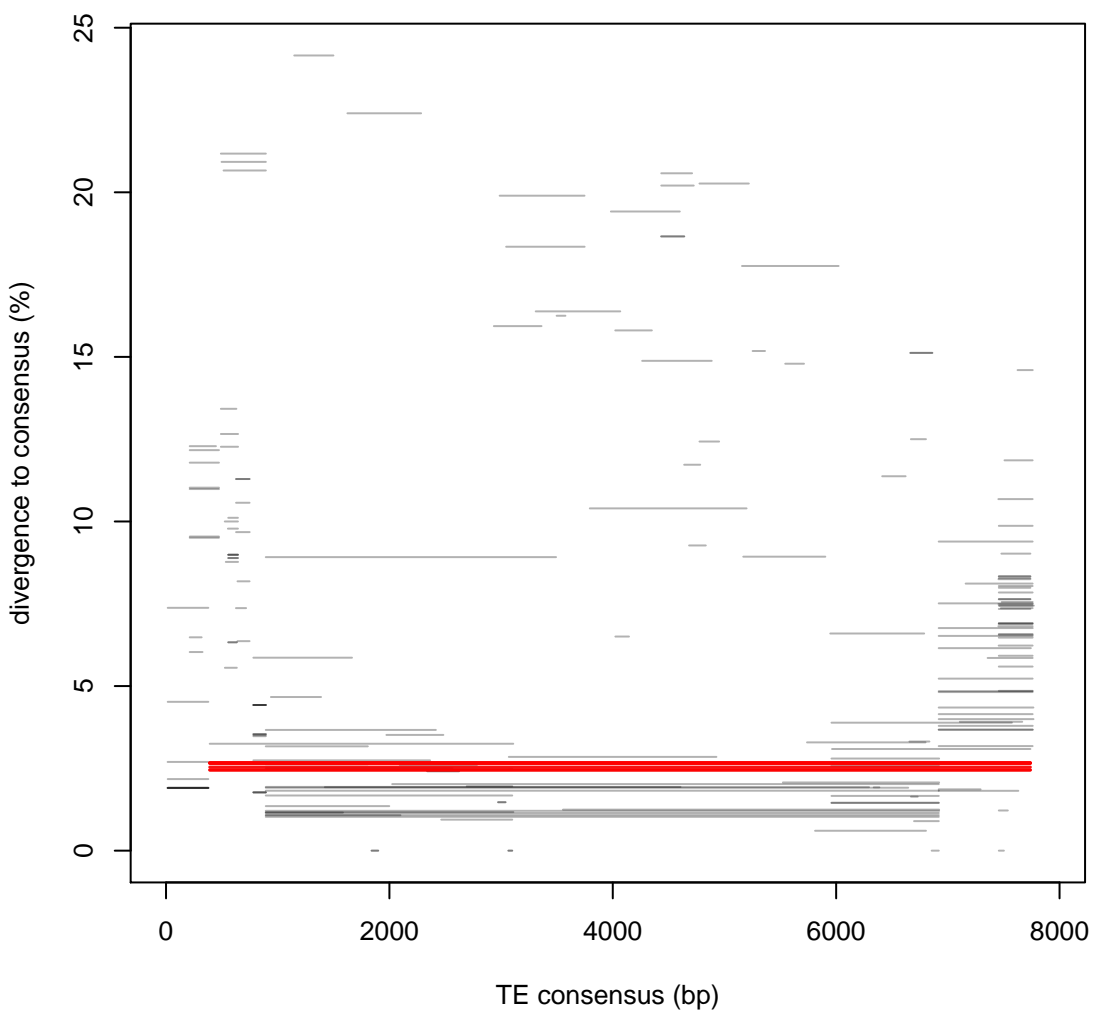
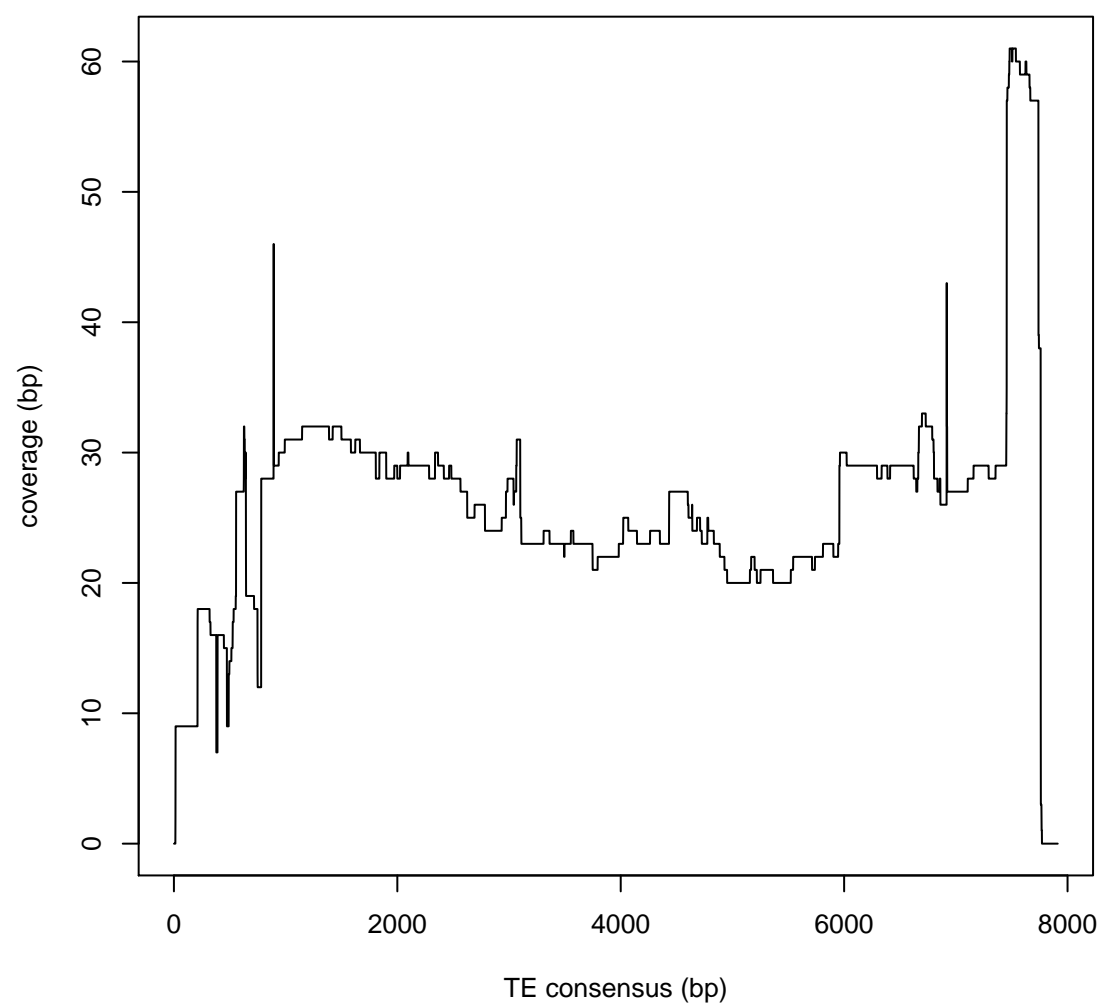


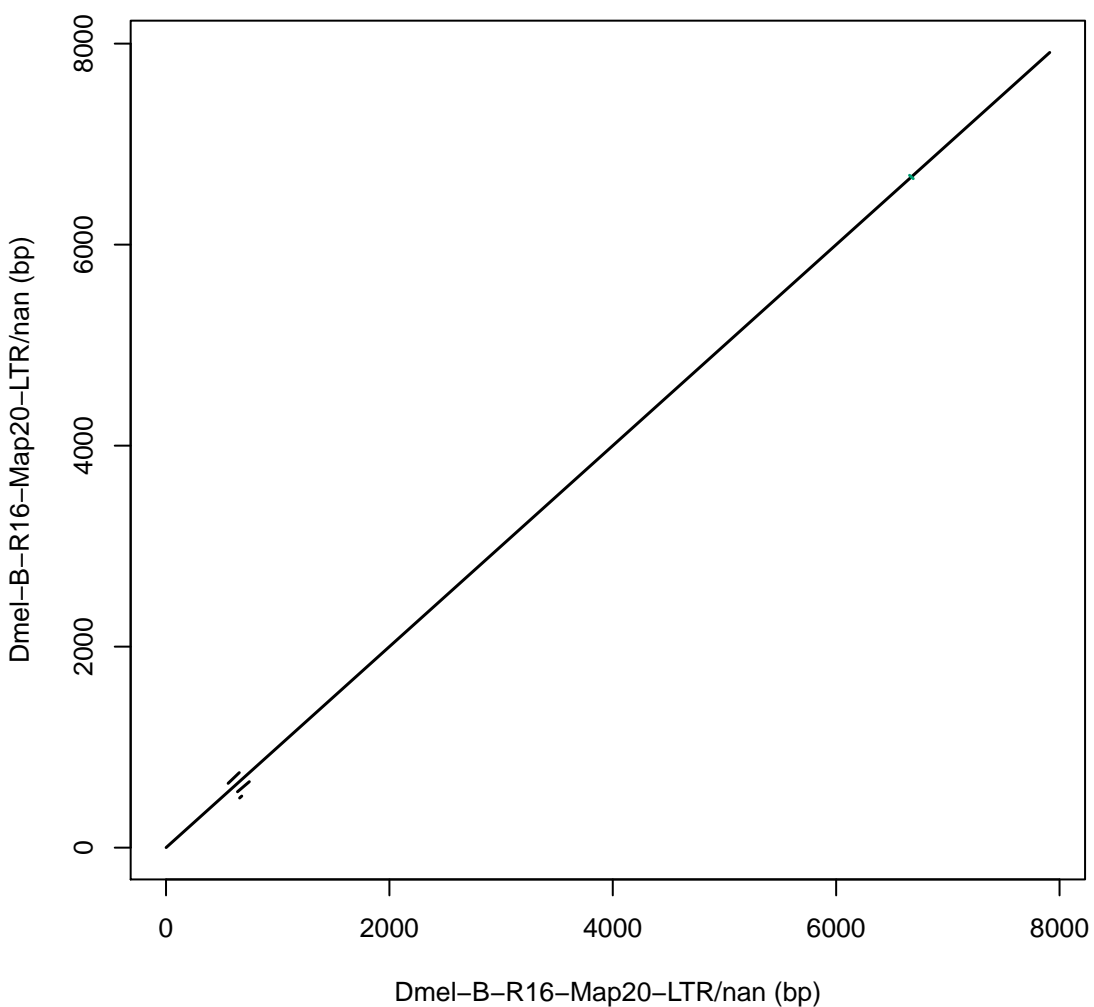
TE: Dmel-B-R16-Map20-LTR/nan
consensus size: 7912bp; fragments: 203; full length: 8 (>=7120.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

