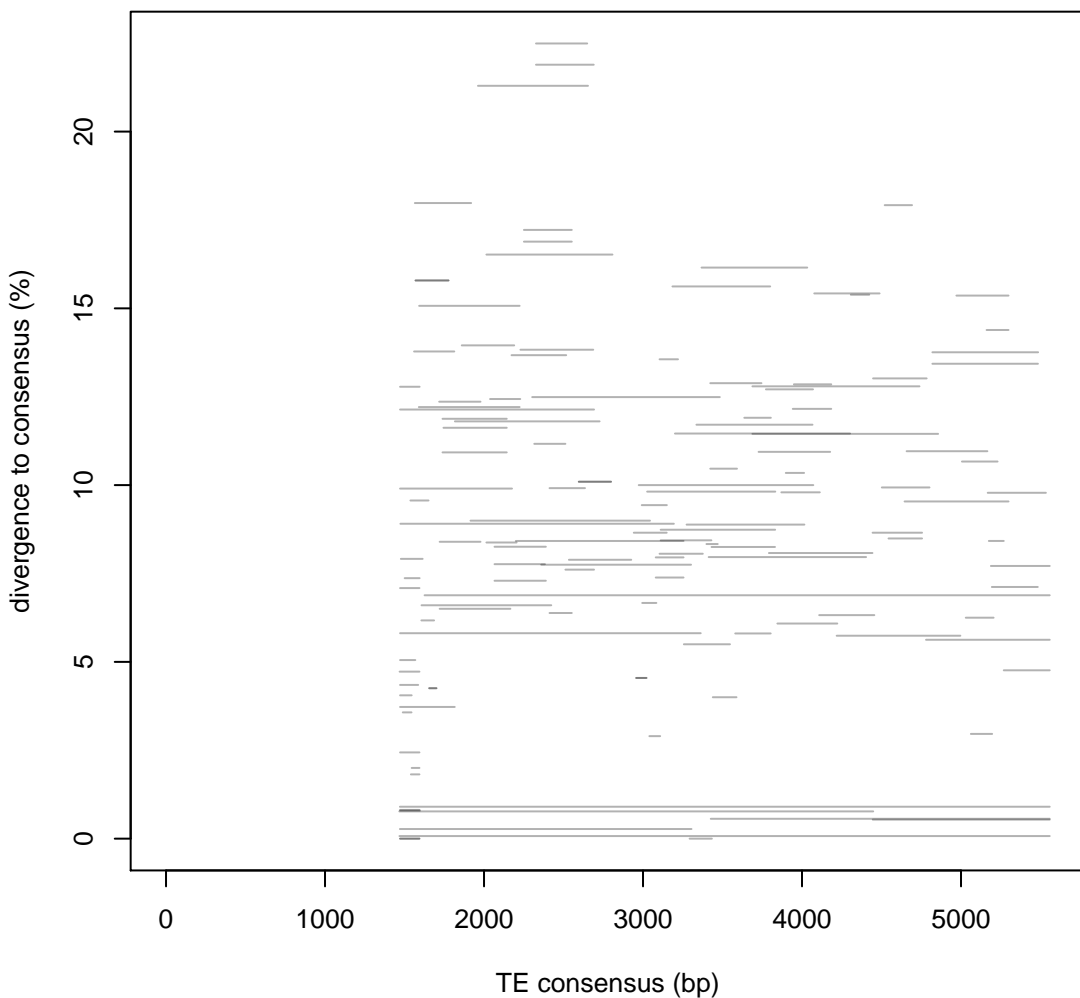
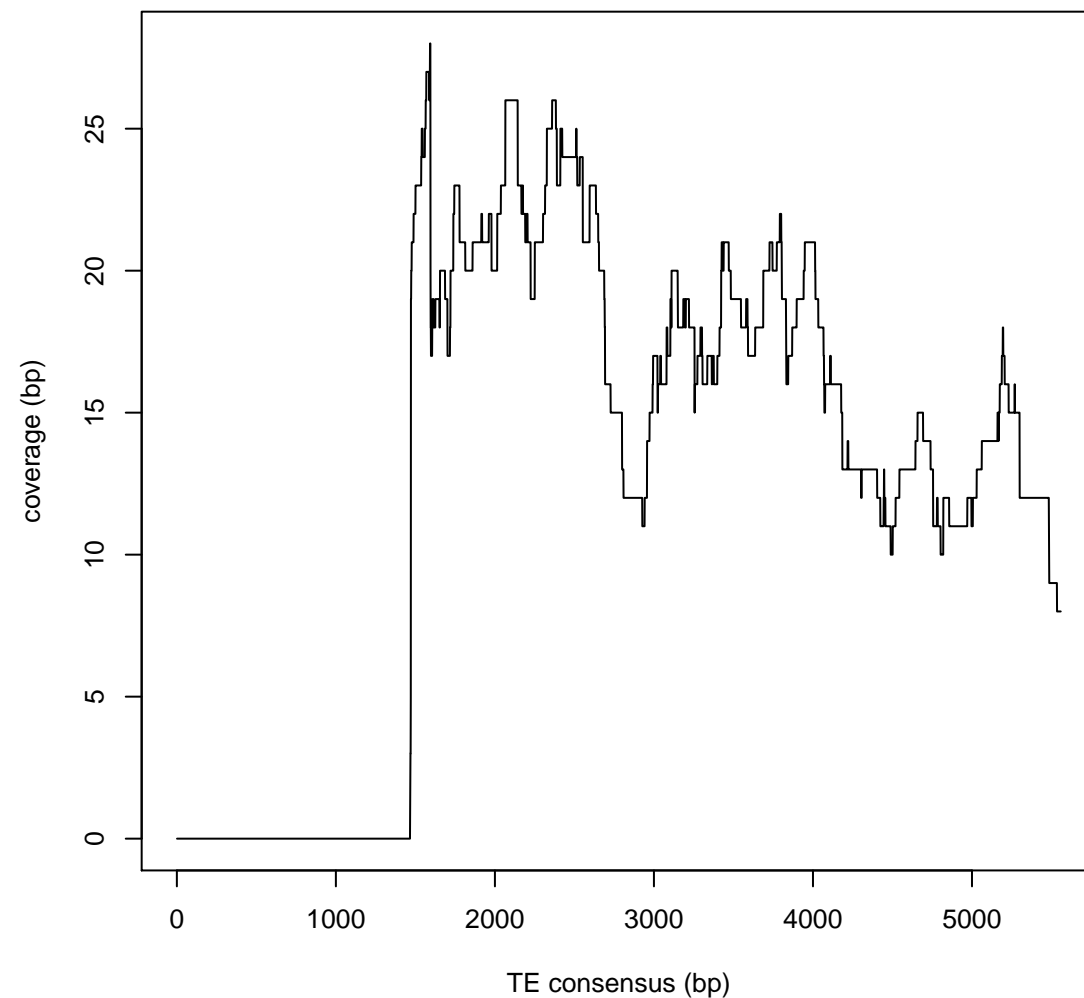


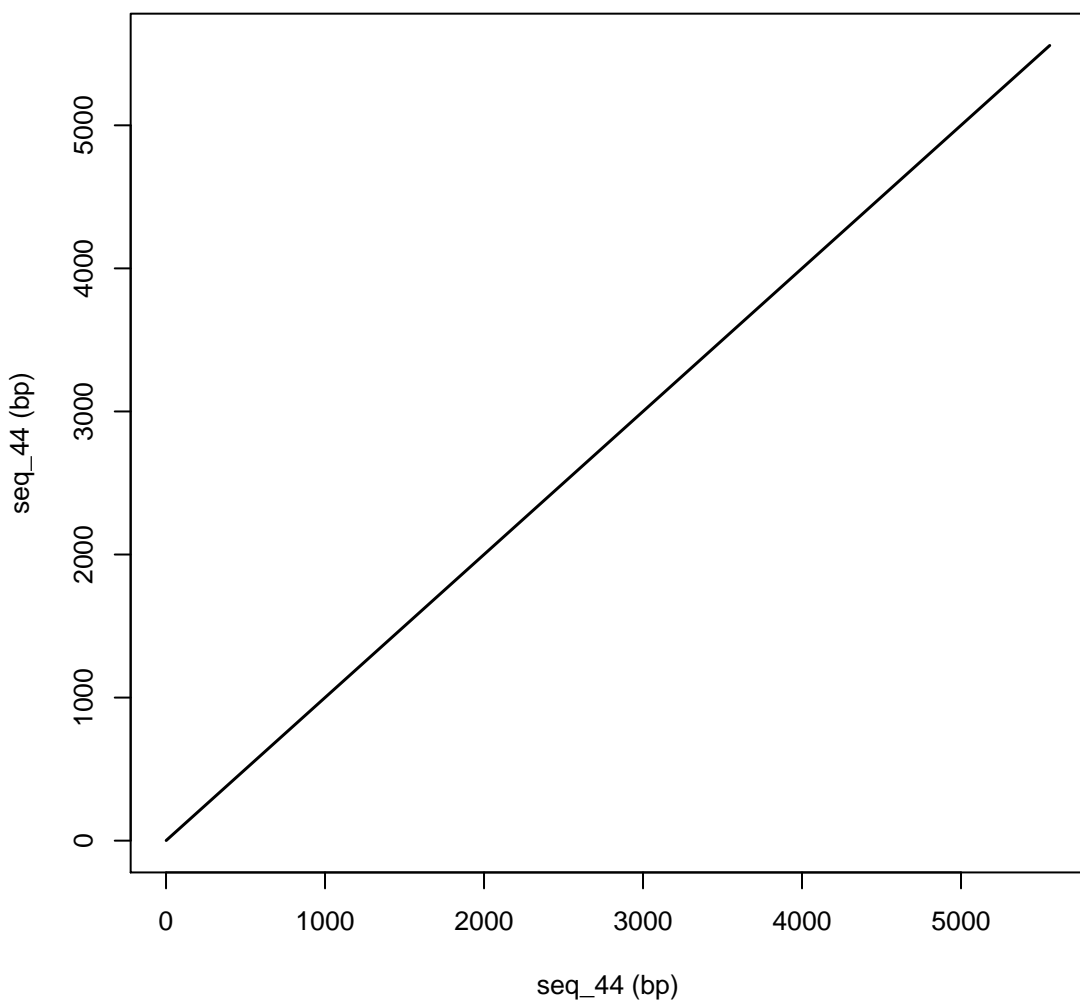
TE: seq_44
consensus size: 5558bp; fragments: 134; full length: 0 (>=5002.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

