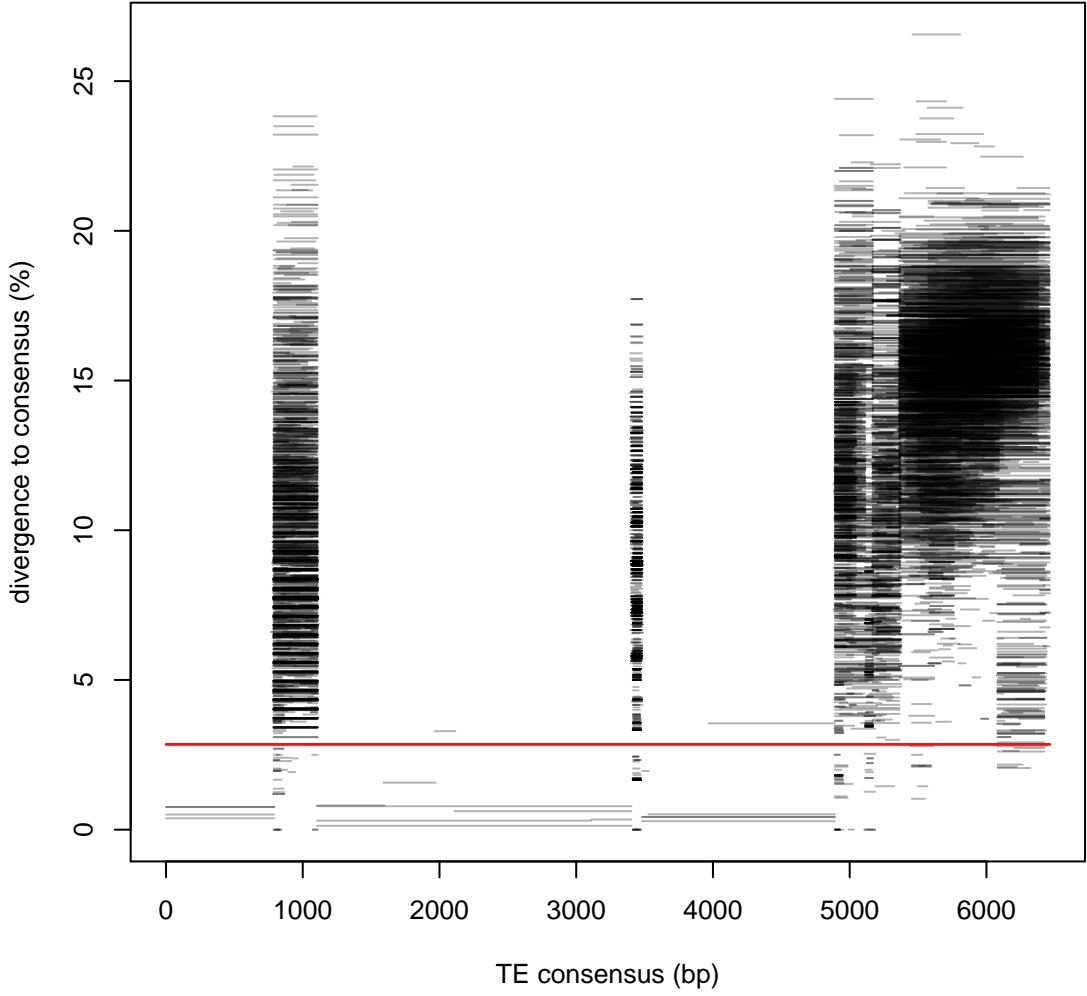
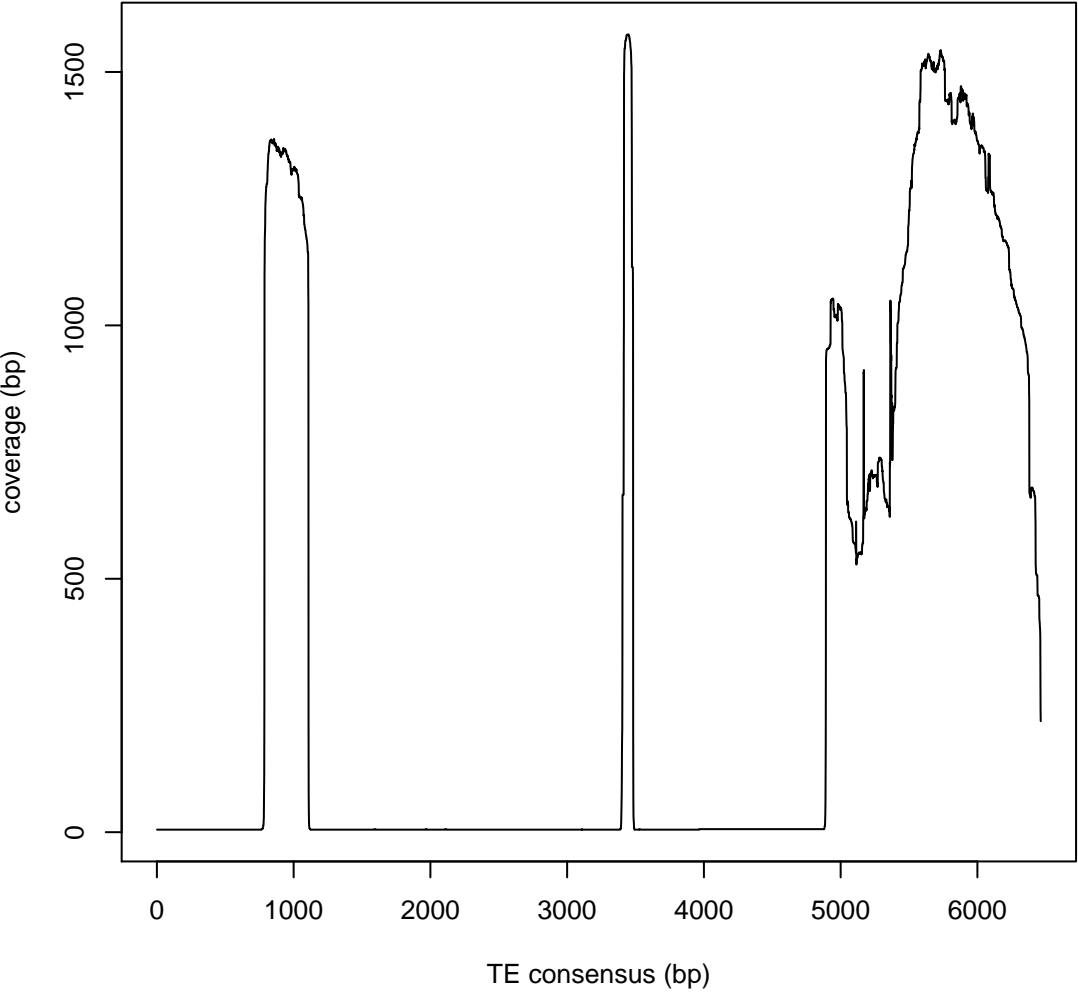


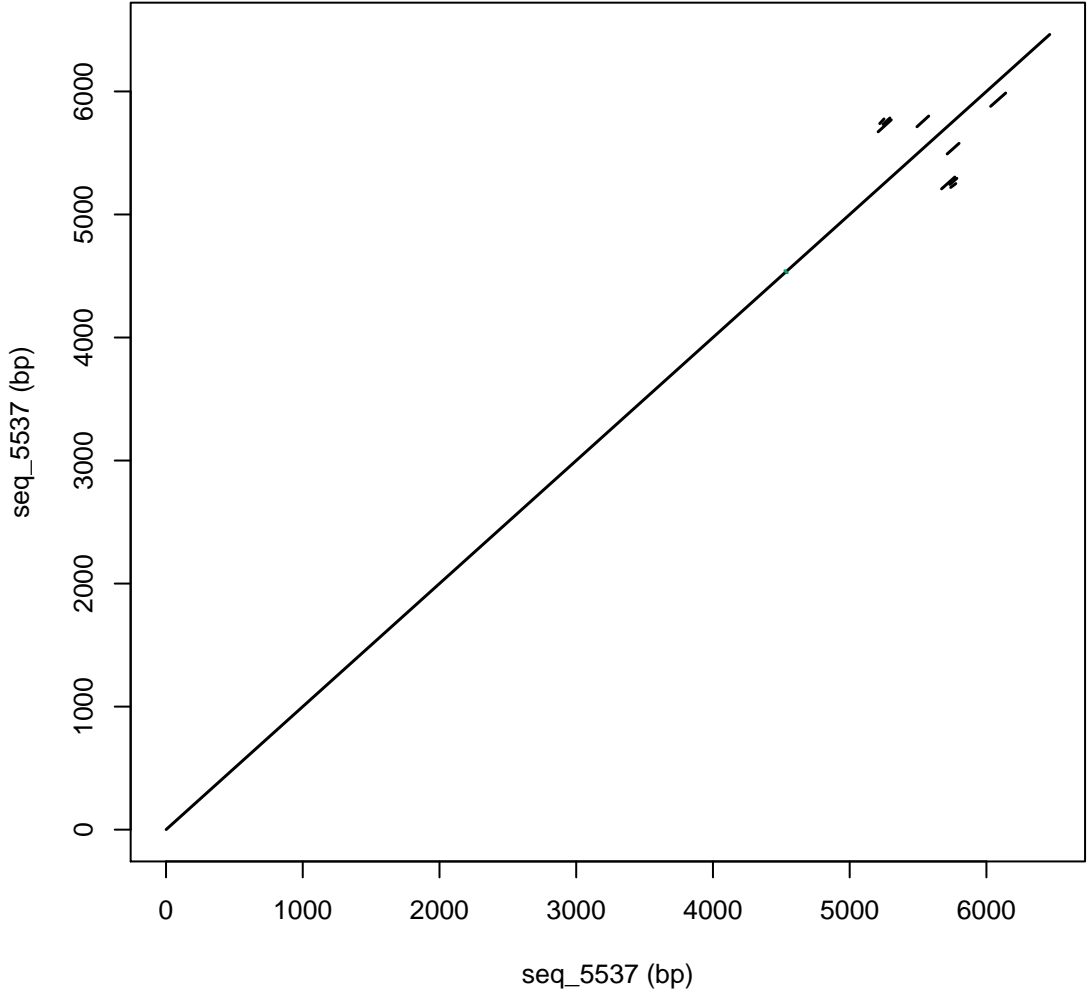
TE: seq_5537
consensus size: 6463bp; fragments: 7490; full length: 1 (>=5816.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

