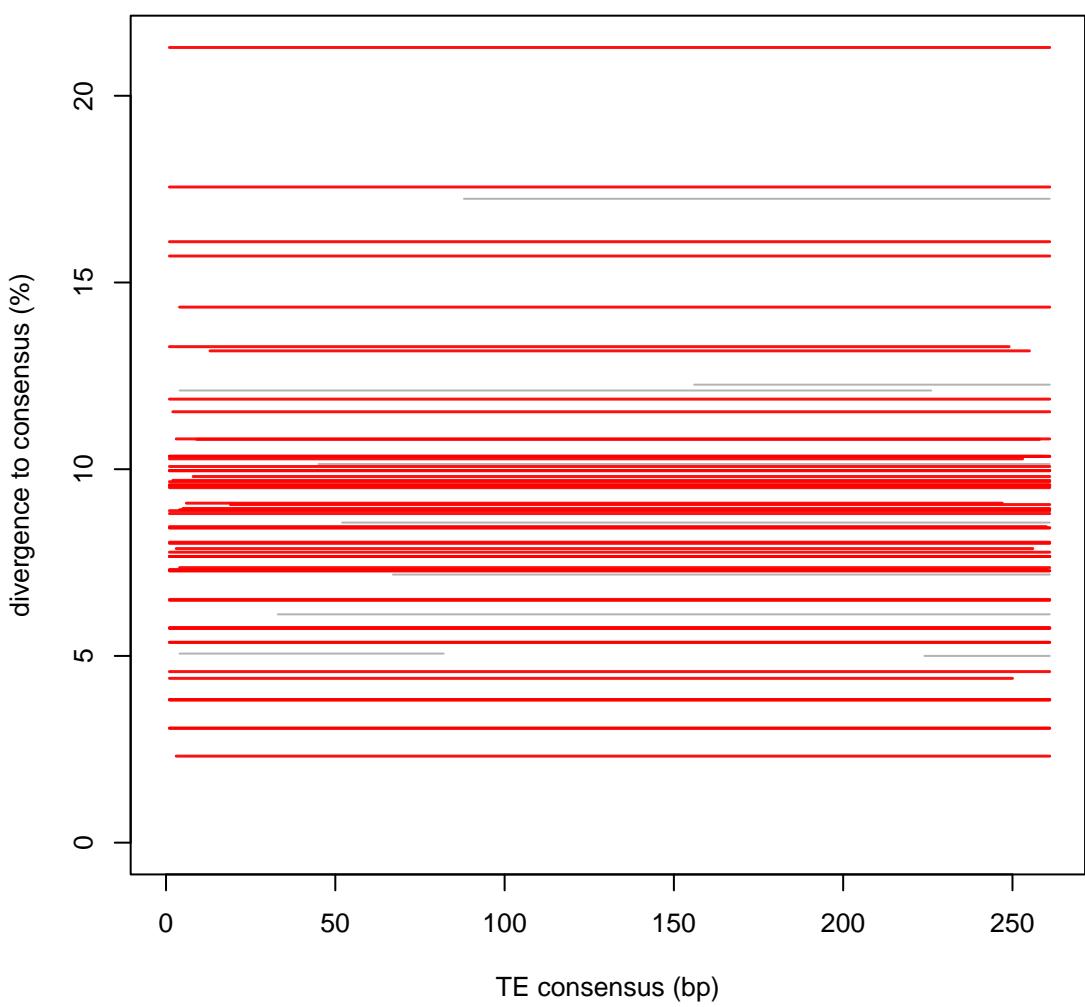
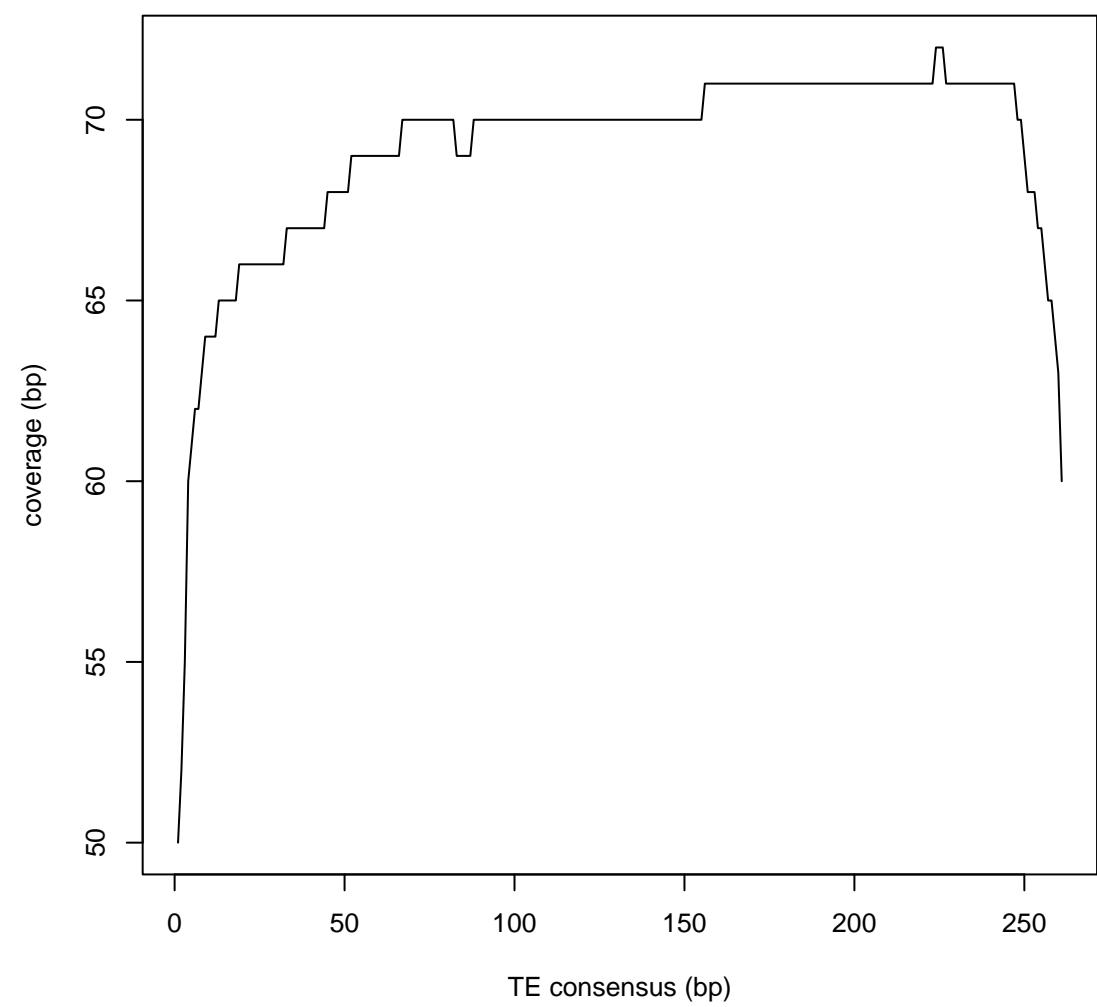


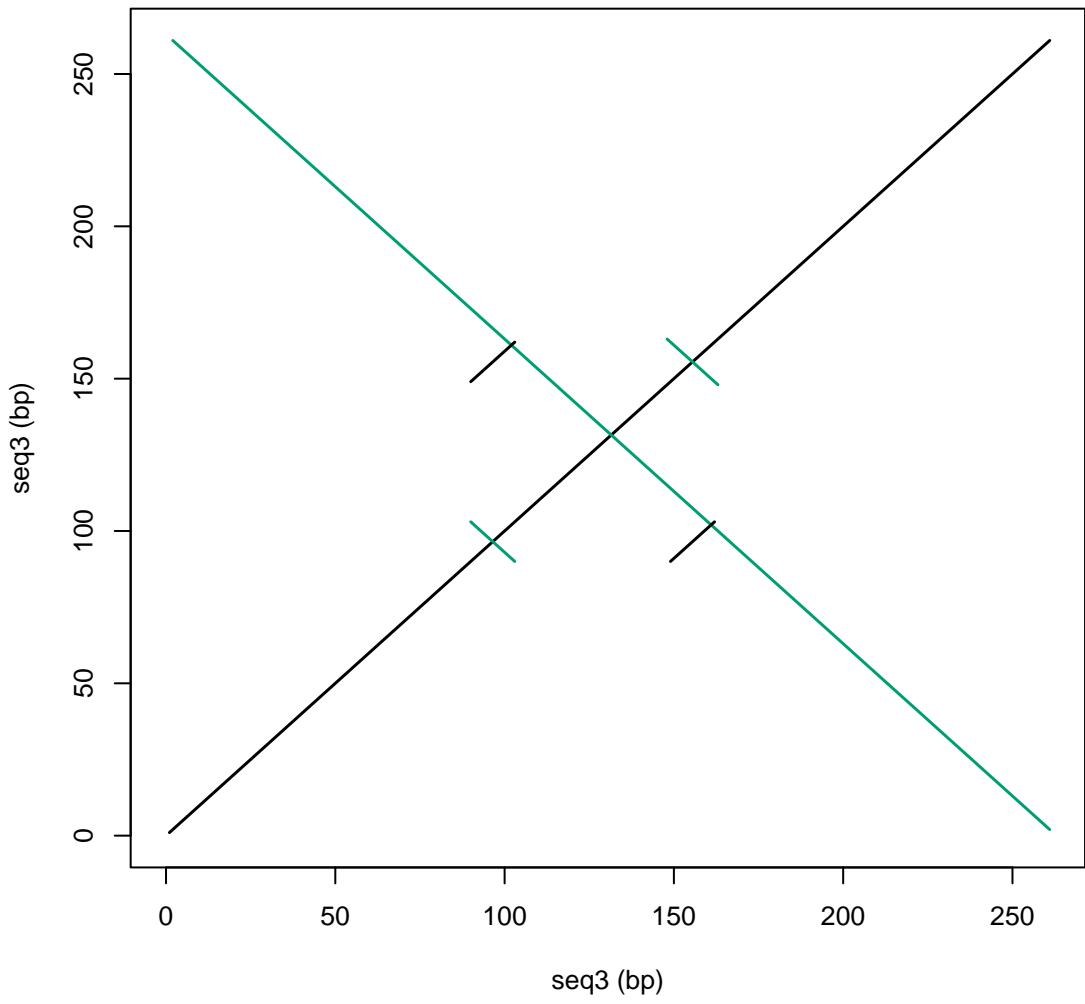
TE: seq3
consensus size: 261bp; fragments: 73; full length: 64 (≥ 234.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

