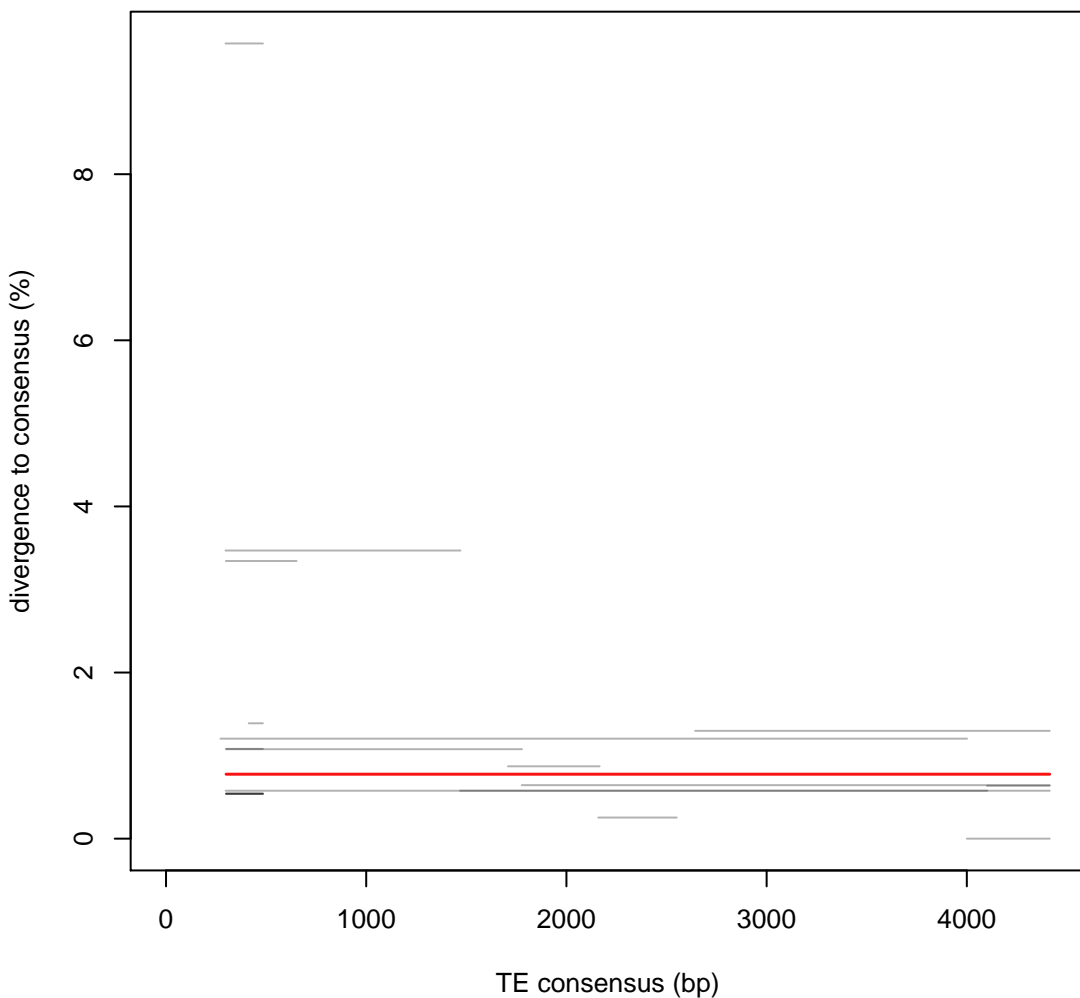
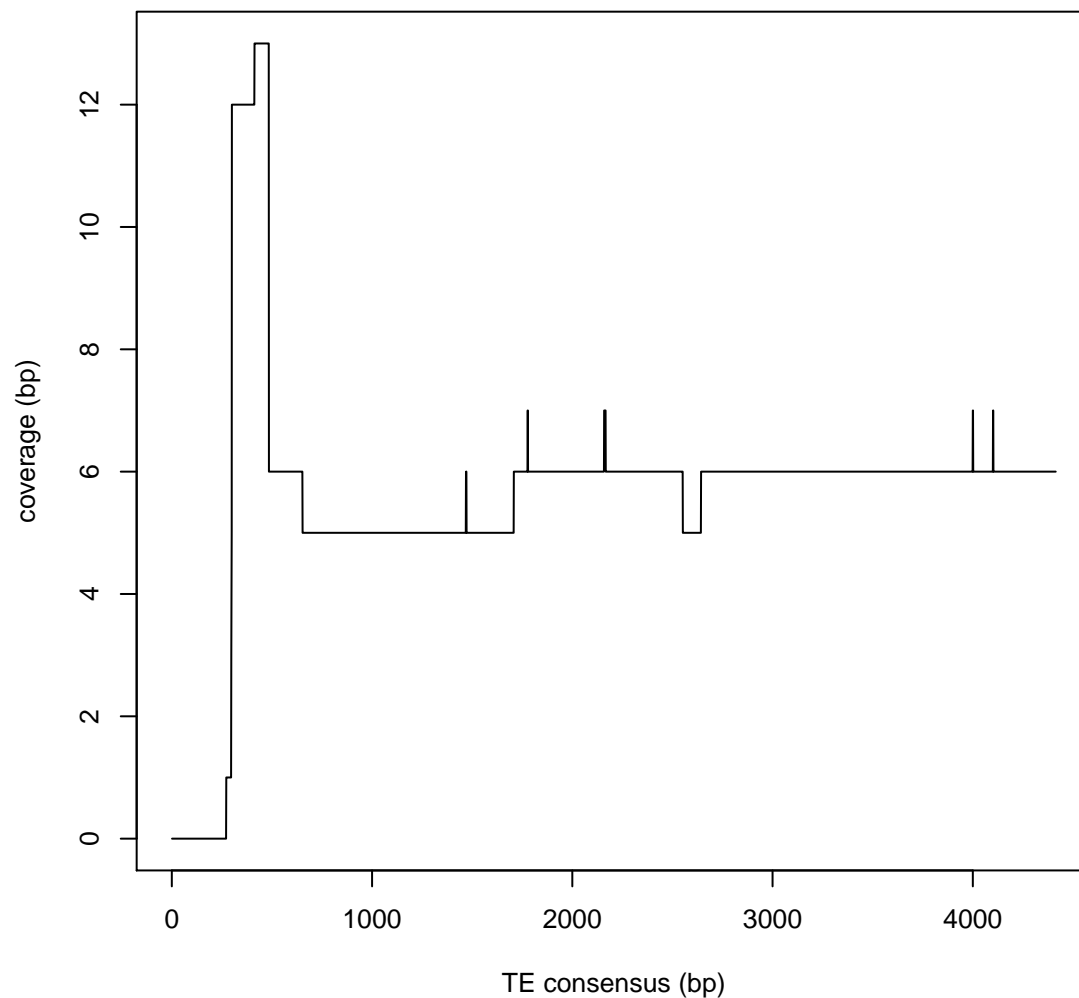


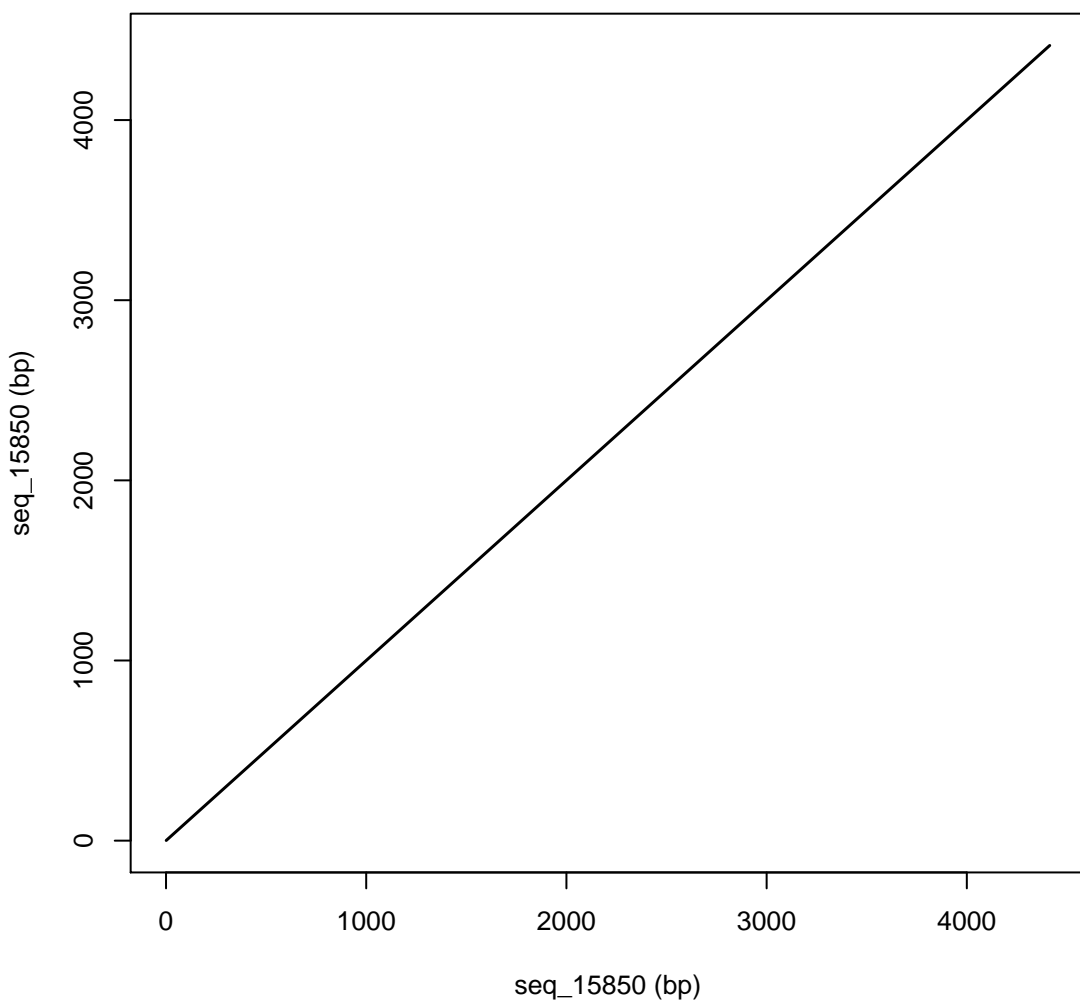
TE: seq_15850
consensus size: 4414bp; fragments: 20; full length: 1 (≥ 3972.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

