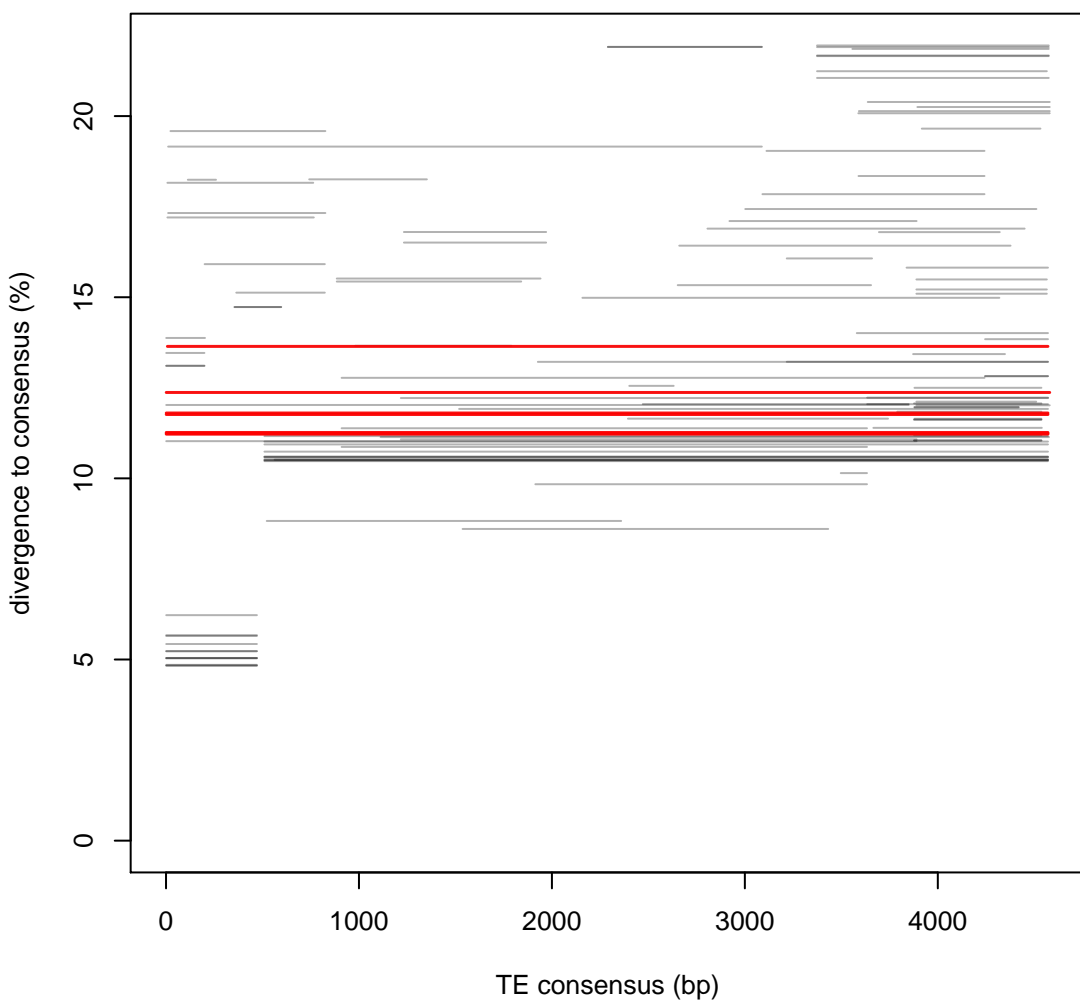
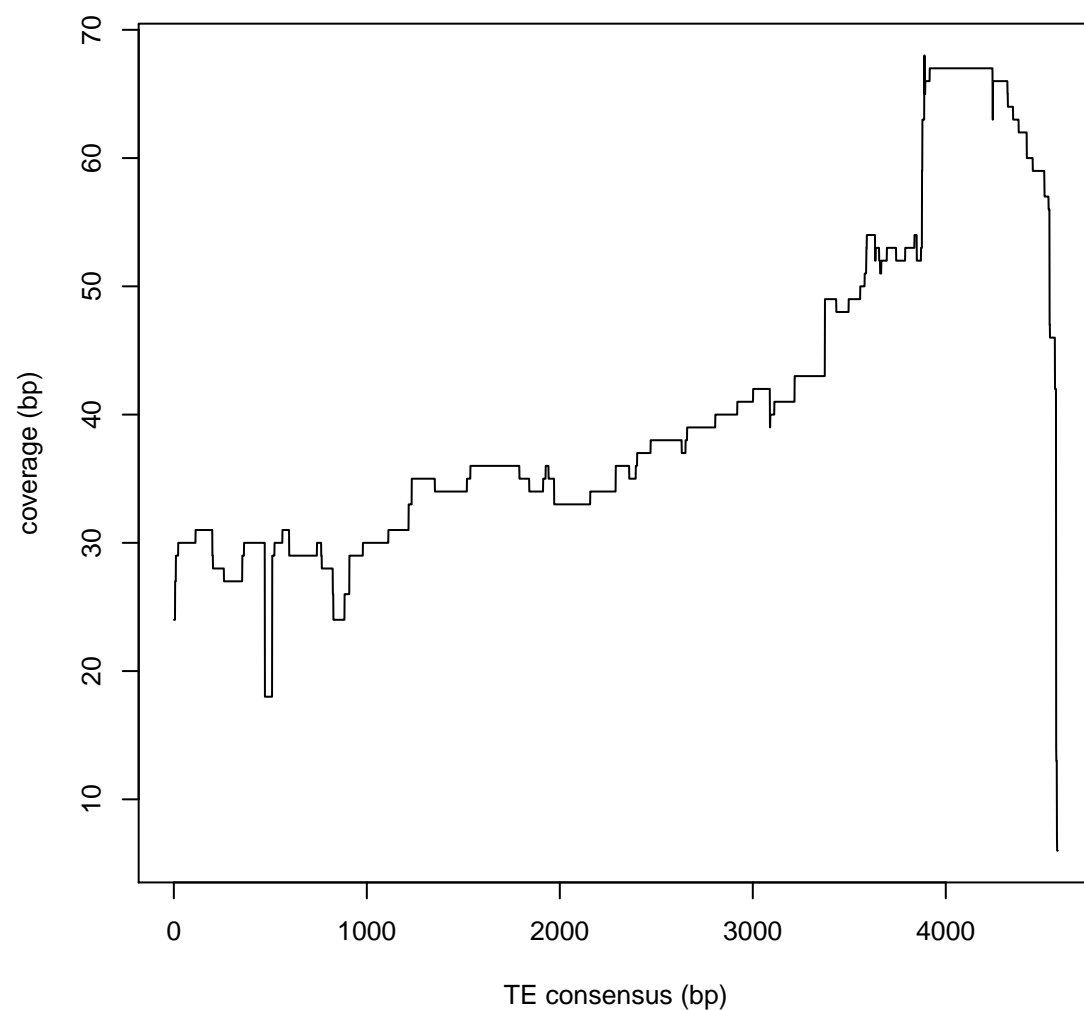


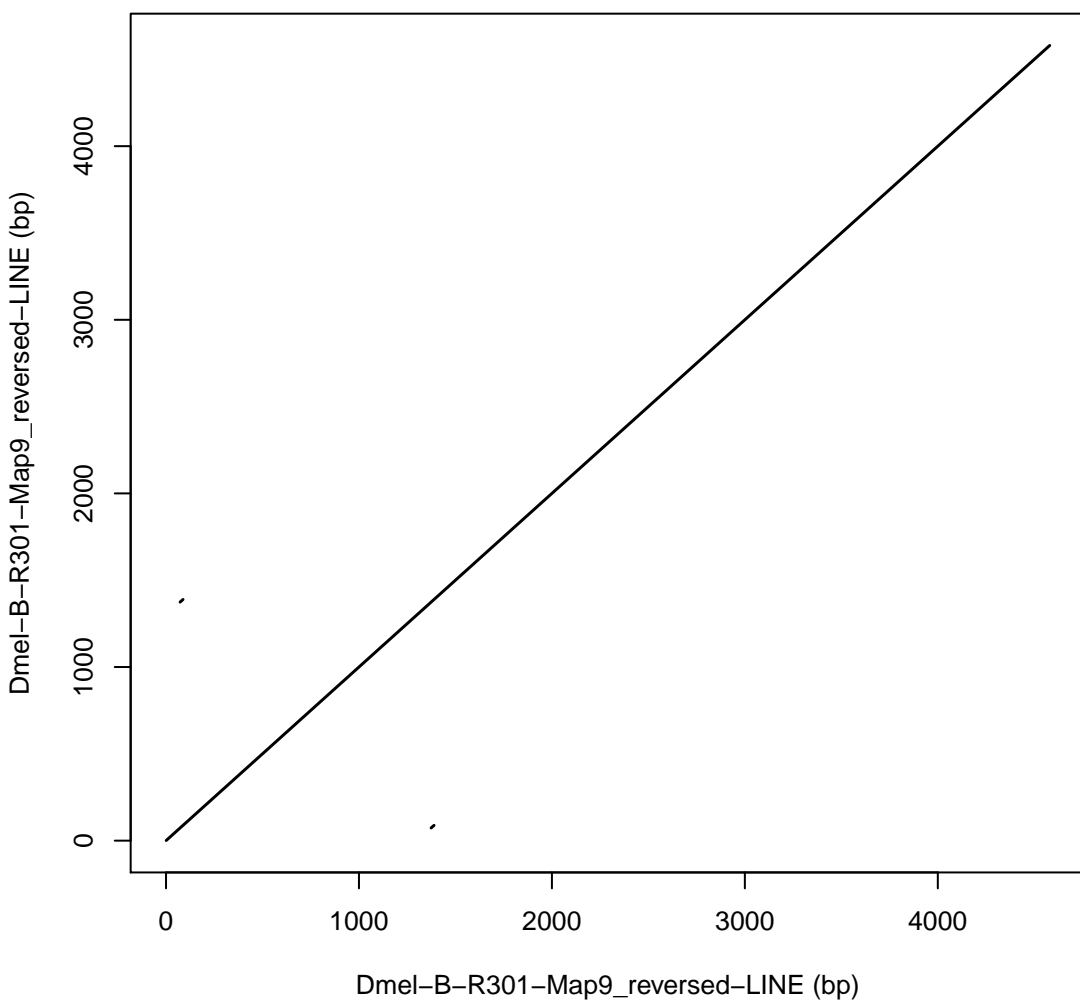
TE: Dmel-B-R301-Map9_reversed-LINE
consensus size: 4580bp; fragments: 119; full length: 7 (>=4122bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

