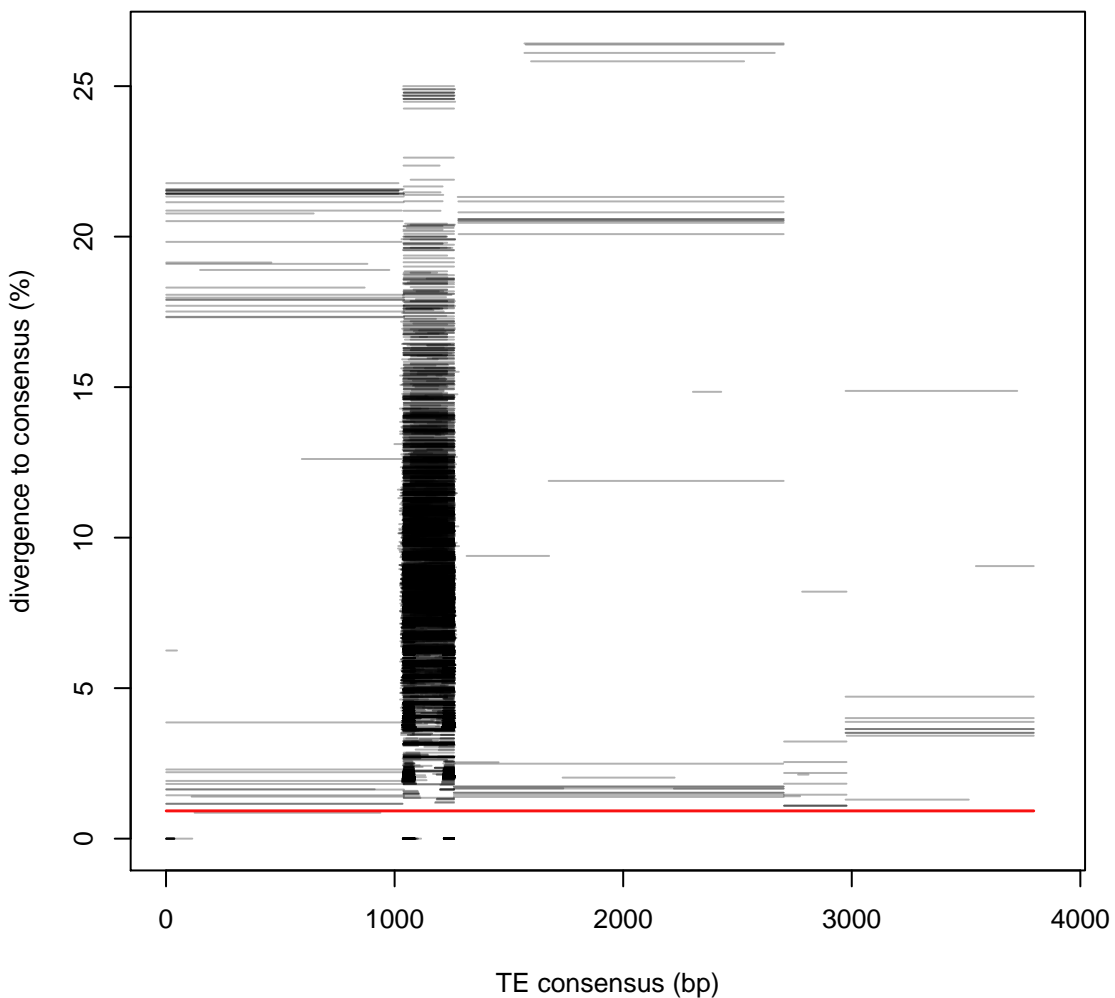
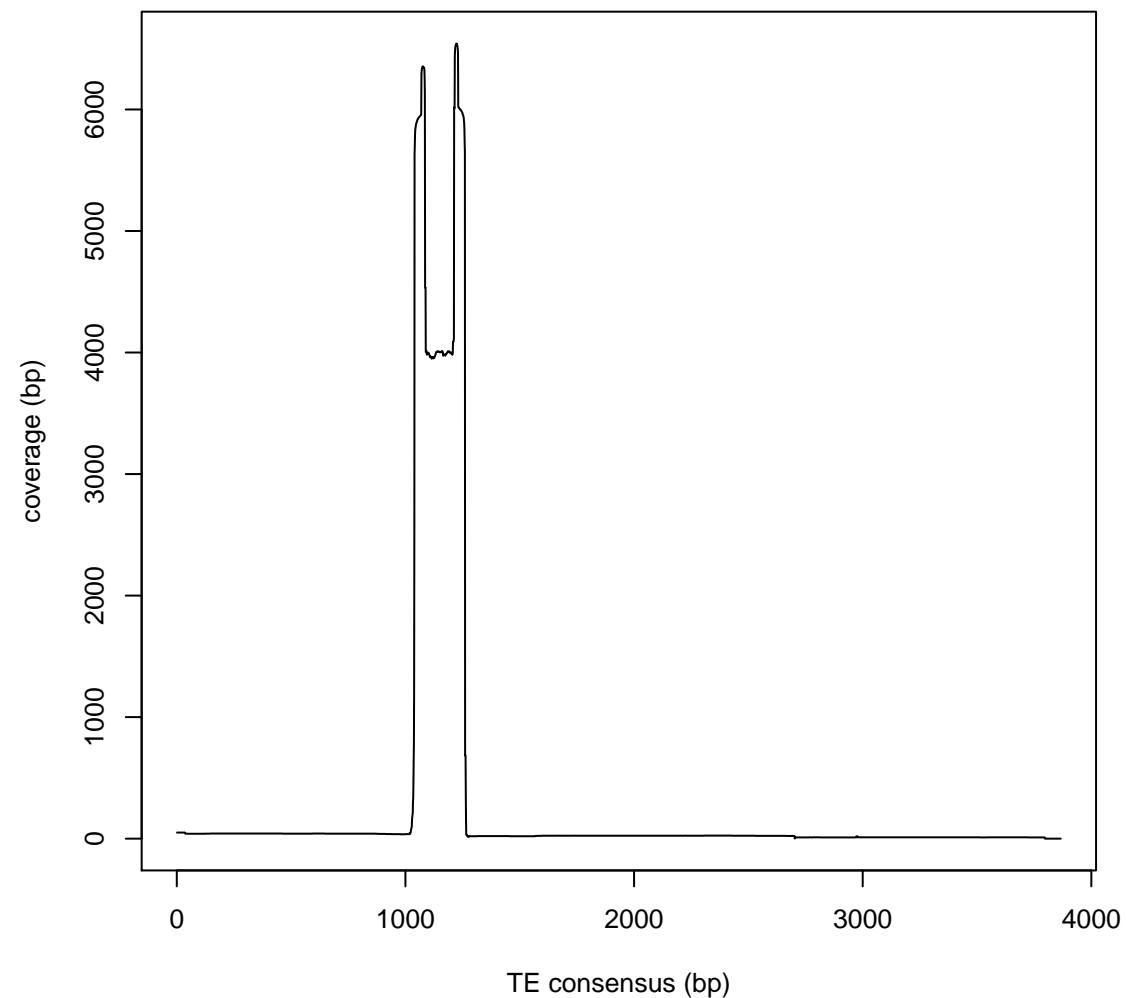


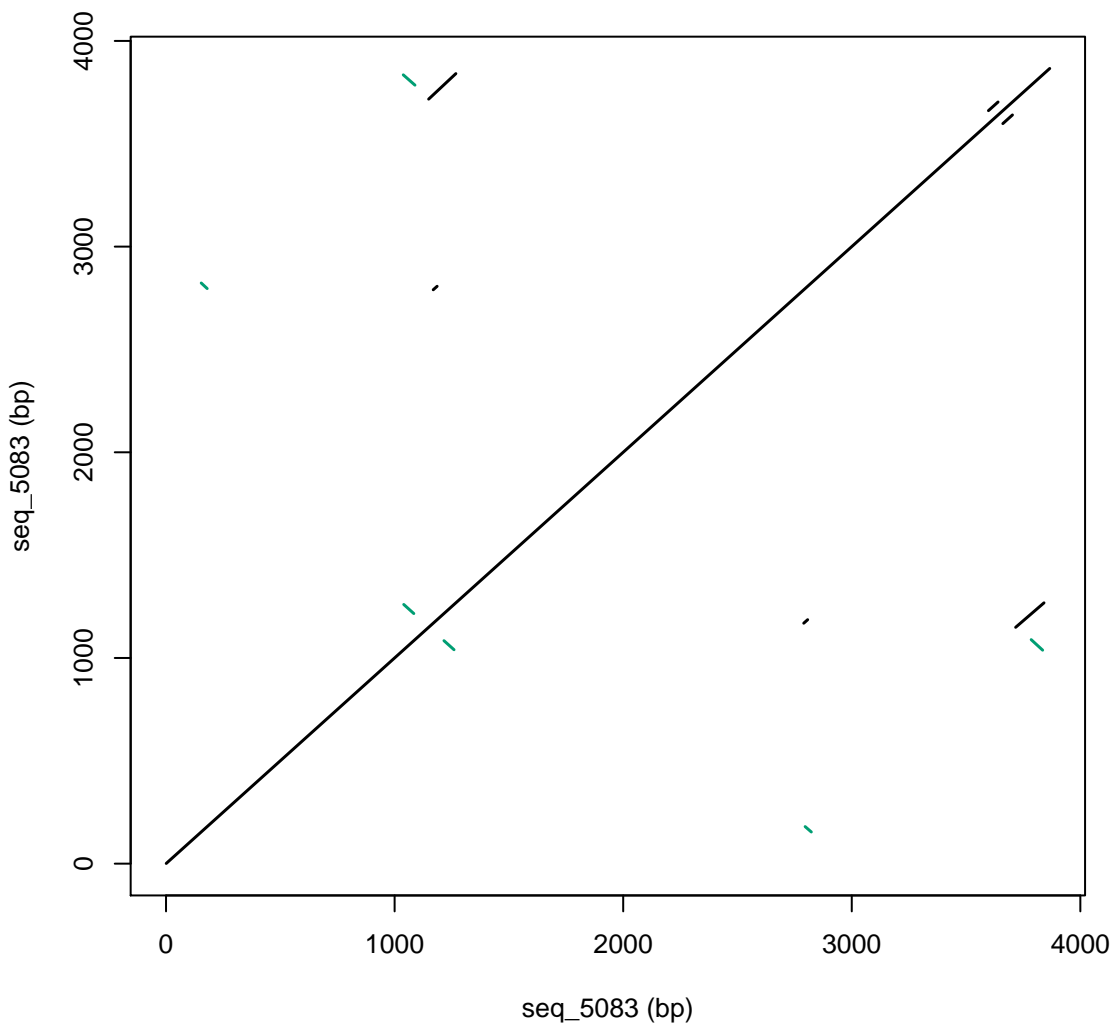
TE: seq_5083
consensus size: 3866bp; fragments: 9528; full length: 1 (≥ 3479.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

