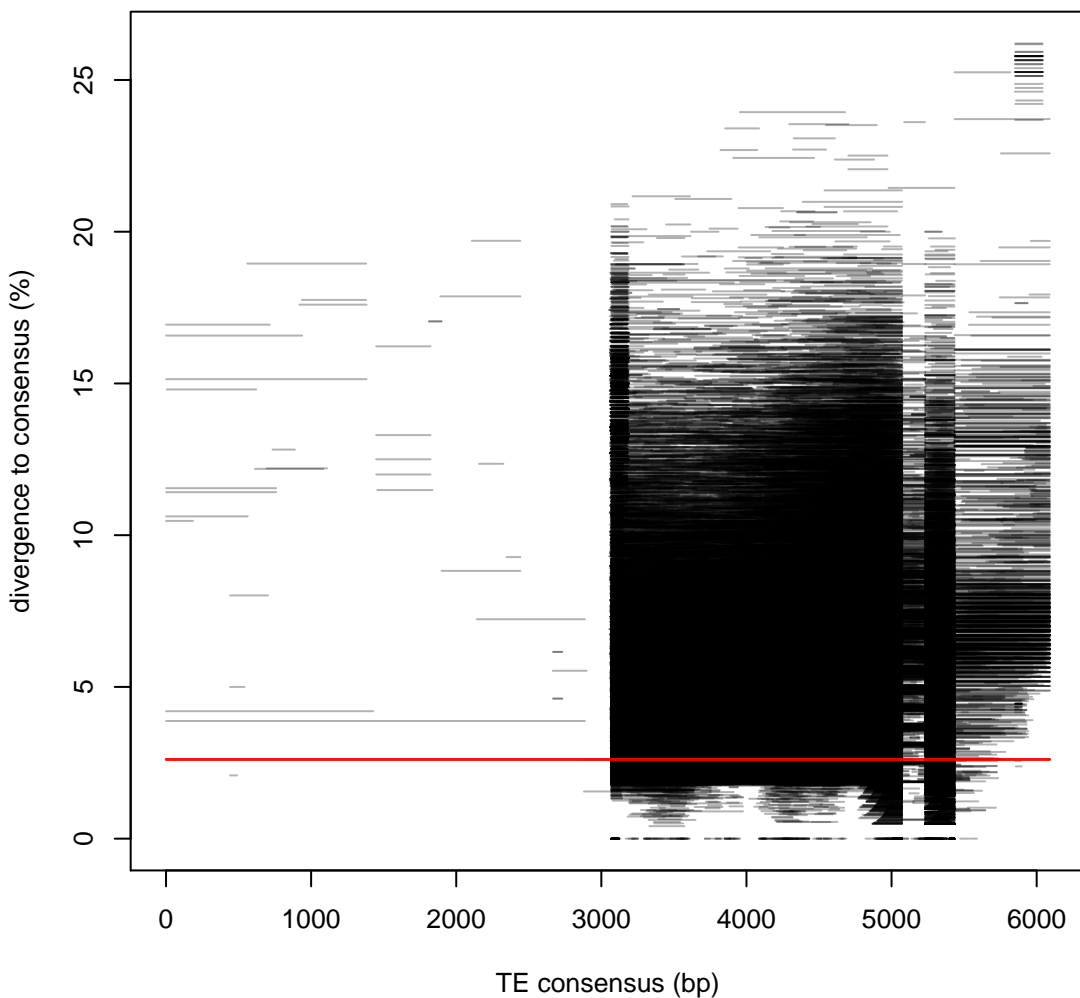
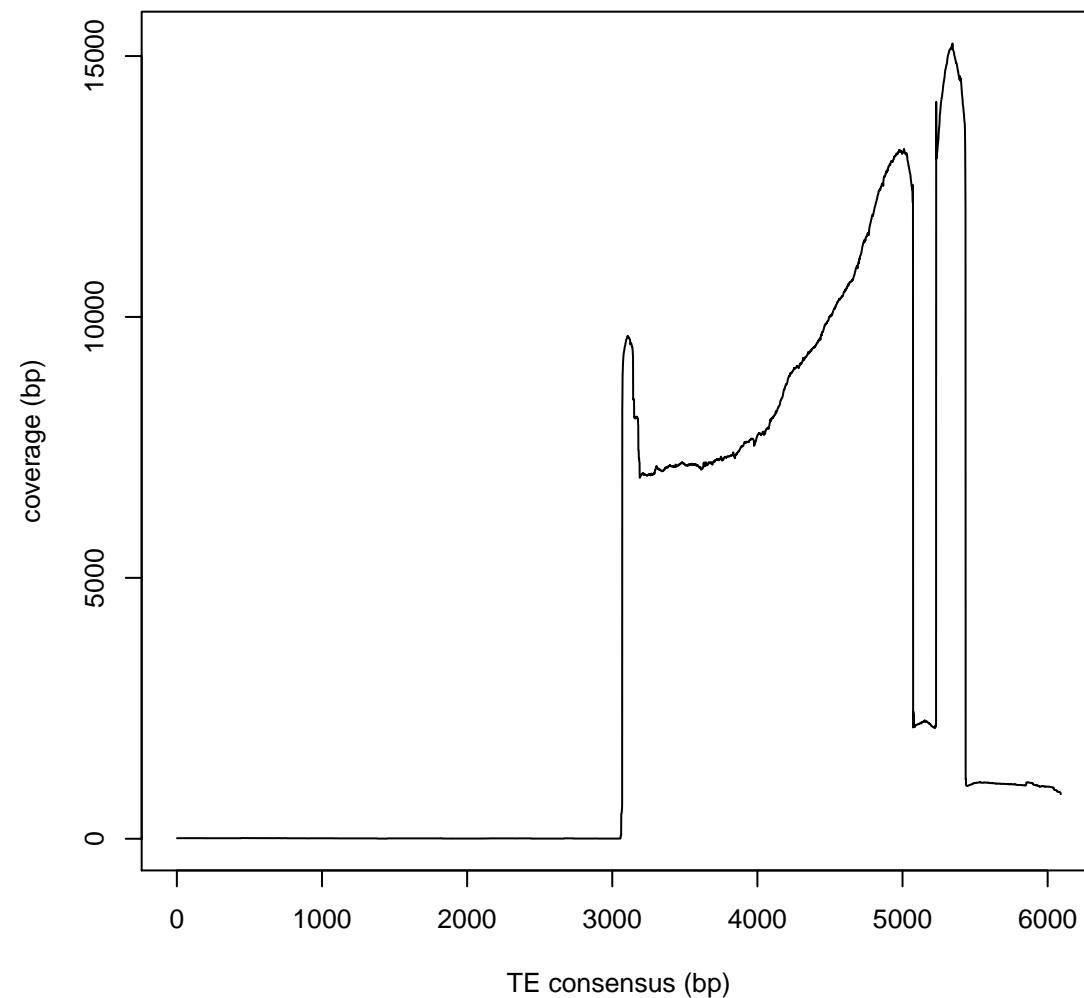


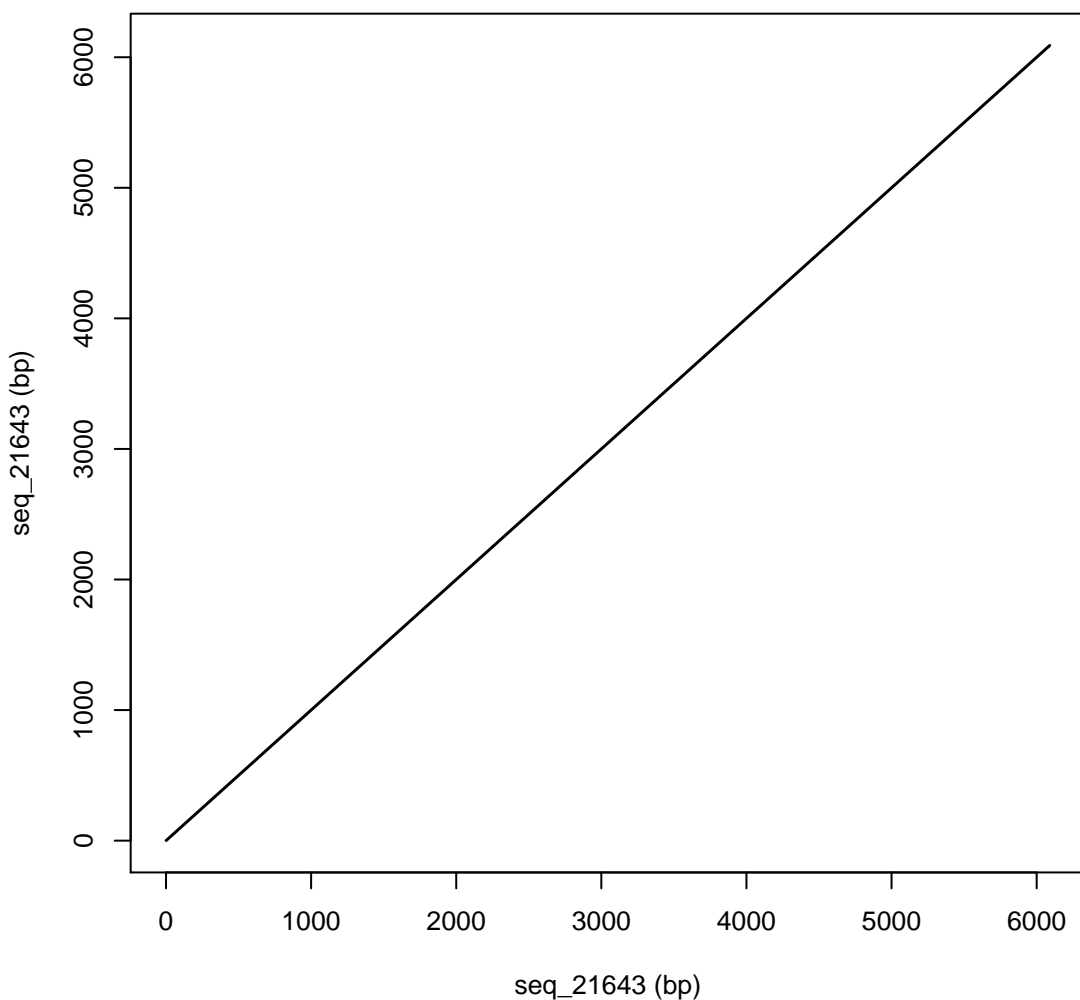
TE: seq_21643
consensus size: 6090bp; fragments: 50990; full length: 1 (>=5481bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

