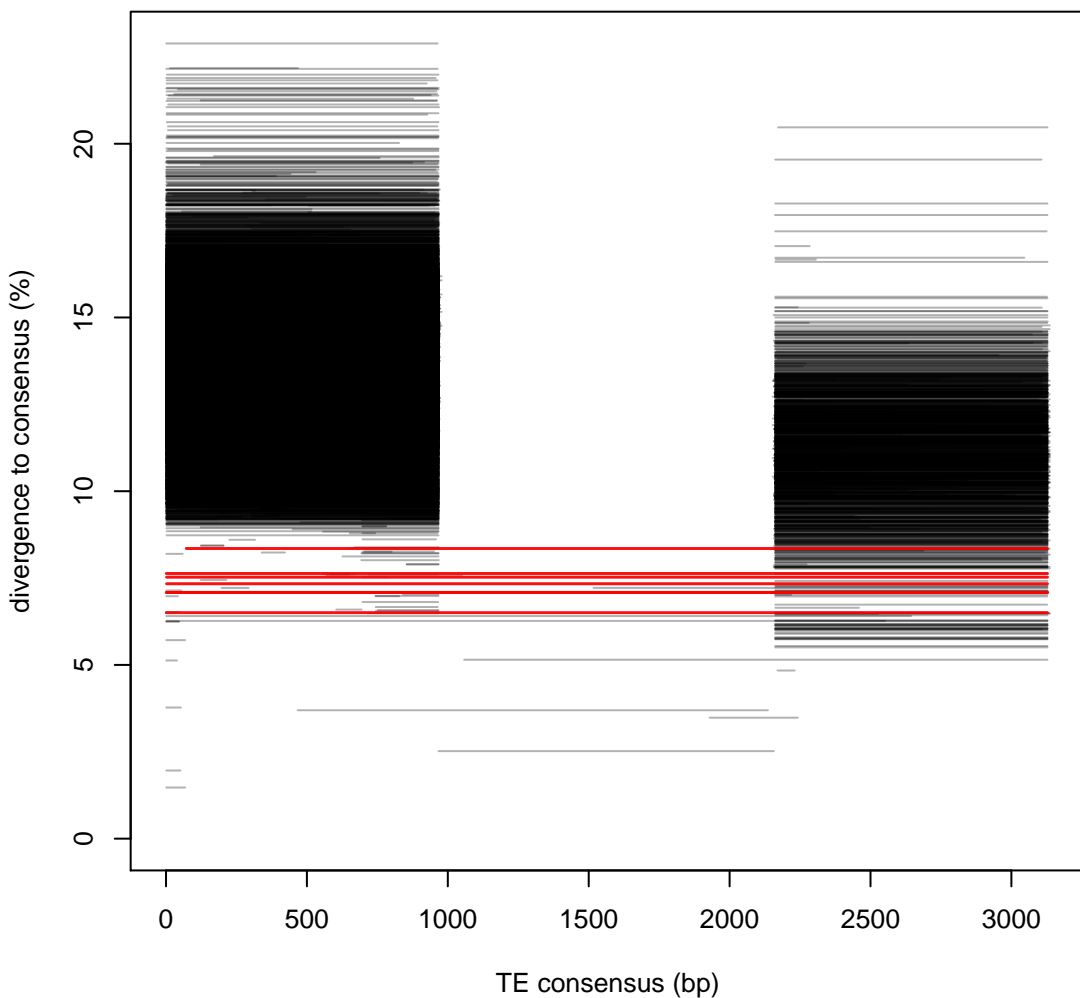
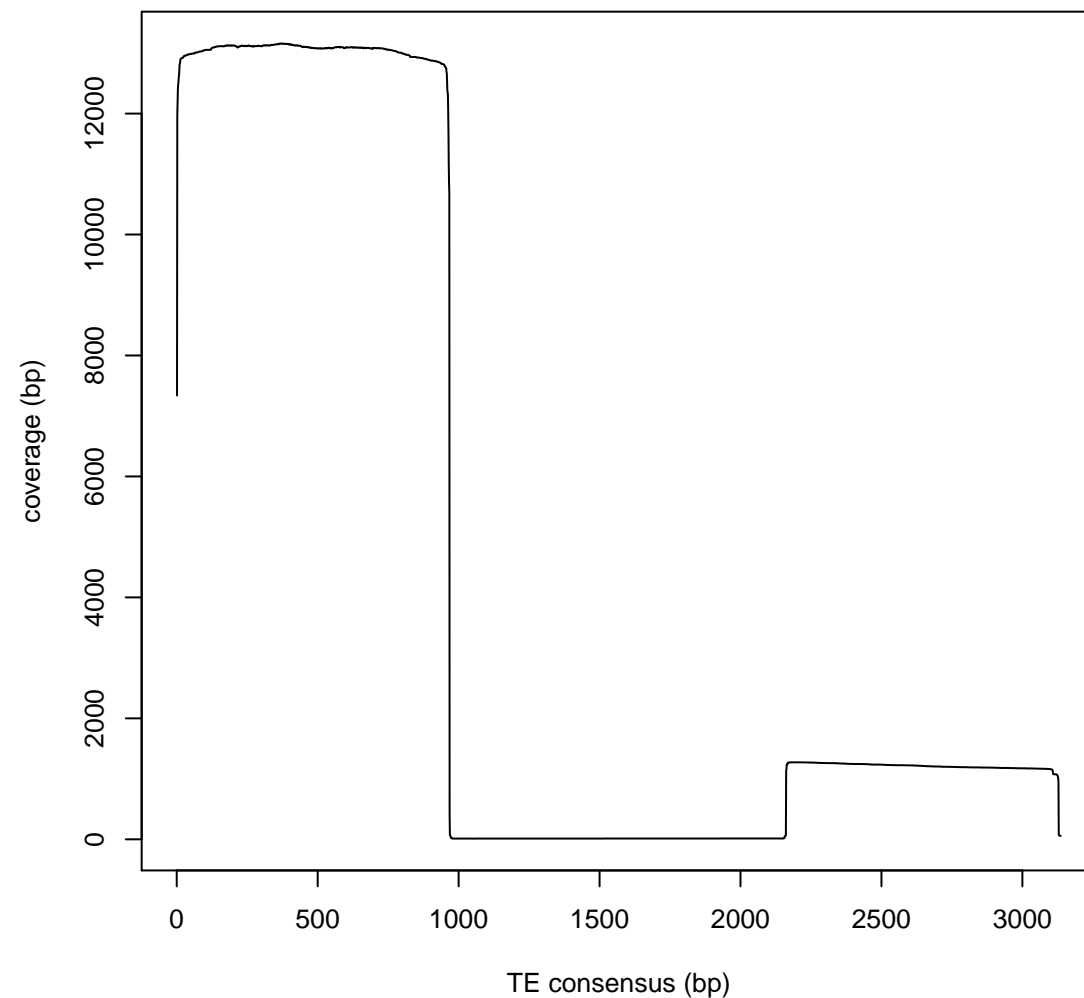


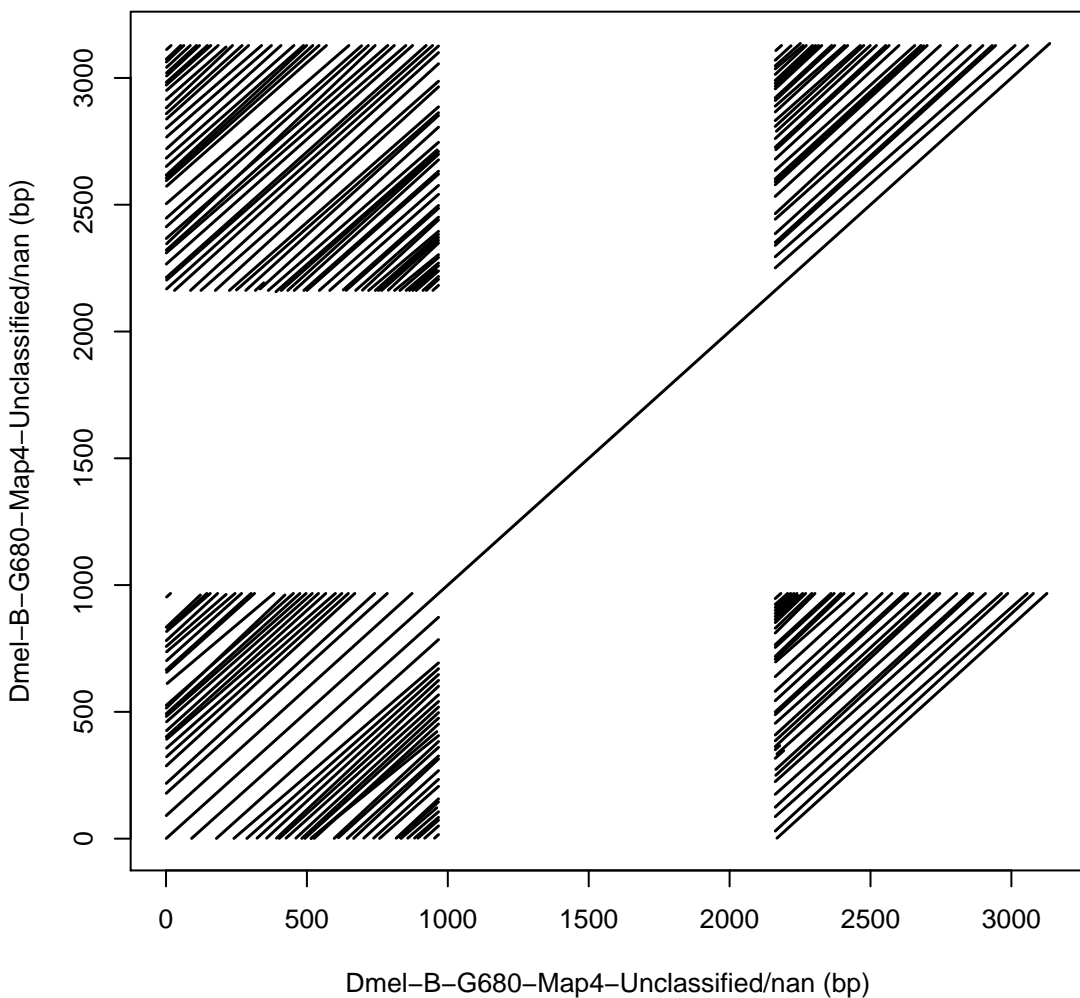
TE: Dmel-B-G680-Map4-Unclassified/nan
consensus size: 3136bp; fragments: 14949; full length: 7 (>=2822.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

