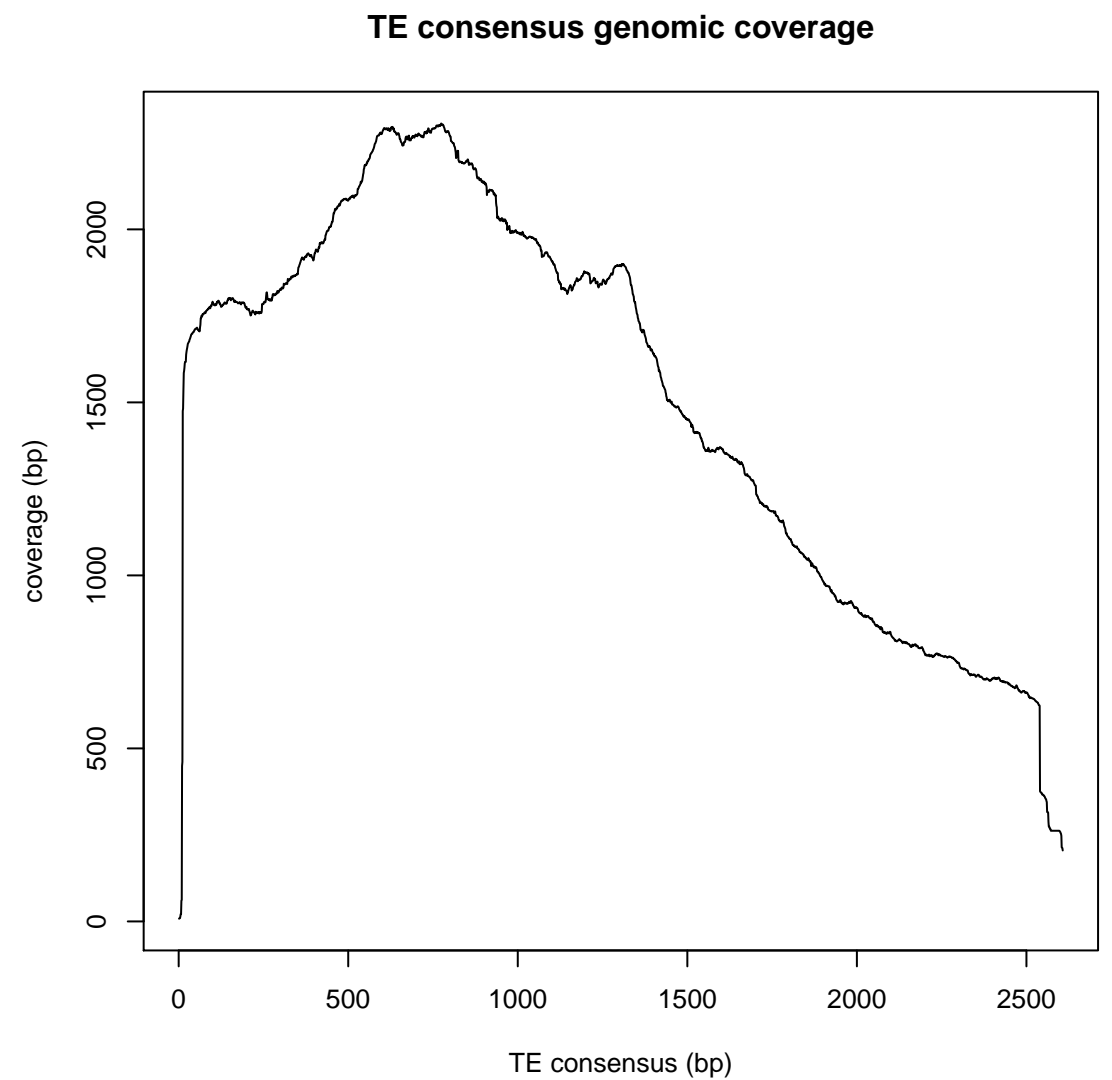
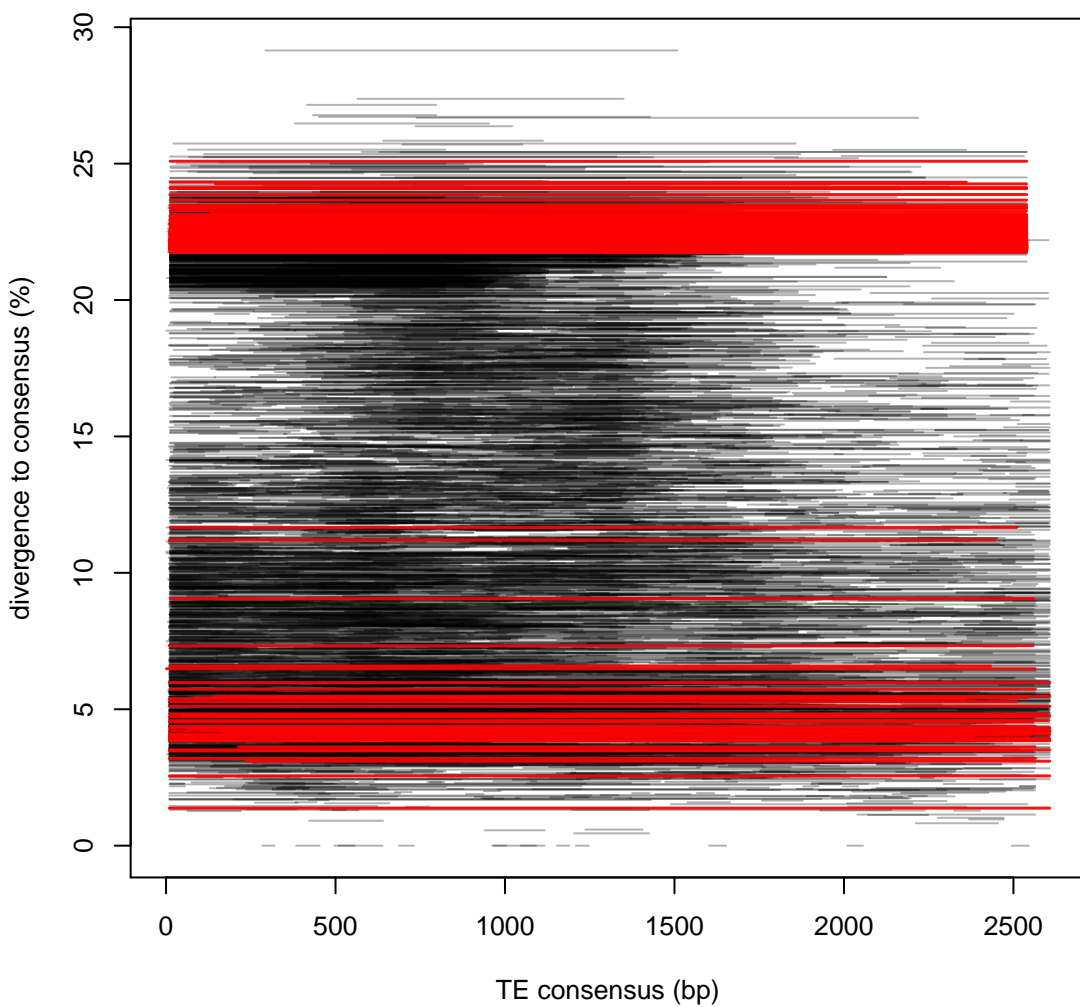
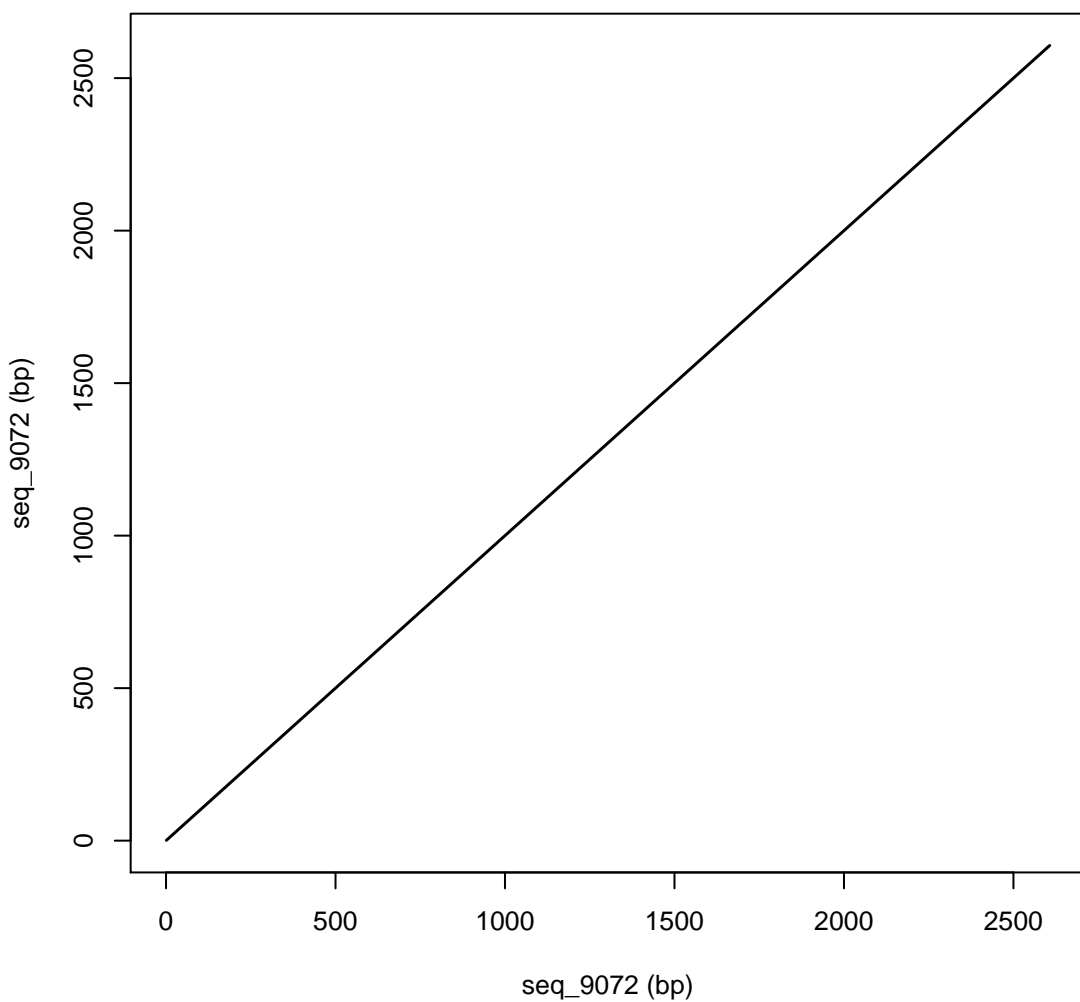


TE: seq_9072
consensus size: 2607bp; fragments: 6009; full length: 276 (≥ 2346.3 bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

