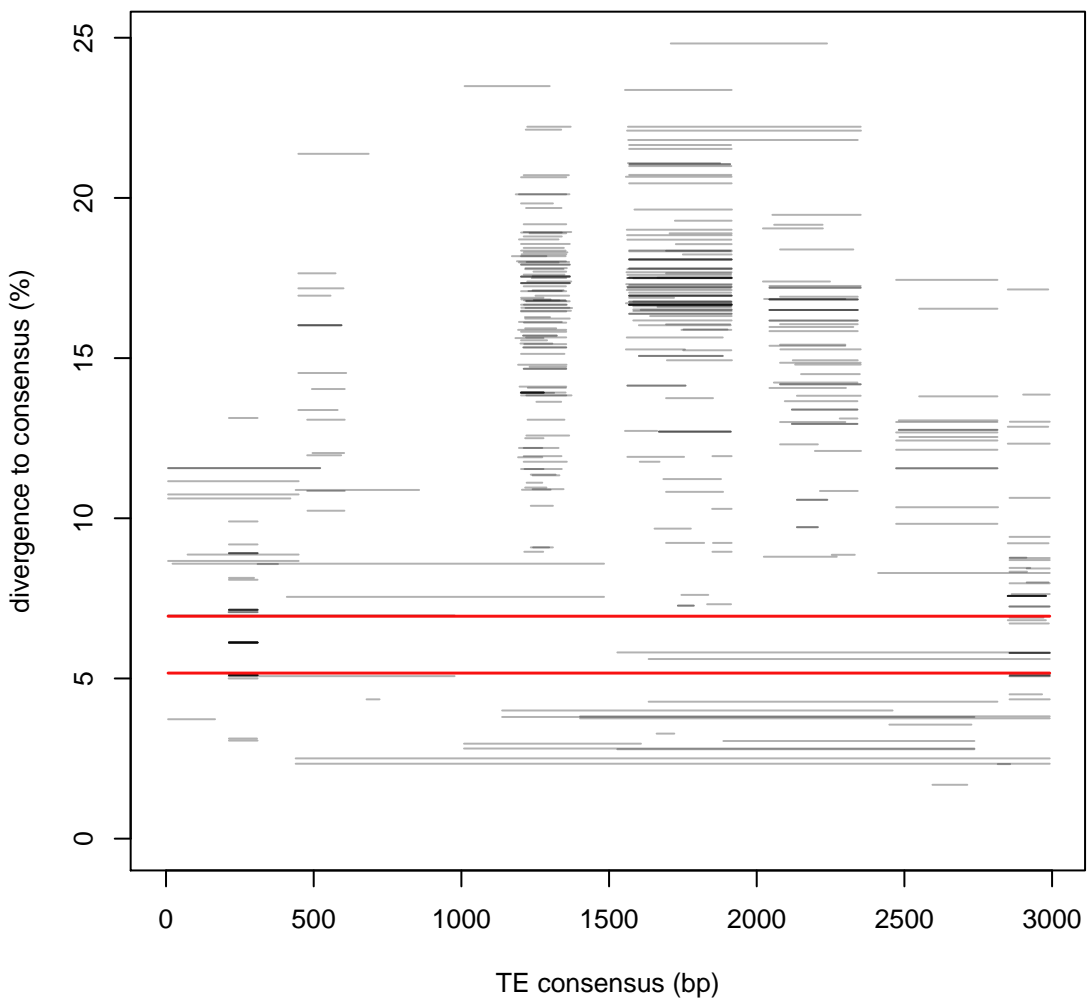
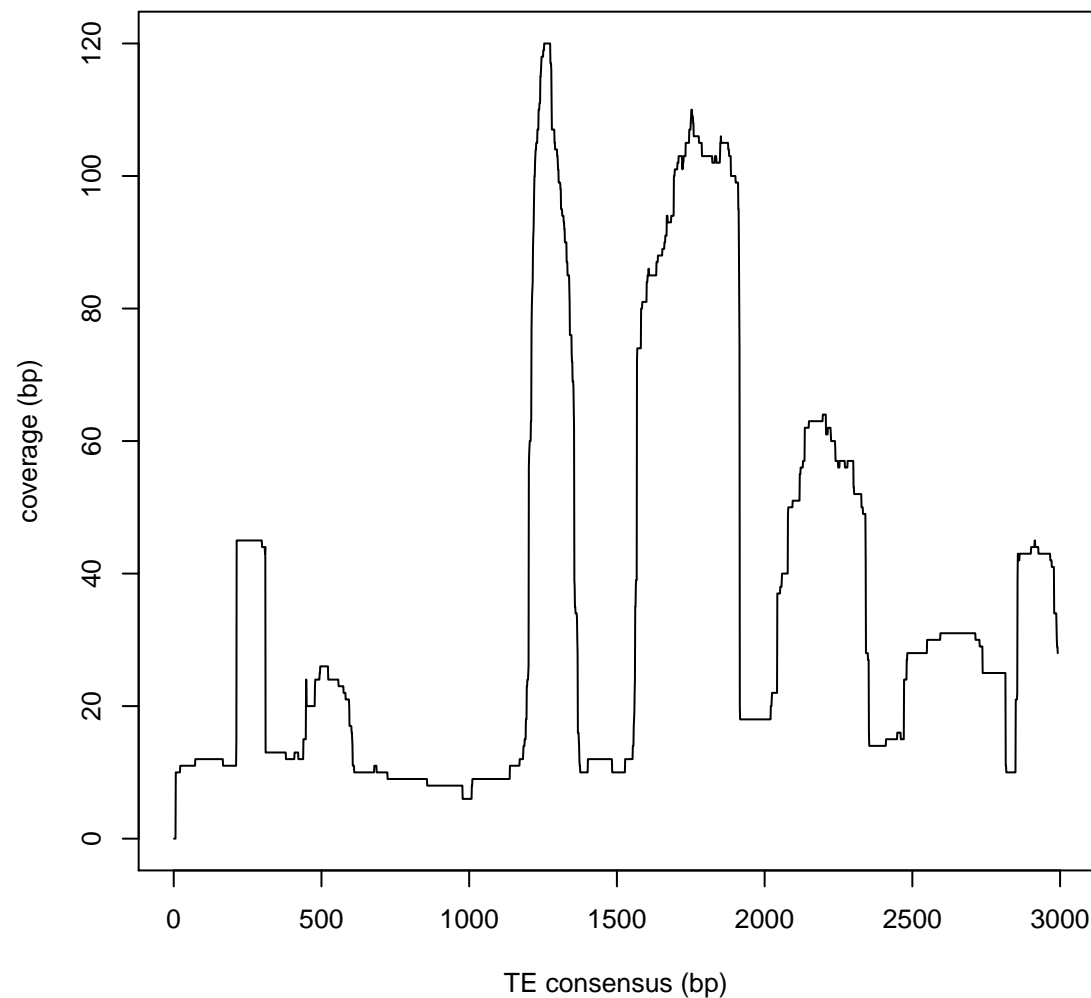


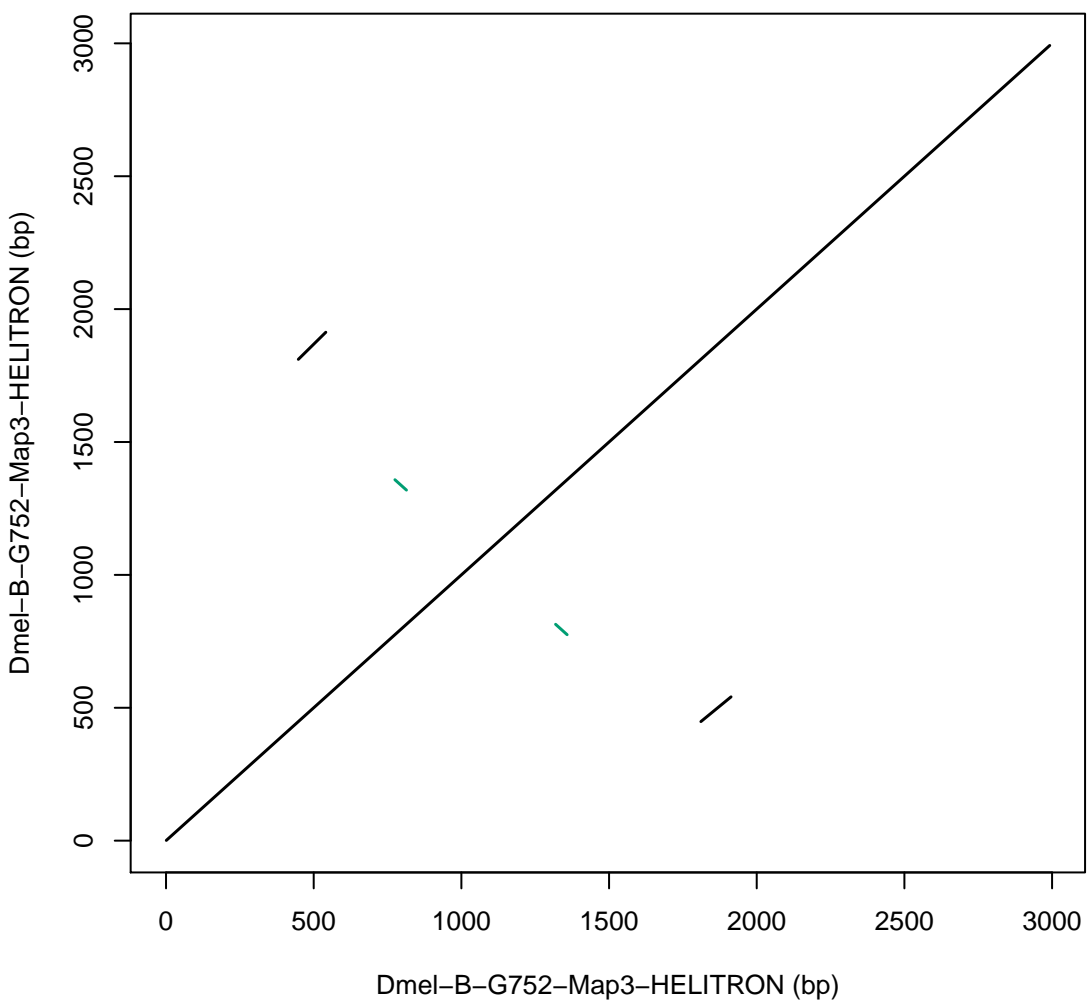
TE: Dmel-B-G752-Map3-HELITRON
consensus size: 2992bp; fragments: 399; full length: 2 (≥ 2692.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

