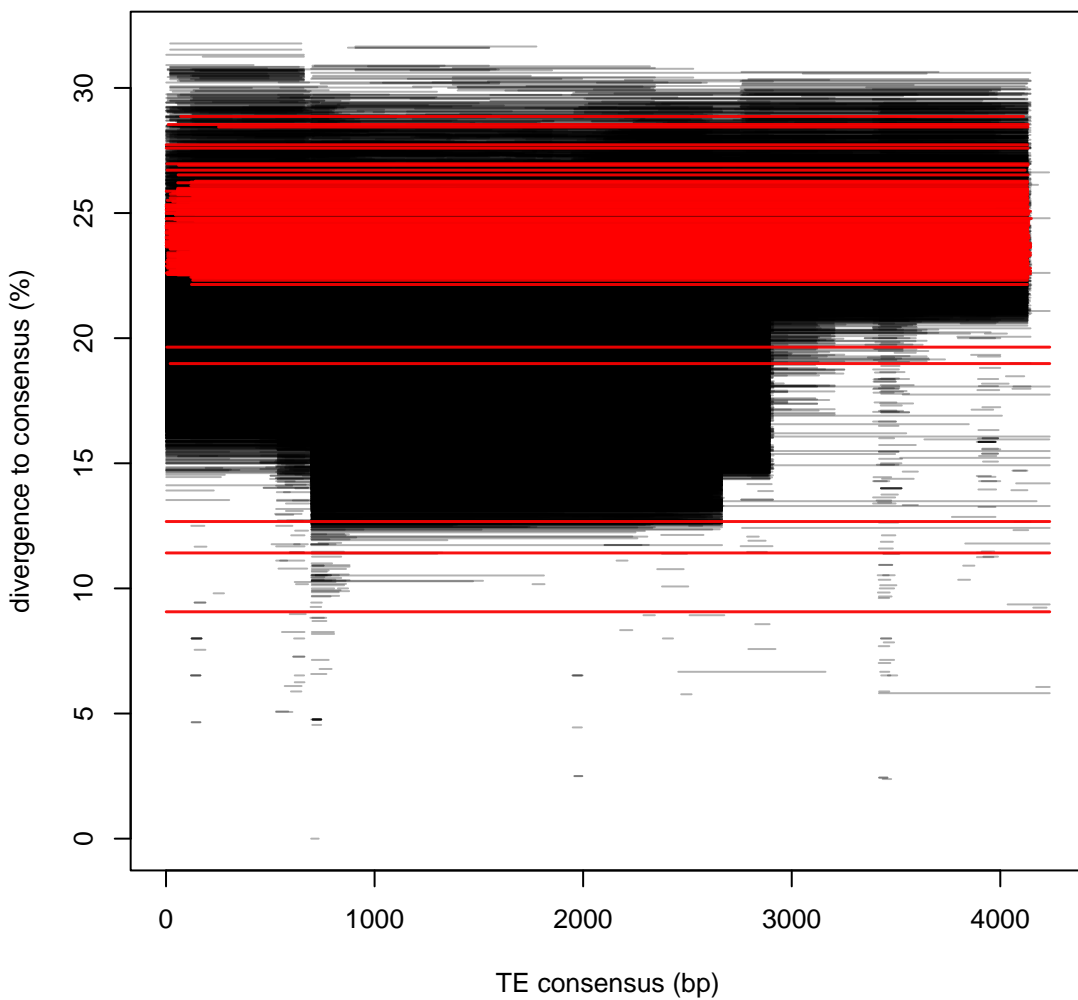
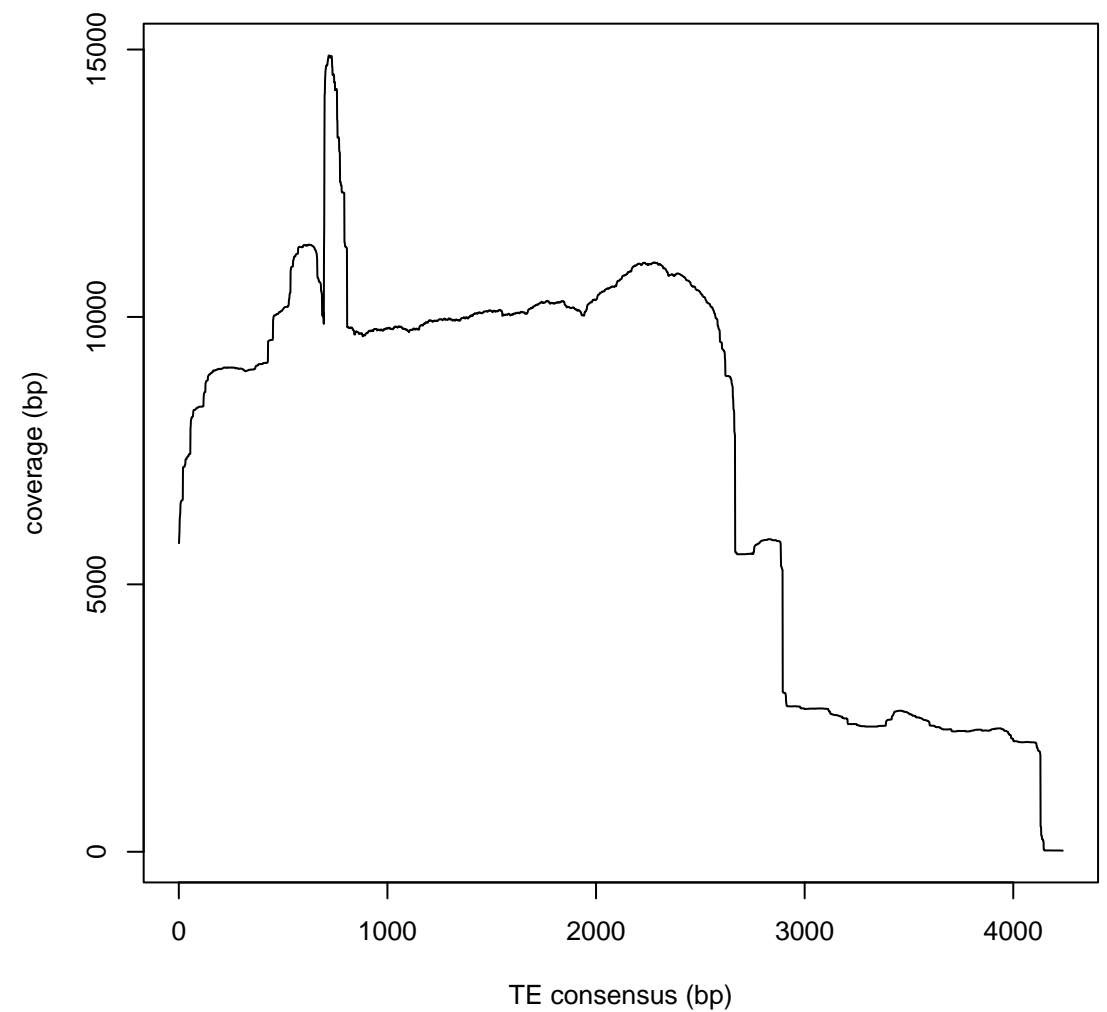


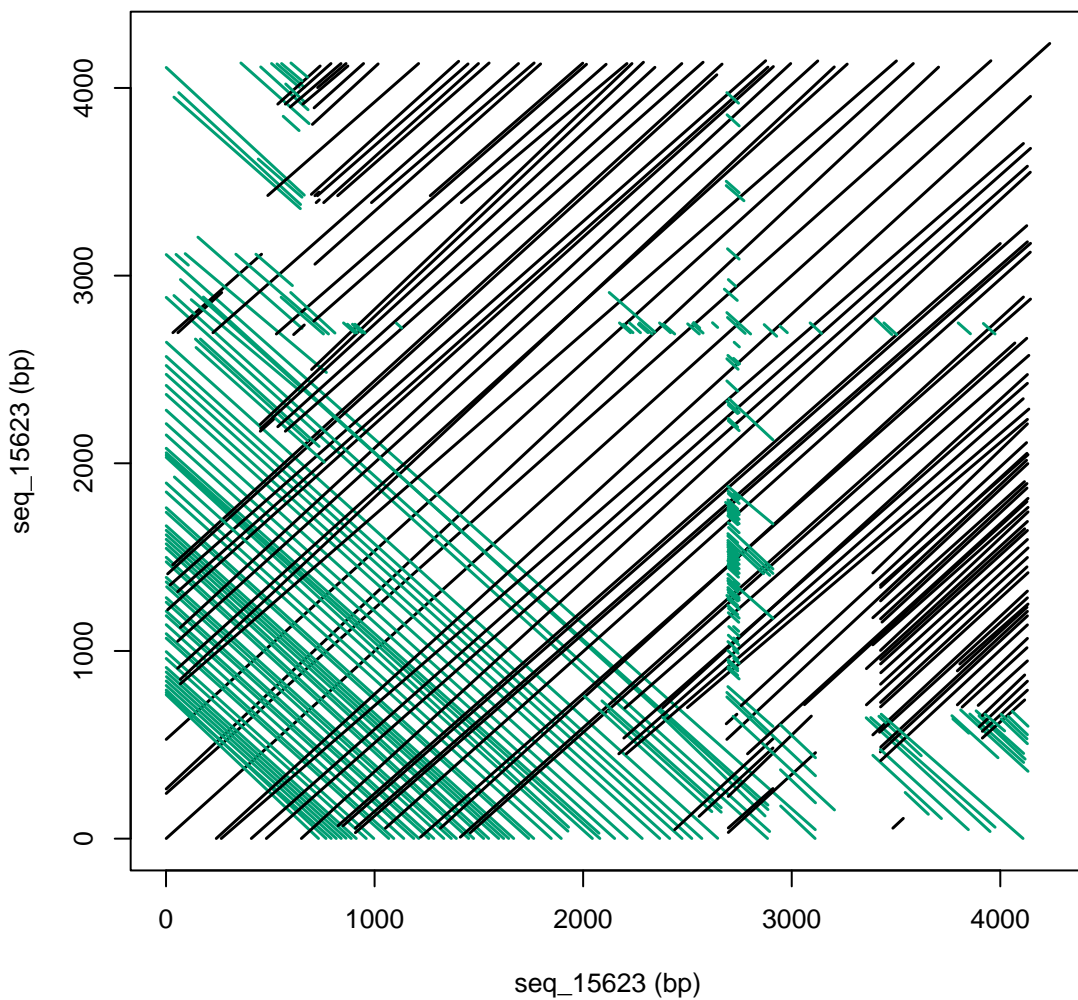
TE: seq_15623
consensus size: 4237bp; fragments: 22246; full length: 237 (>=3813.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

