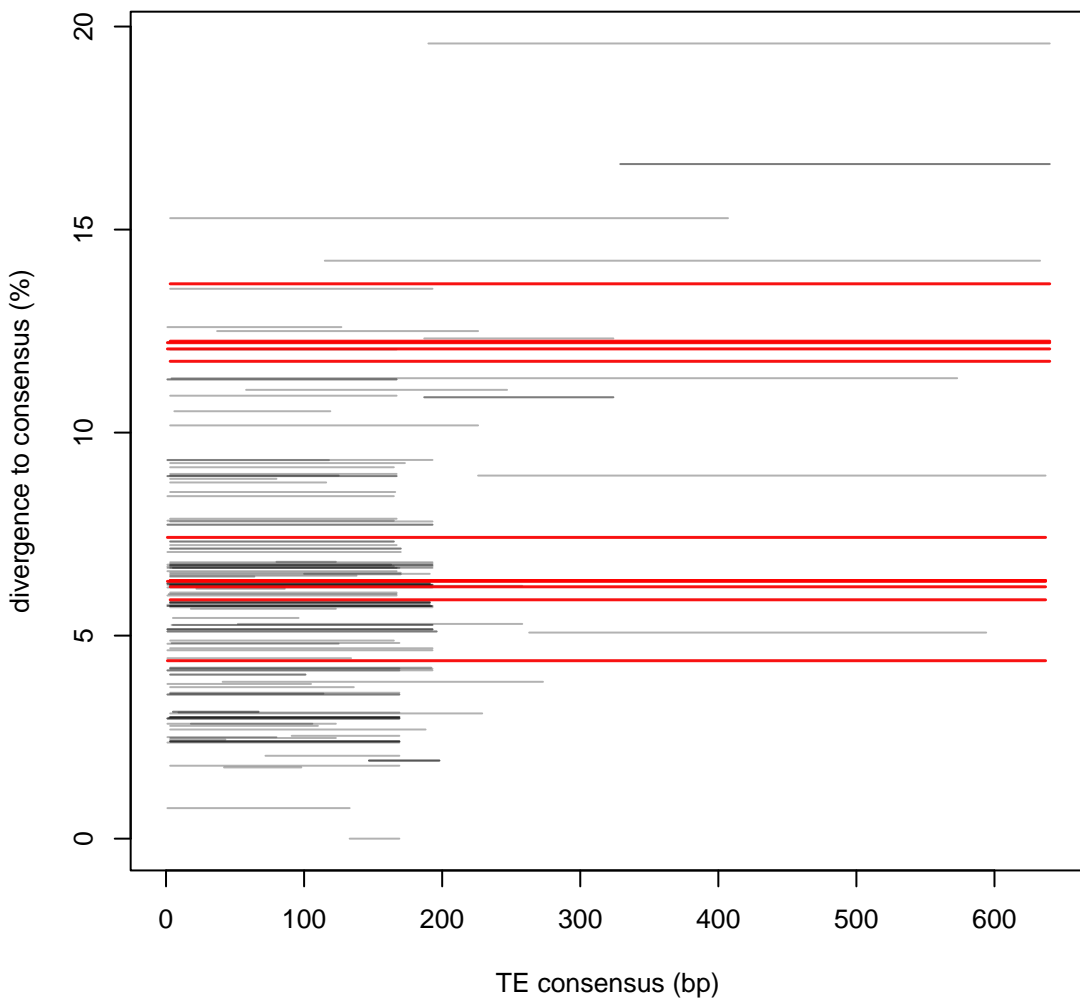
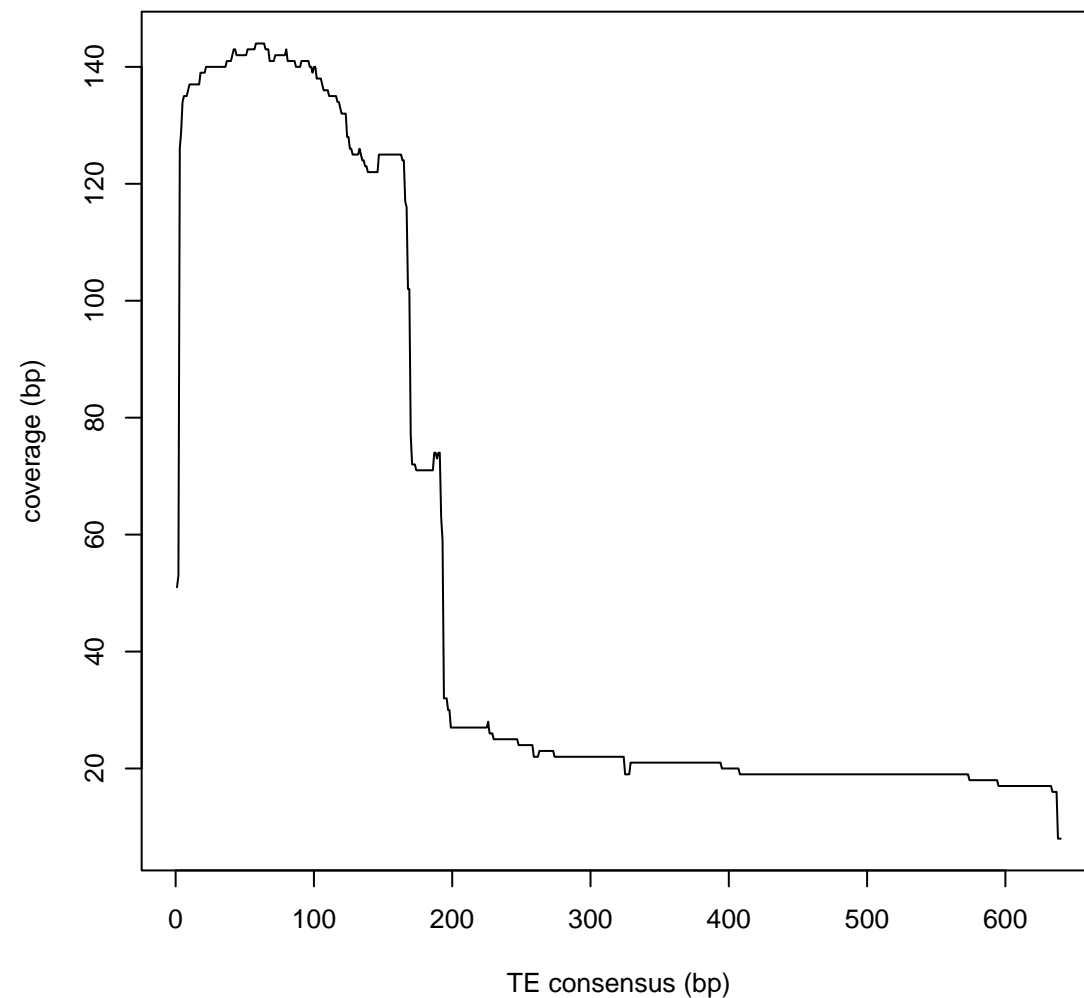


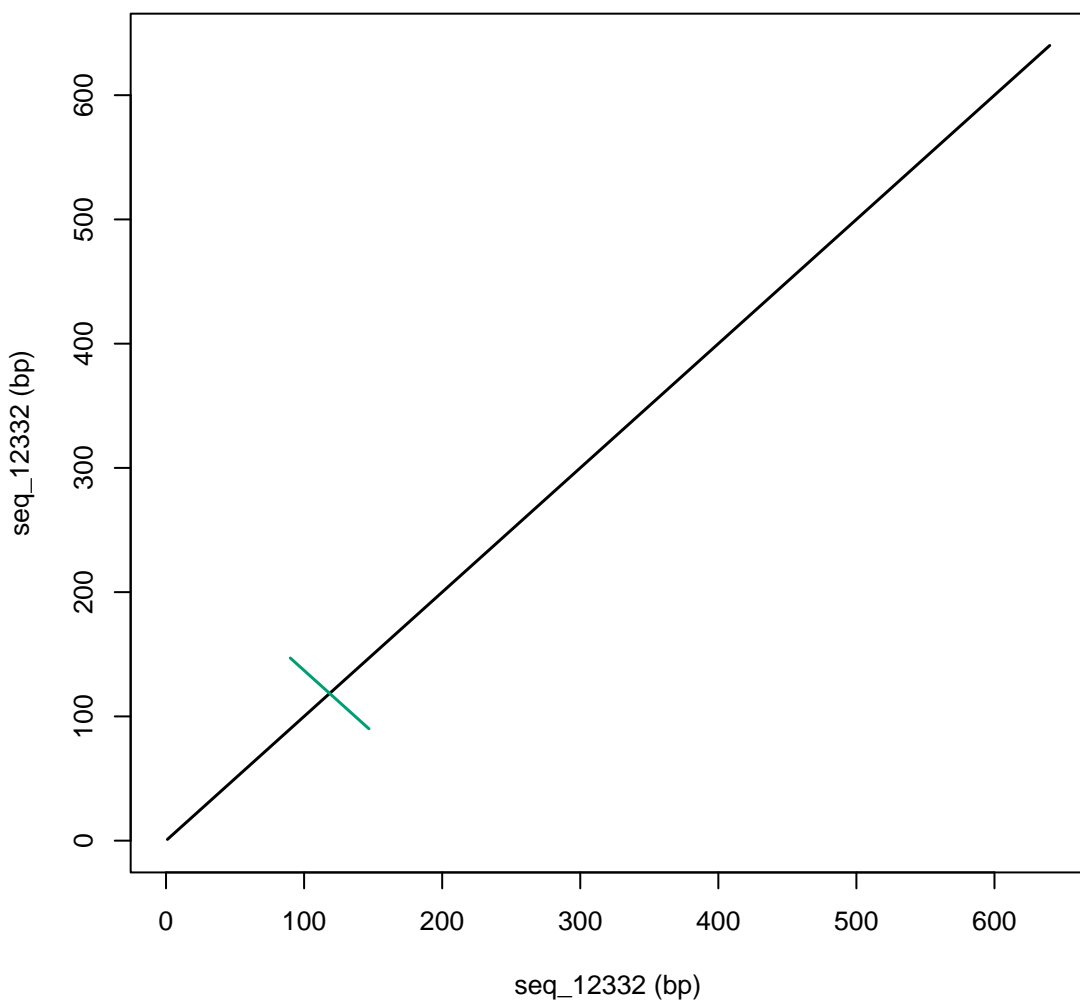
TE: seq_12332
consensus size: 640bp; fragments: 162; full length: 12 (>=576bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

