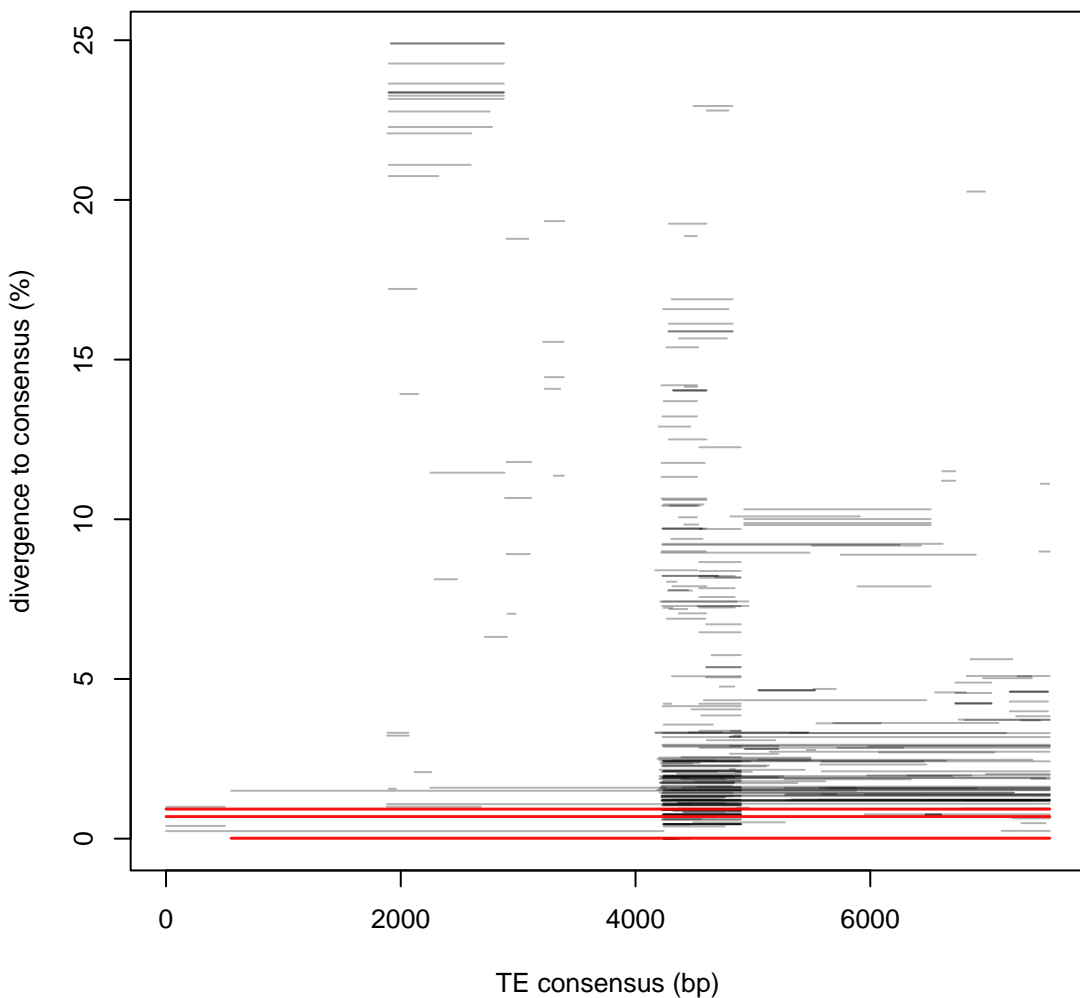
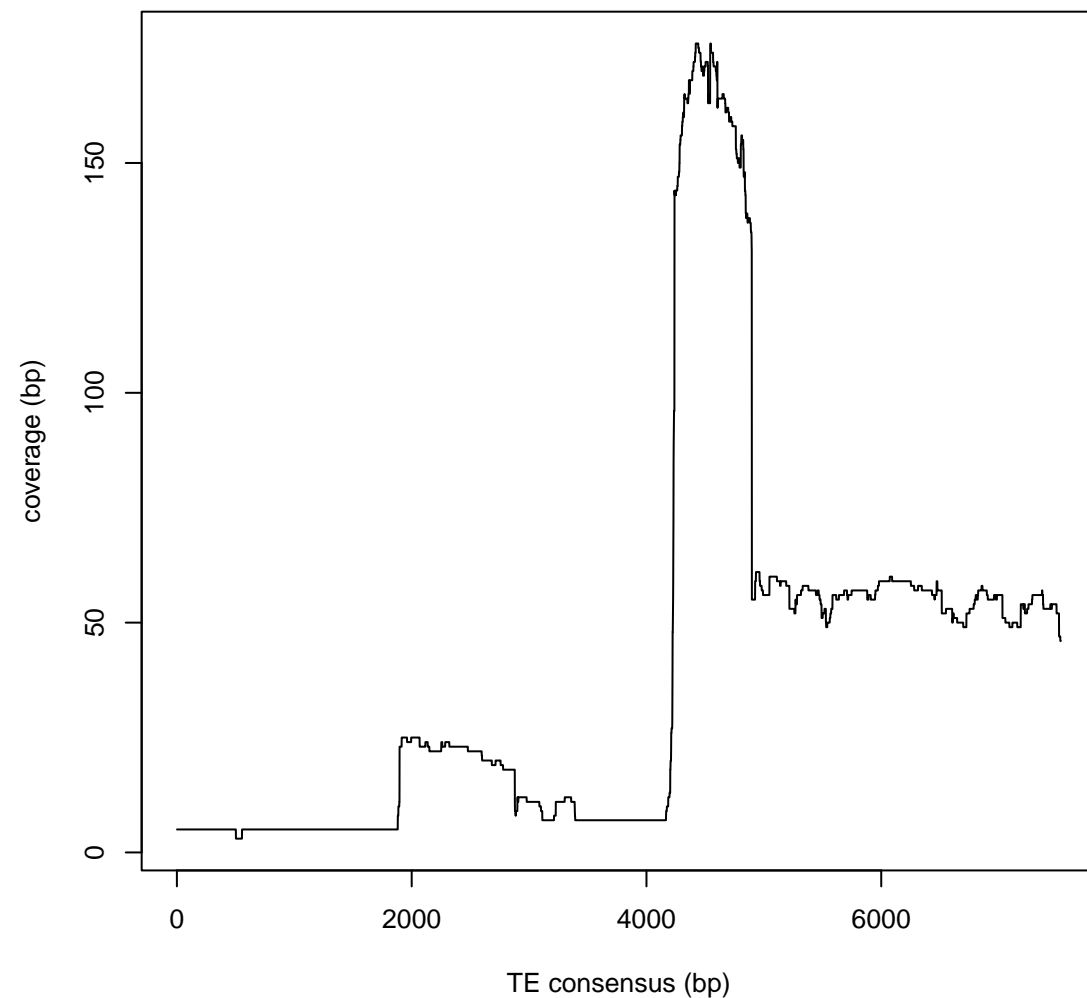


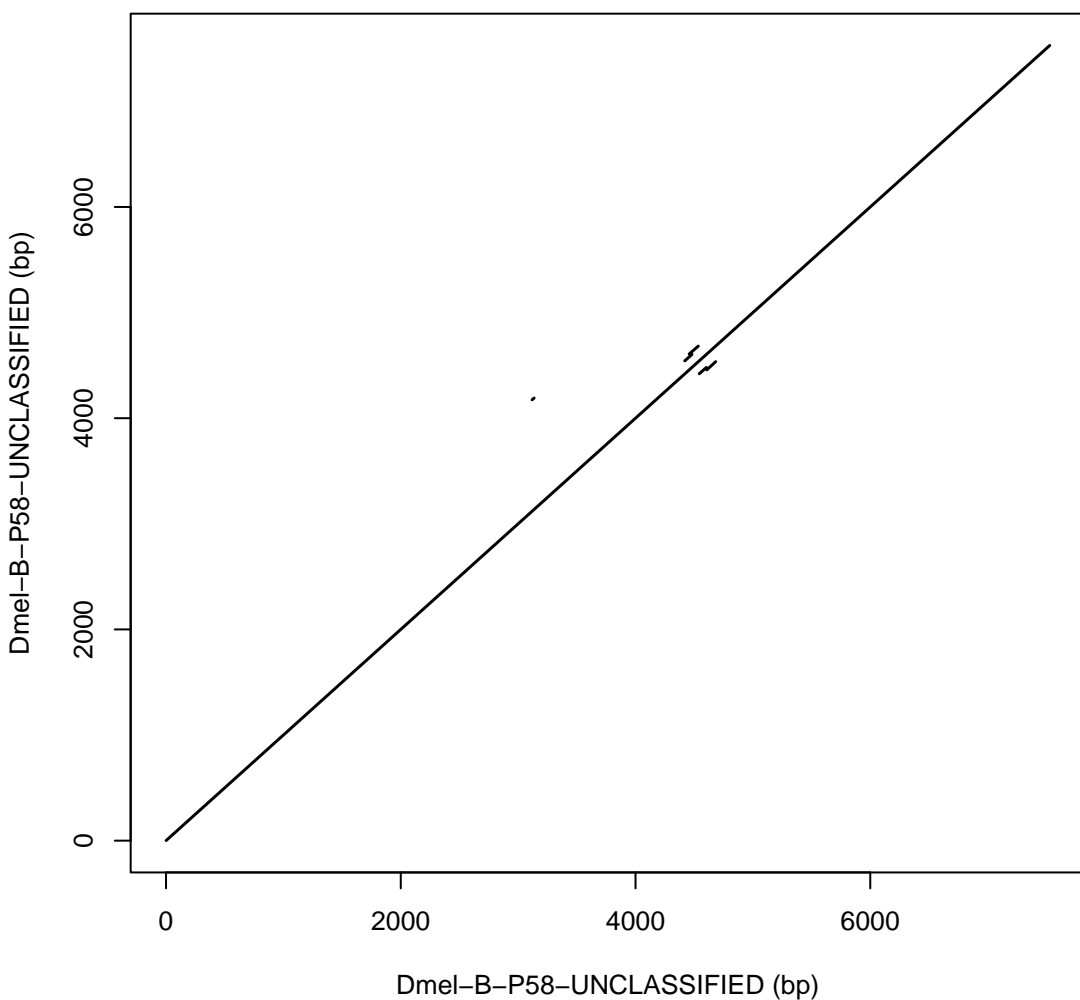
TE: Dmel-B-P58-UNCLASSIFIED
consensus size: 7529bp; fragments: 340; full length: 3 (>=6776.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

