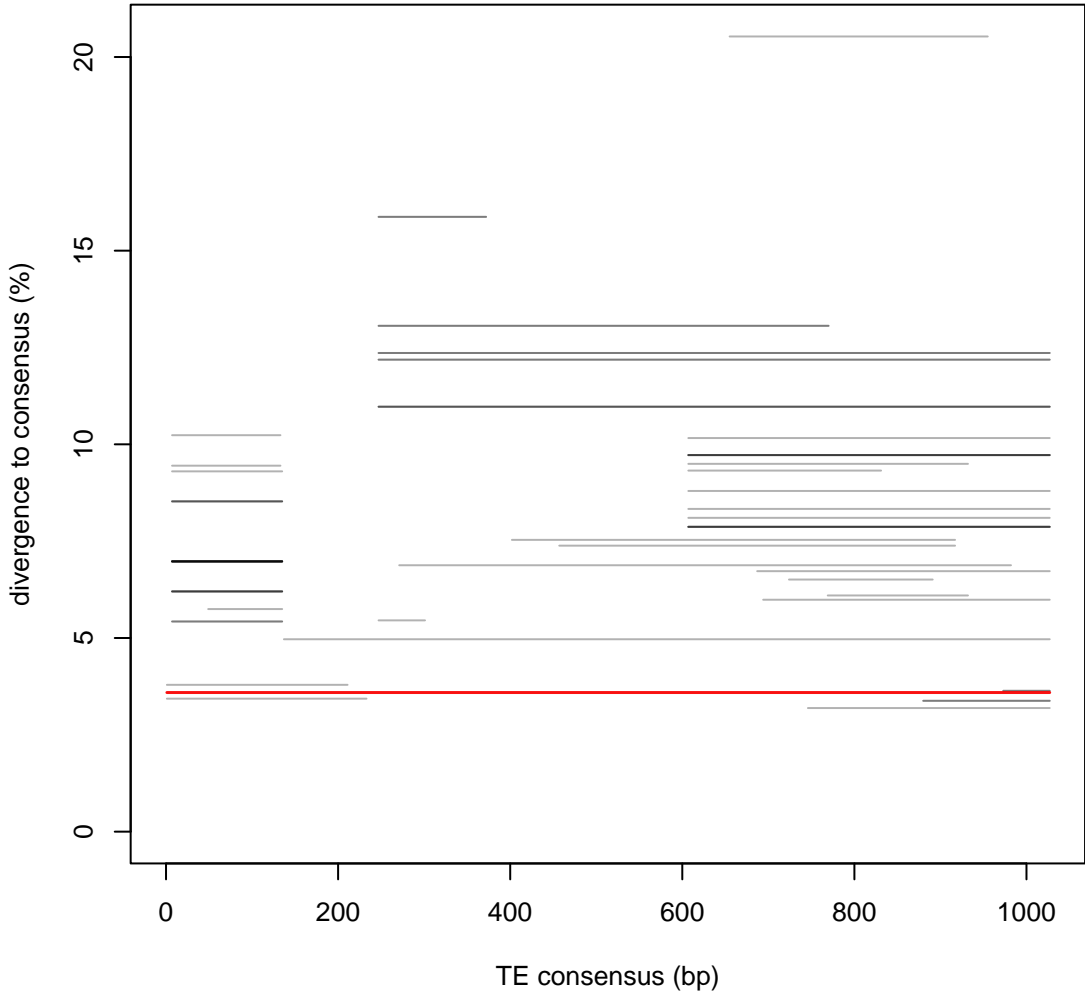
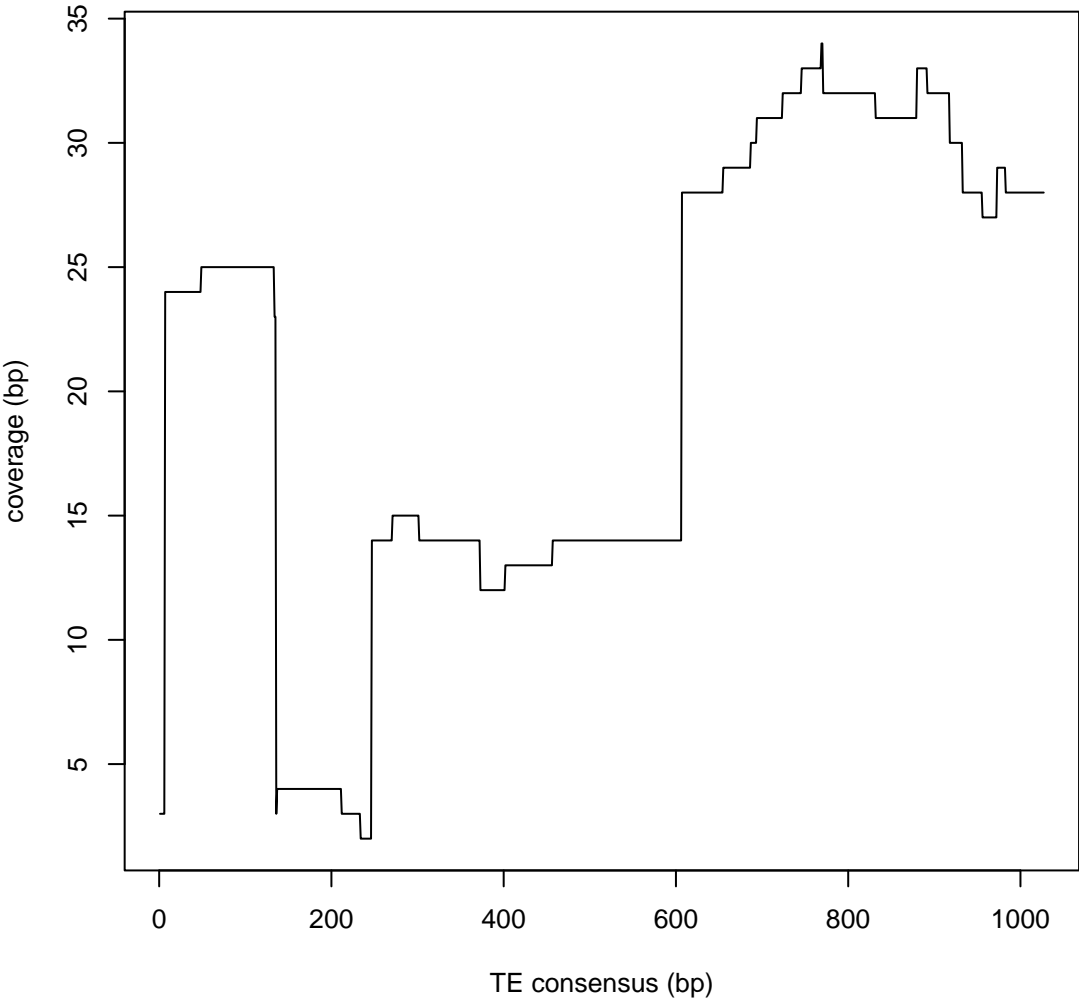


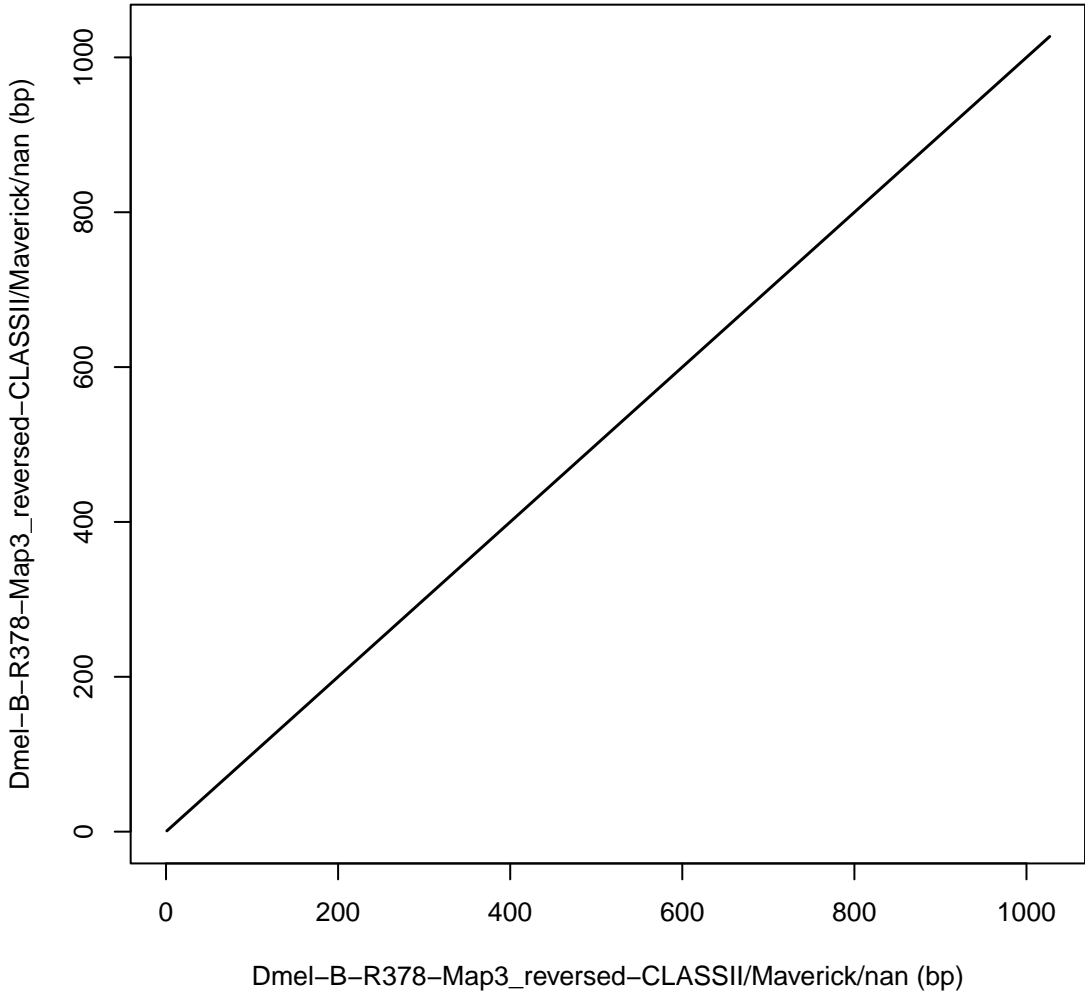
TE: Dmel-B-R378-Map3_reversed-CLASSII/Maverick/nan
consensus size: 1027bp; fragments: 65; full length: 1 (>=924.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

