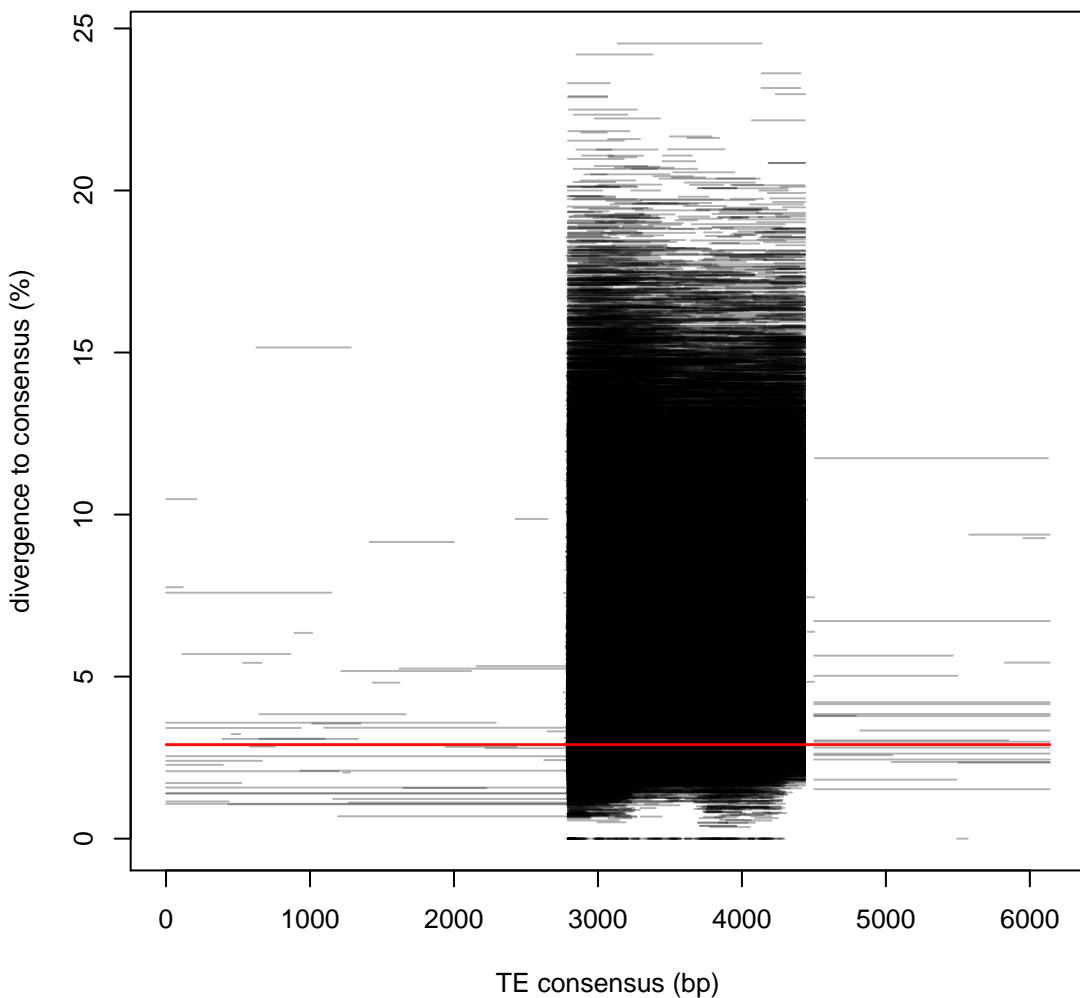
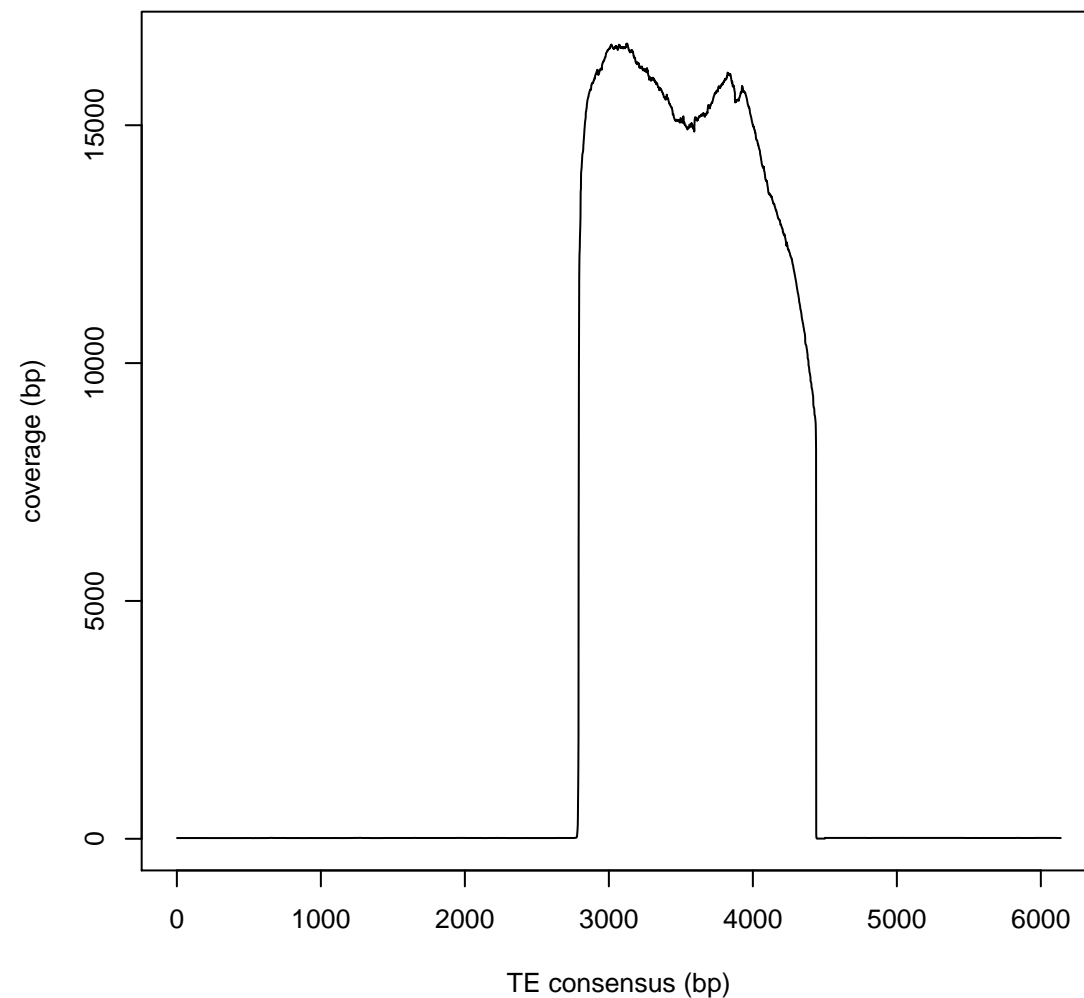


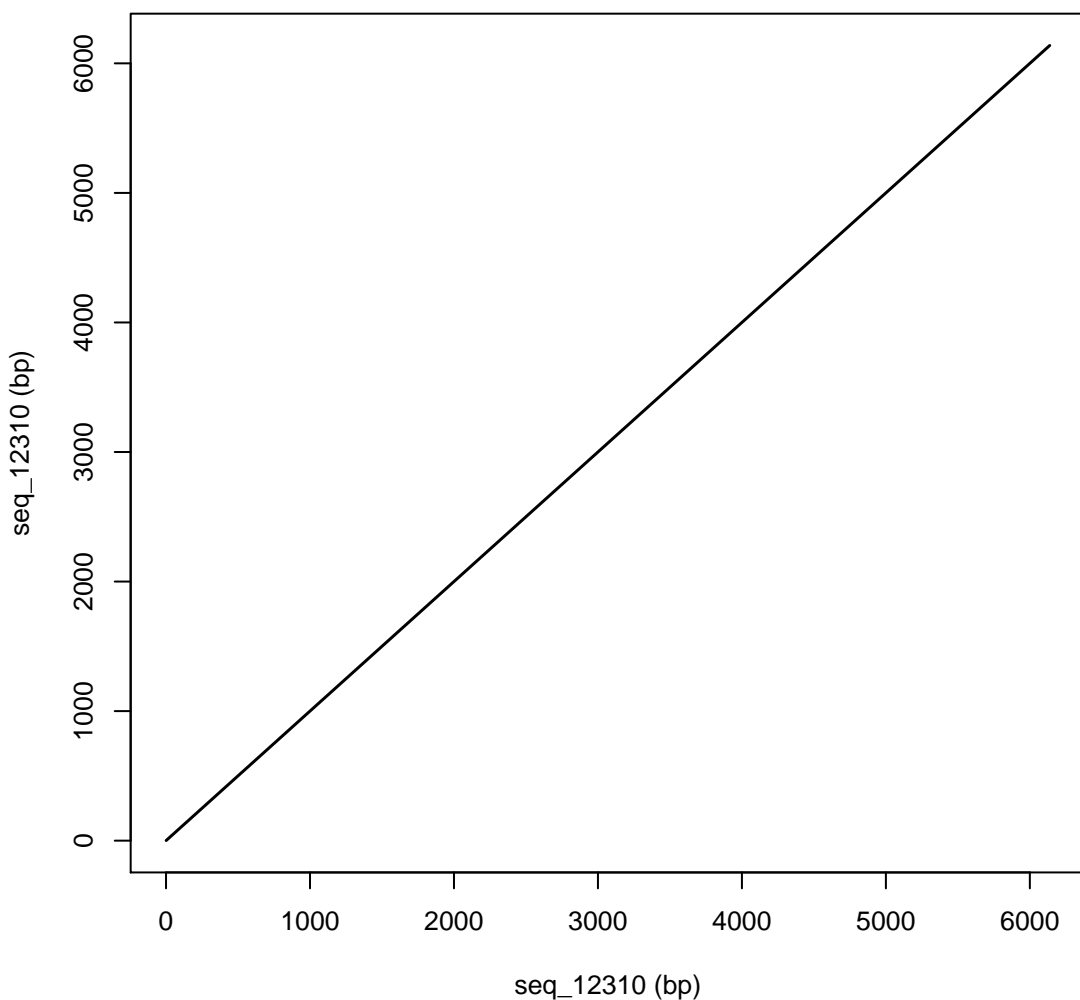
TE: seq_12310
consensus size: 6138bp; fragments: 38705; full length: 1 (>=5524.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

