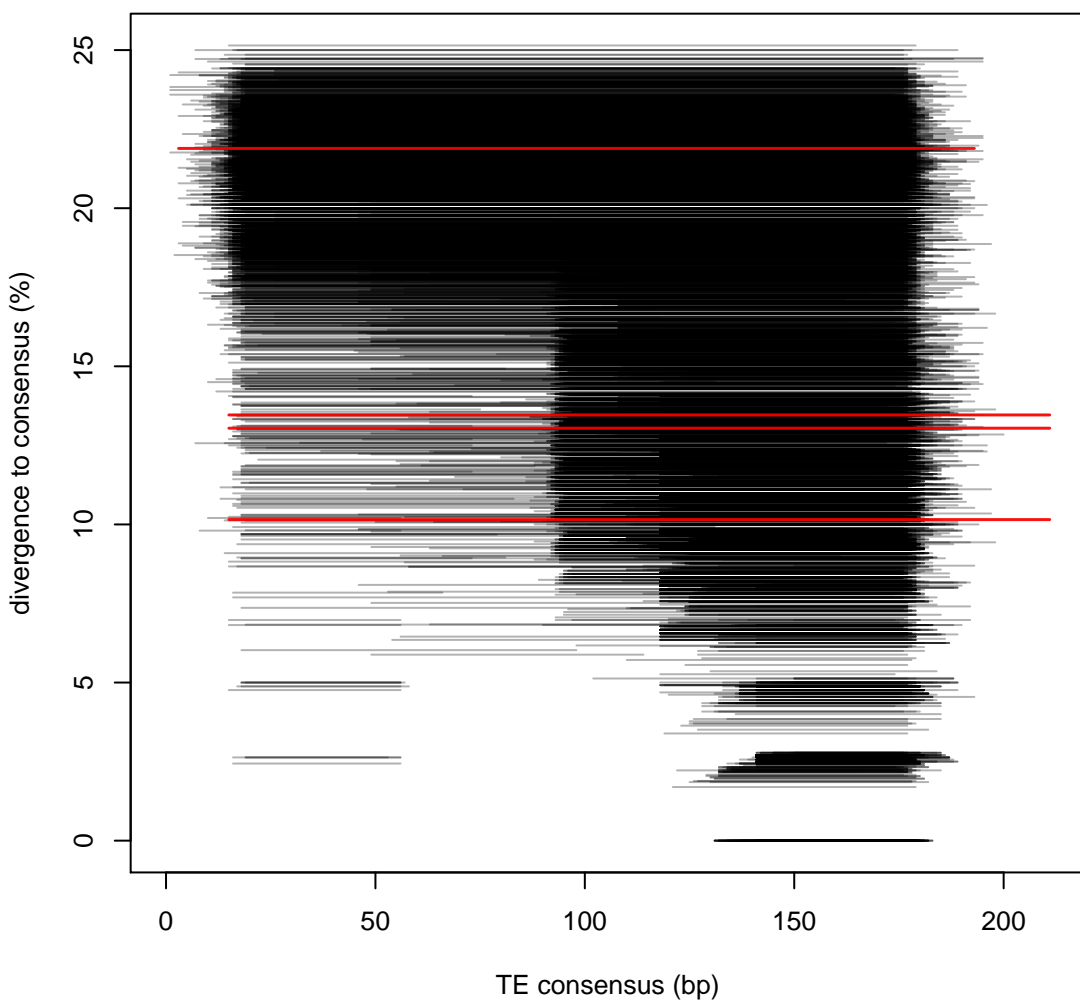
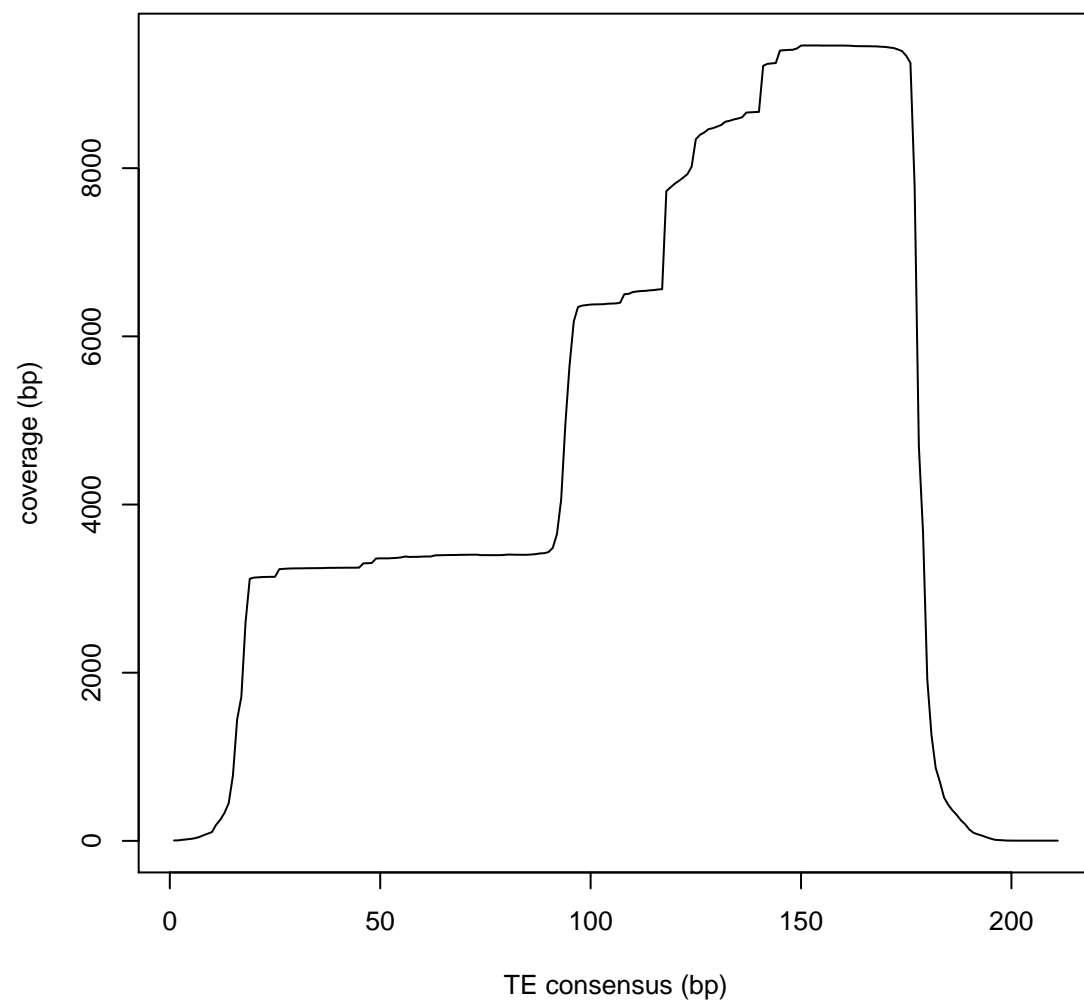


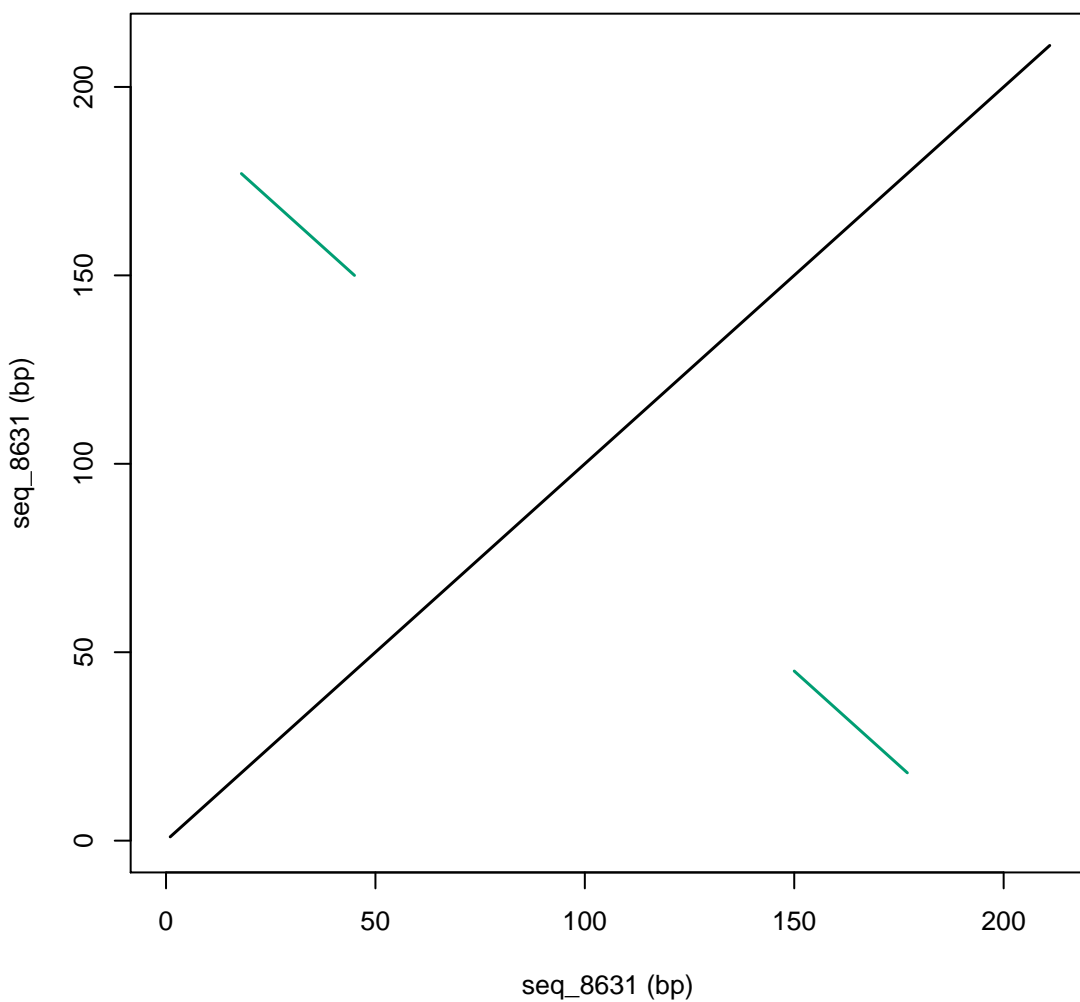
TE: seq_8631
consensus size: 211bp; fragments: 9512; full length: 4 (≥ 189.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

