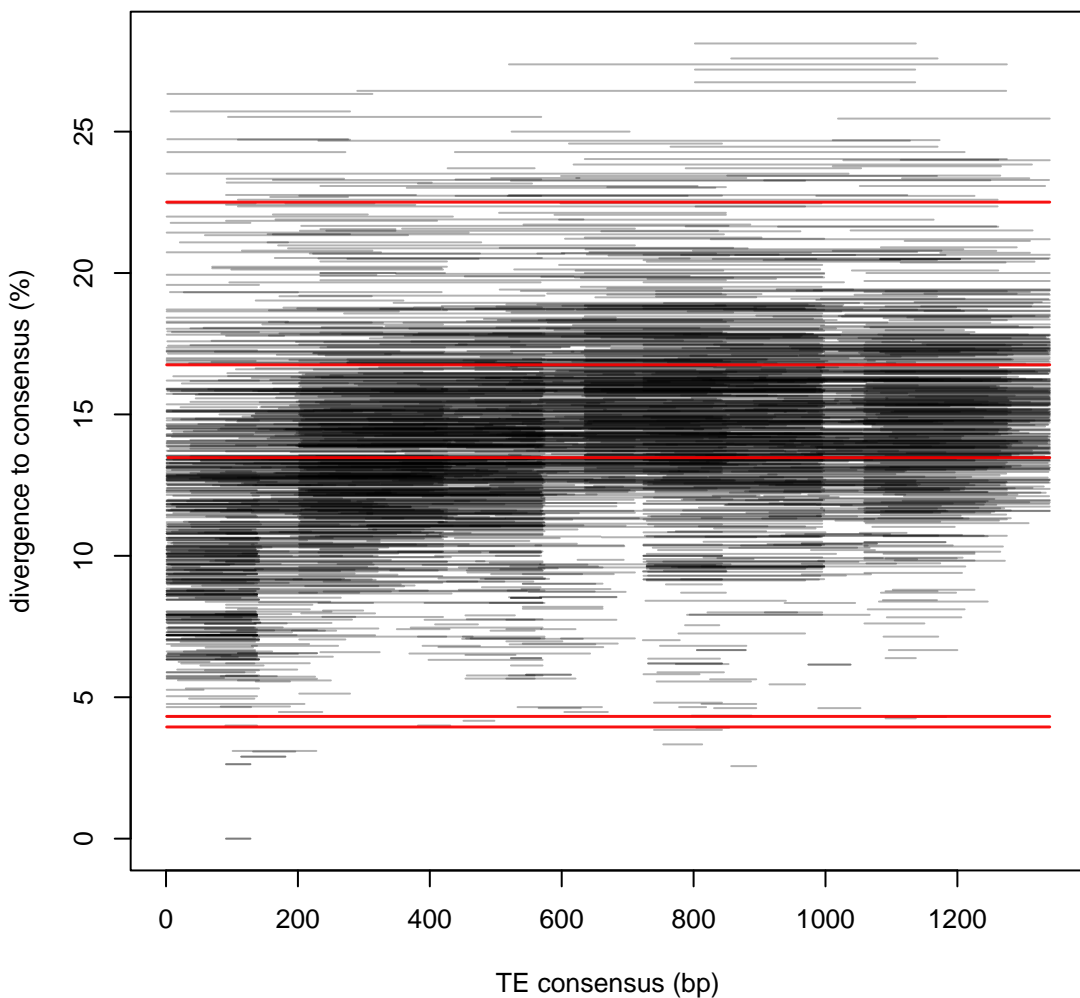
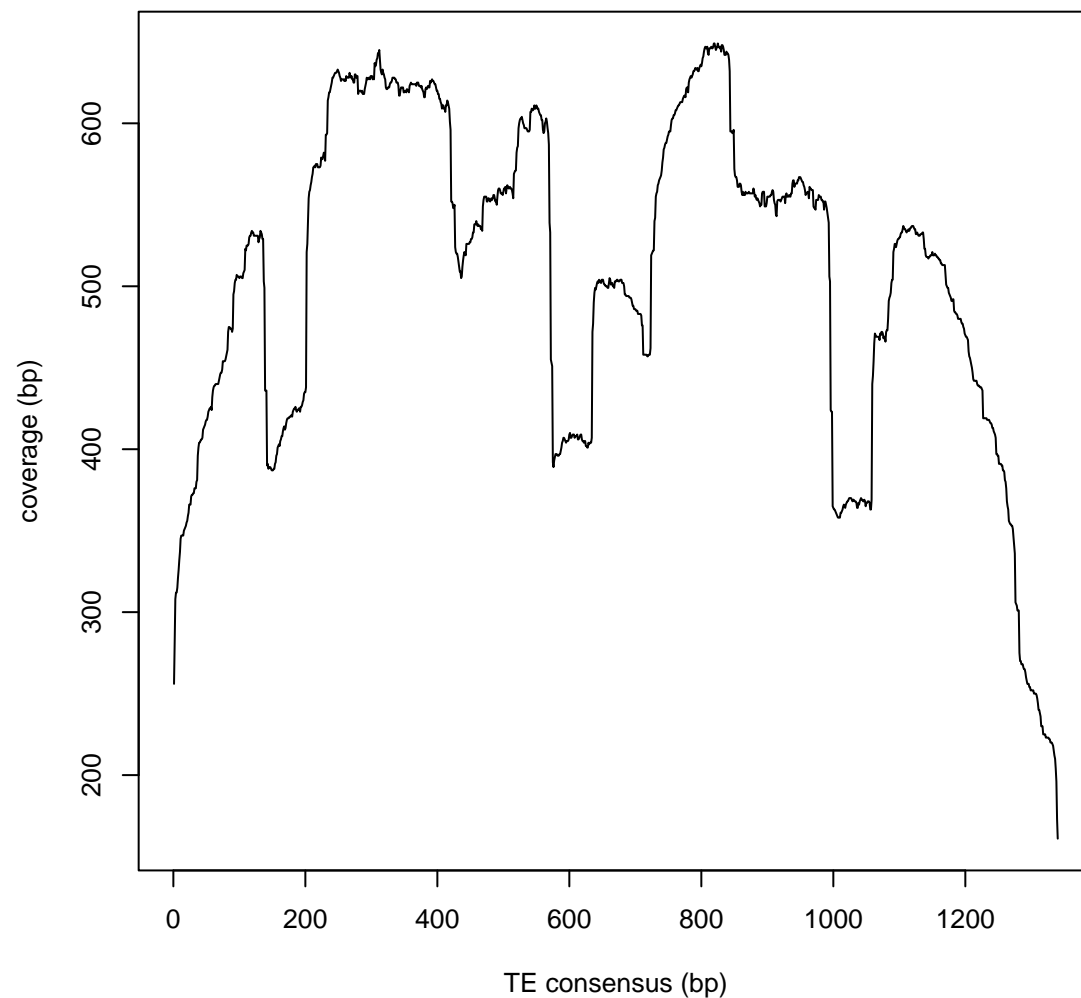


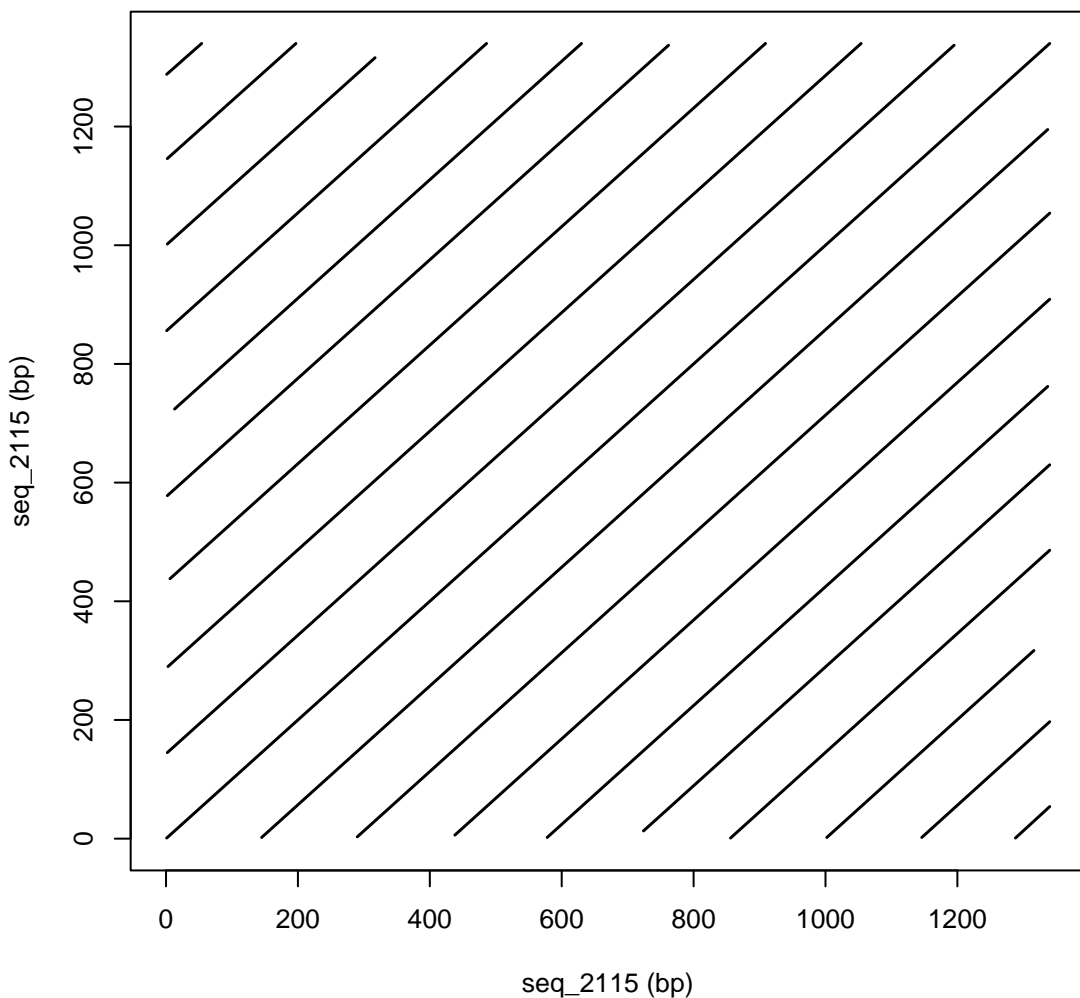
TE: seq_2115
consensus size: 1340bp; fragments: 2146; full length: 5 (≥ 1206 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

