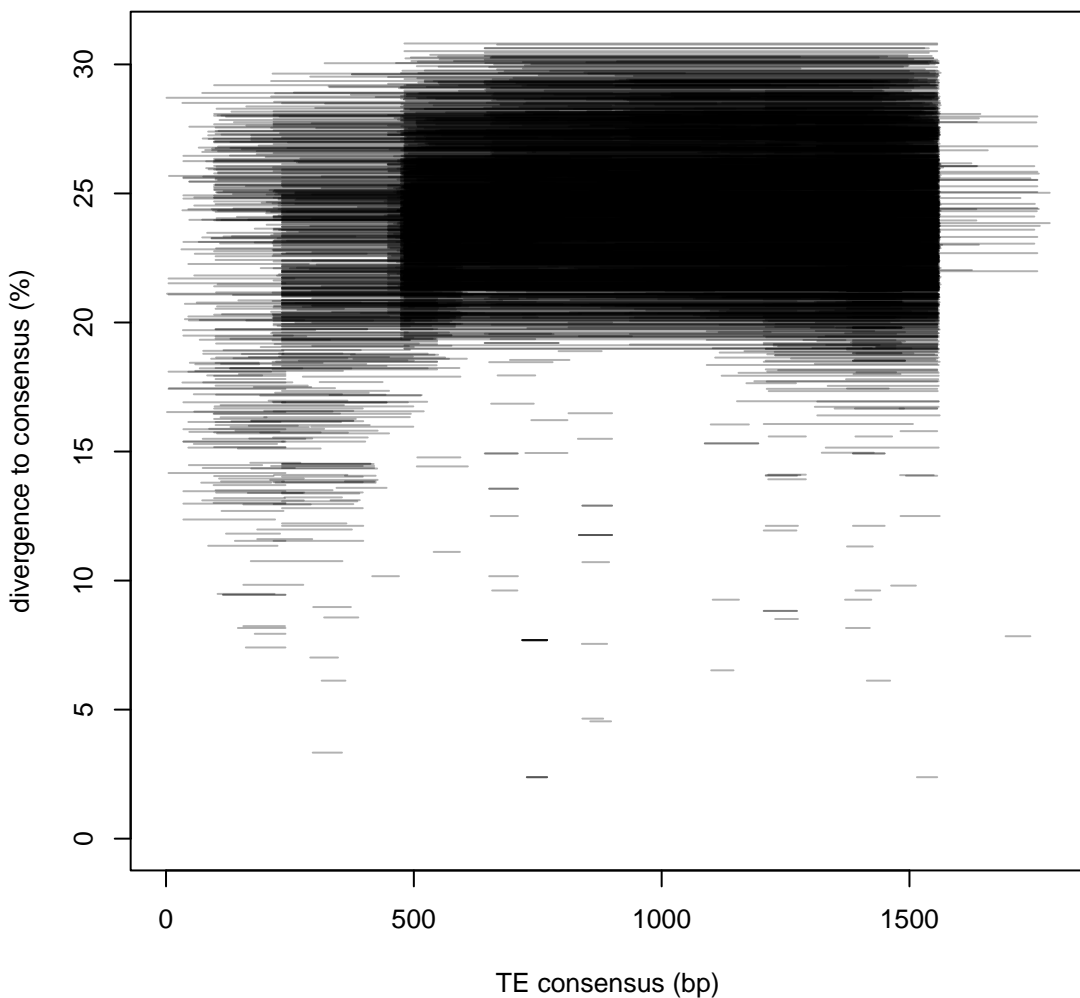
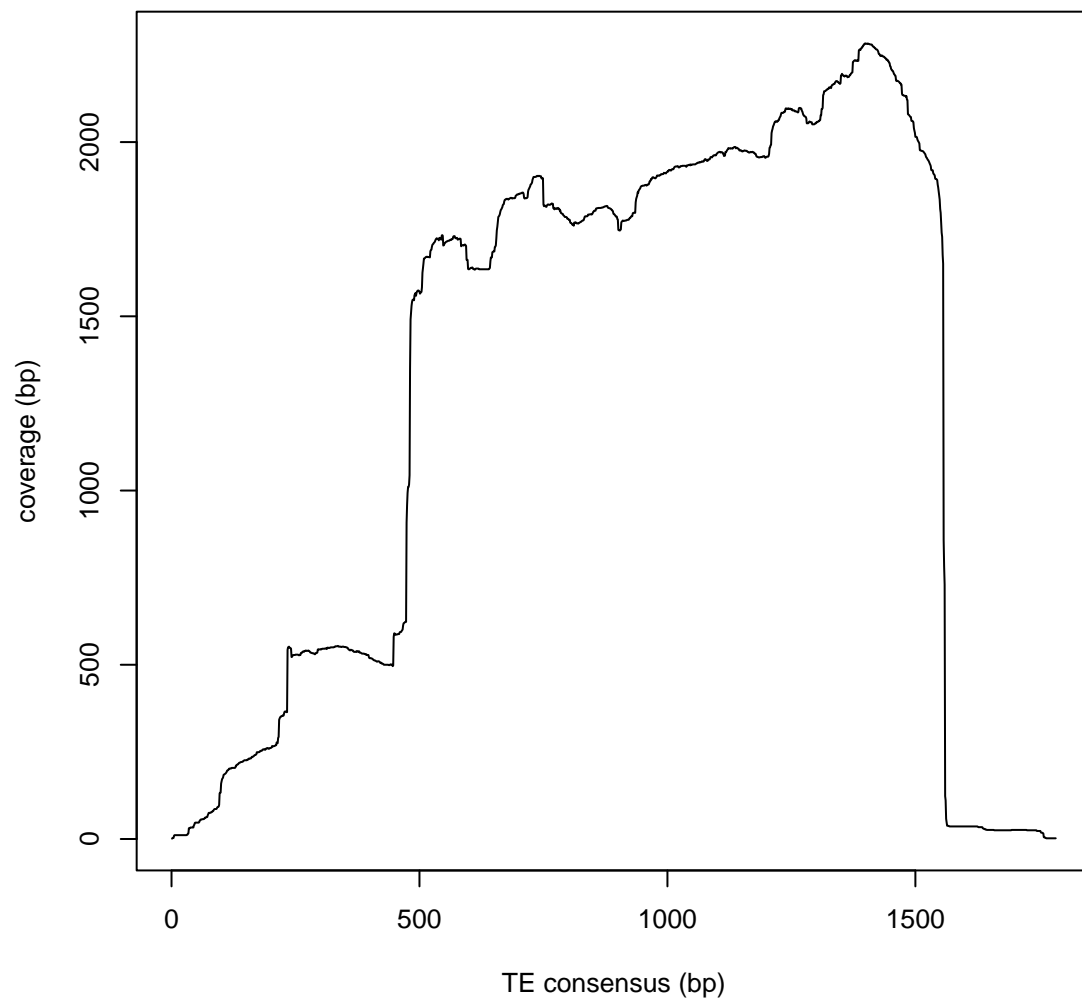


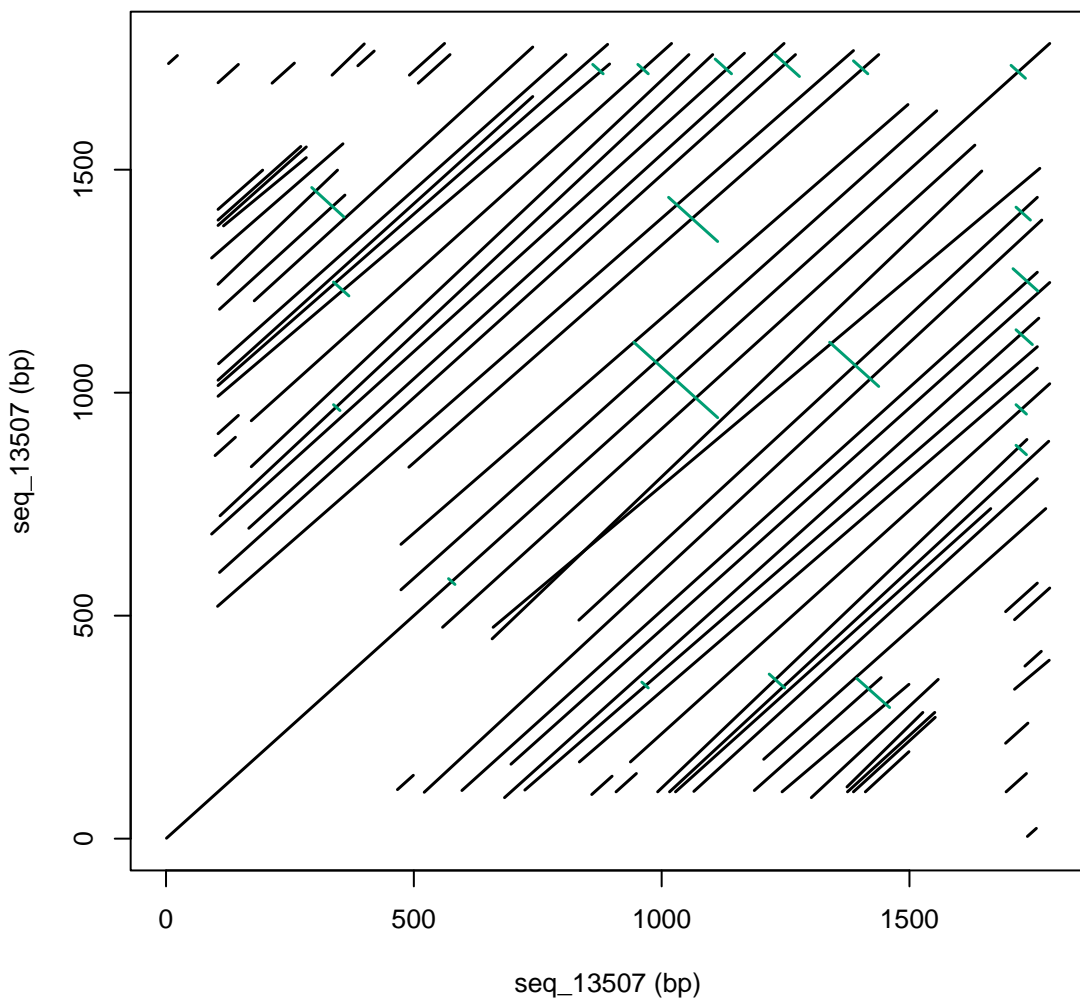
TE: seq_13507
consensus size: 1783bp; fragments: 3161; full length: 0 (>=1604.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

