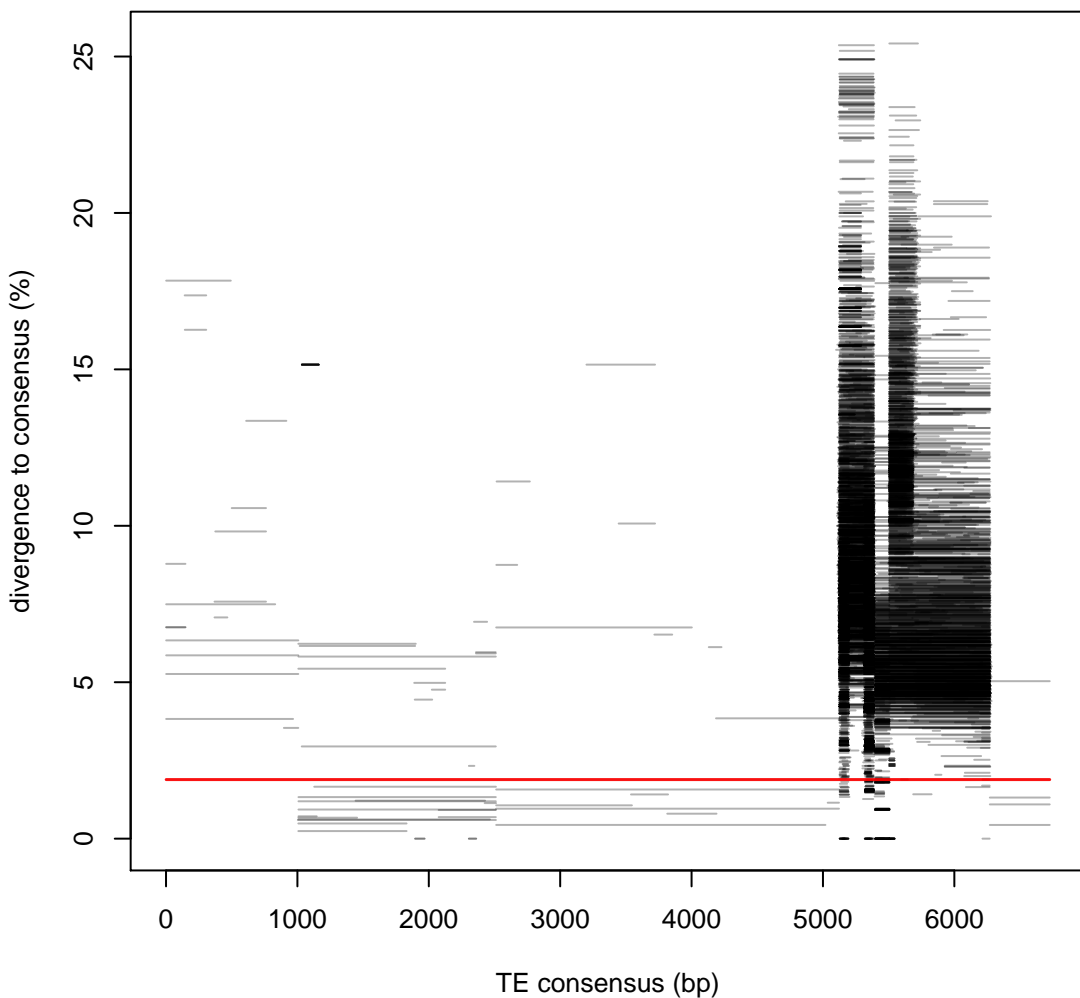
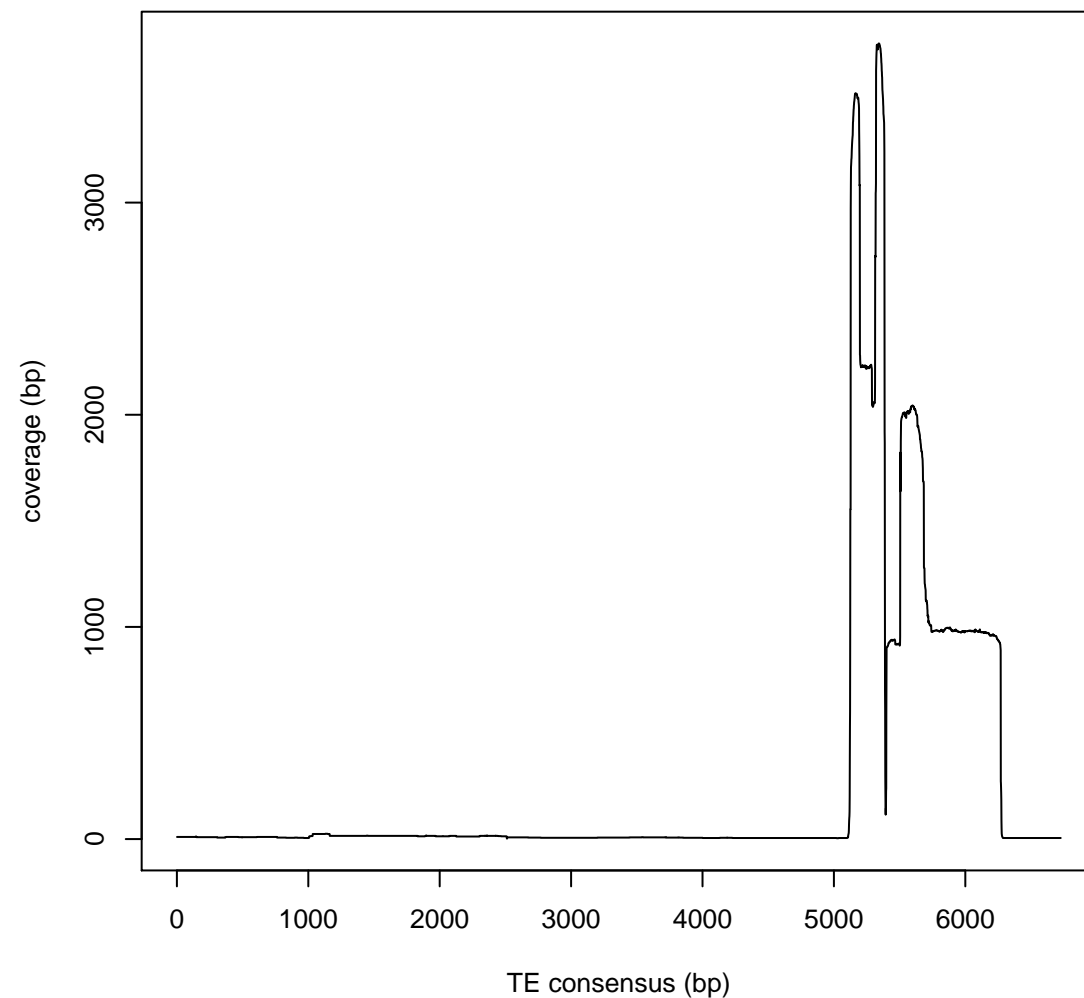


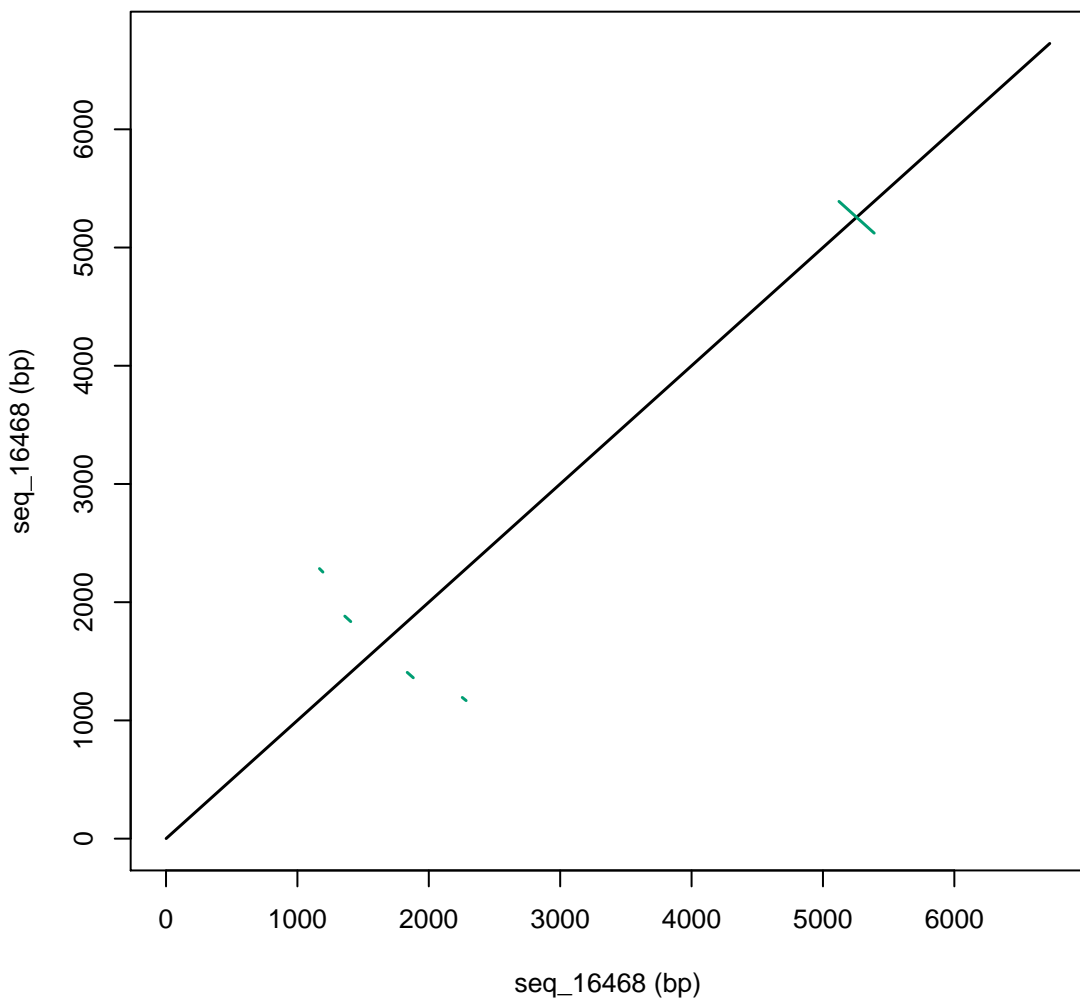
TE: seq_16468
consensus size: 6726bp; fragments: 8482; full length: 1 (>=6053.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

