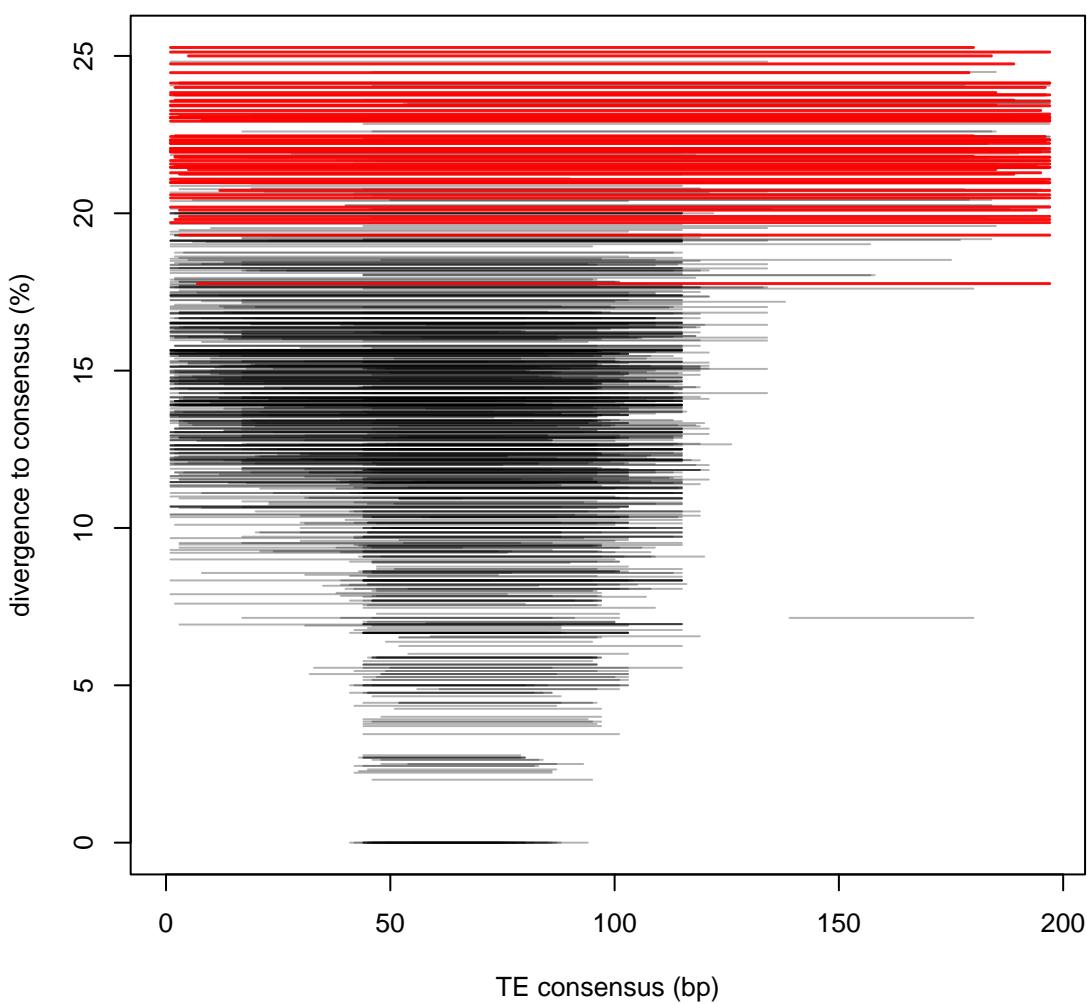
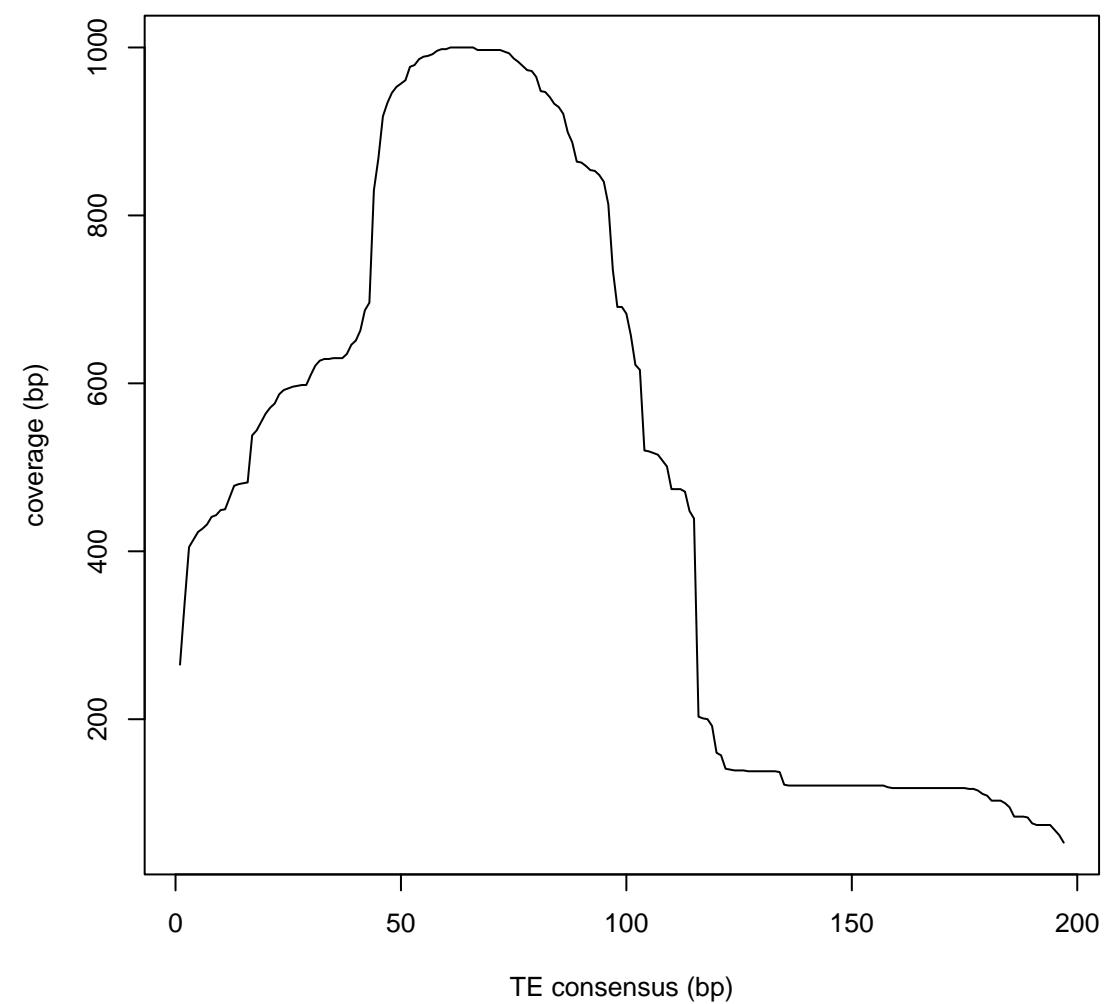


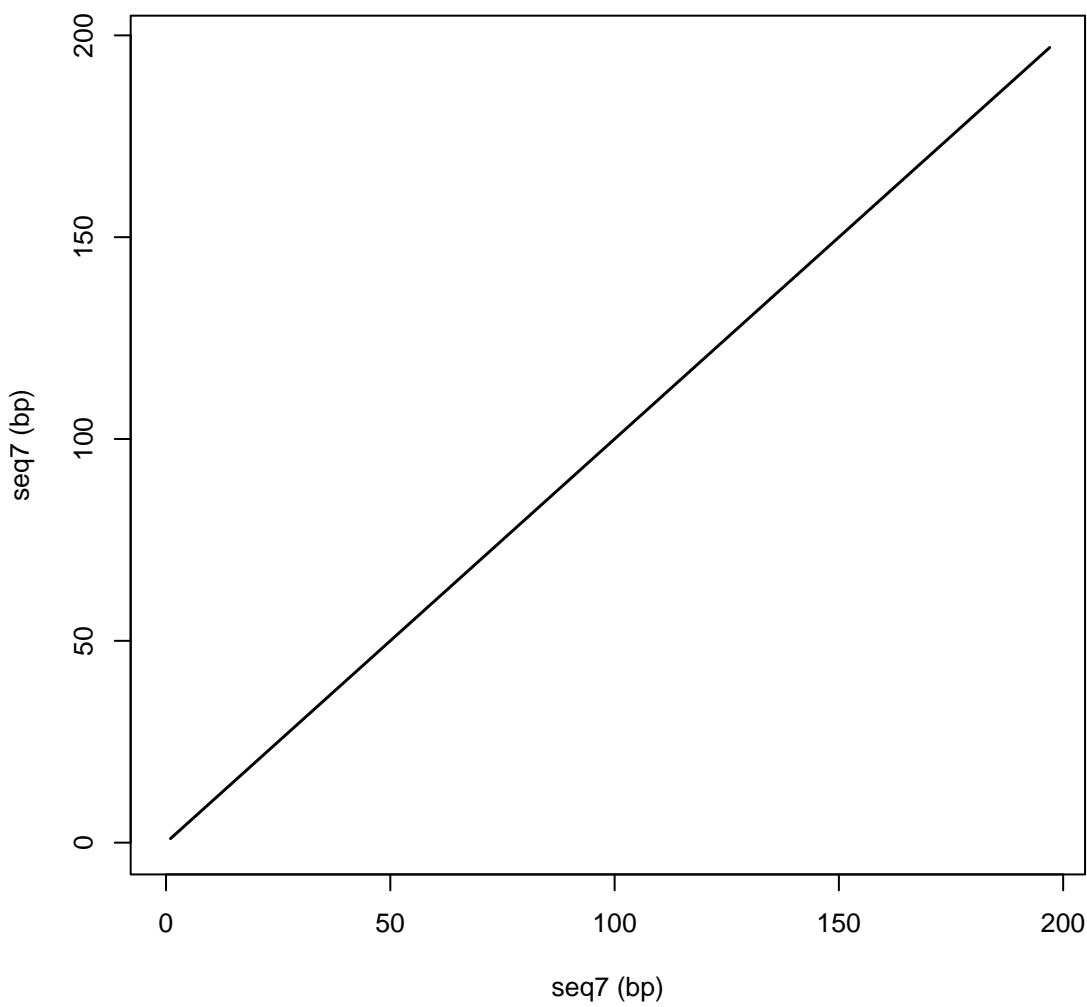
TE: seq7
consensus size: 197bp; fragments: 1001; full length: 63 (≥ 177.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

