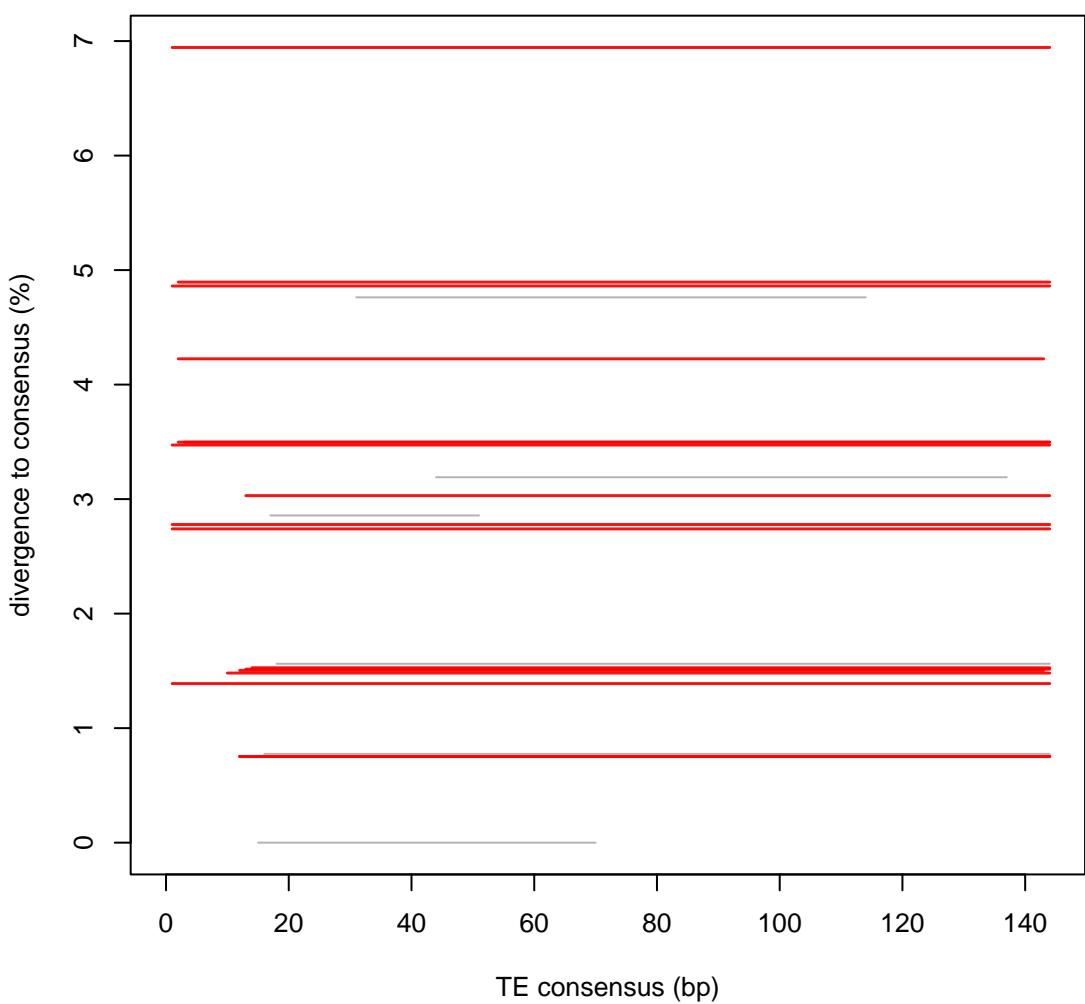
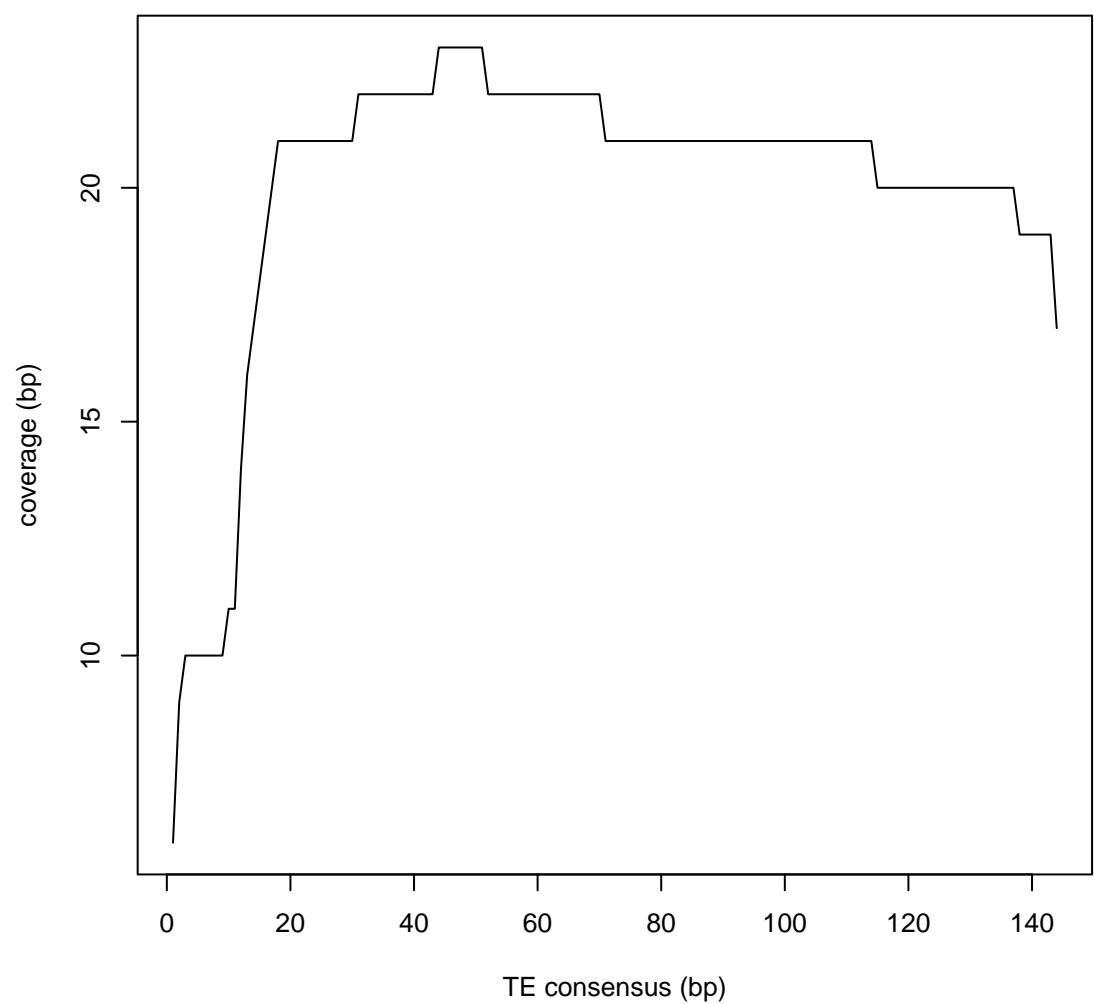


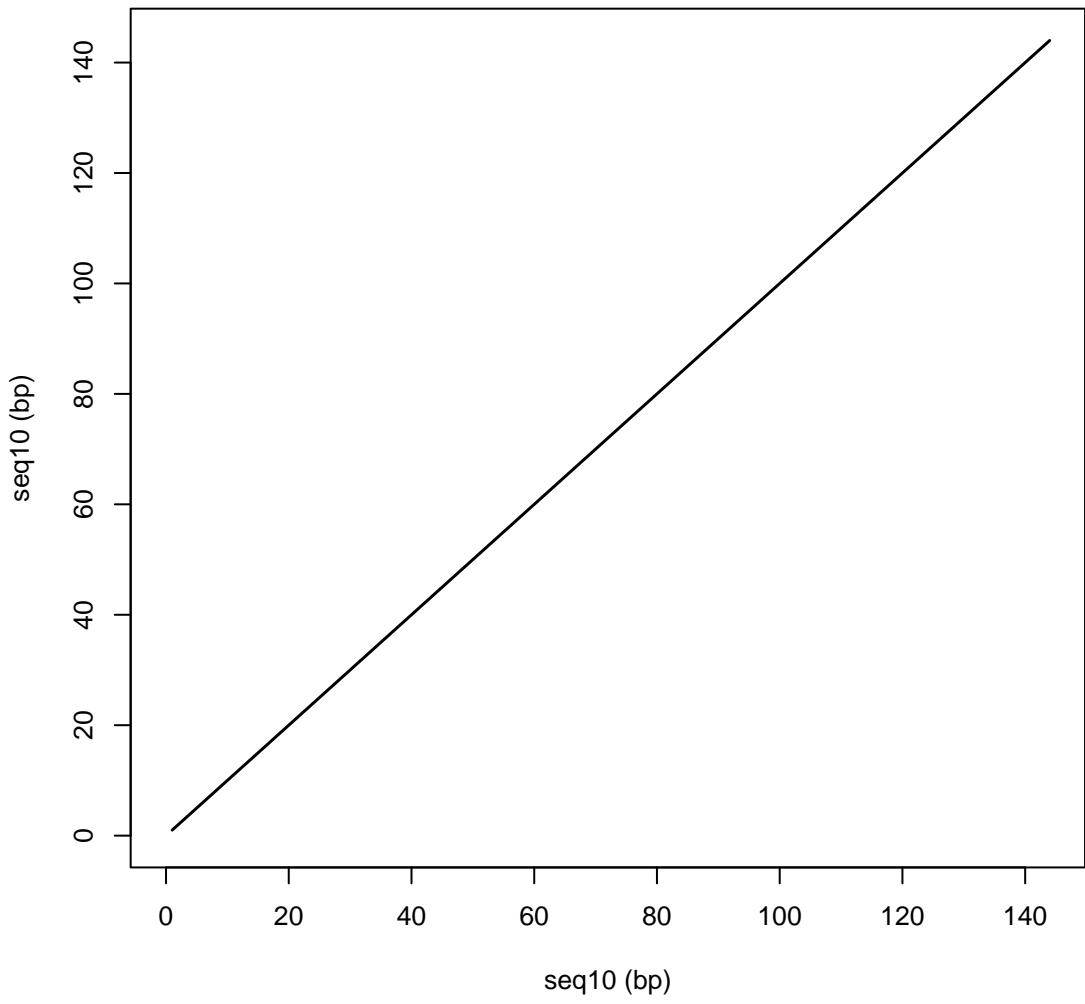
TE: seq10
consensus size: 144bp; fragments: 23; full length: 17 (≥ 129.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

