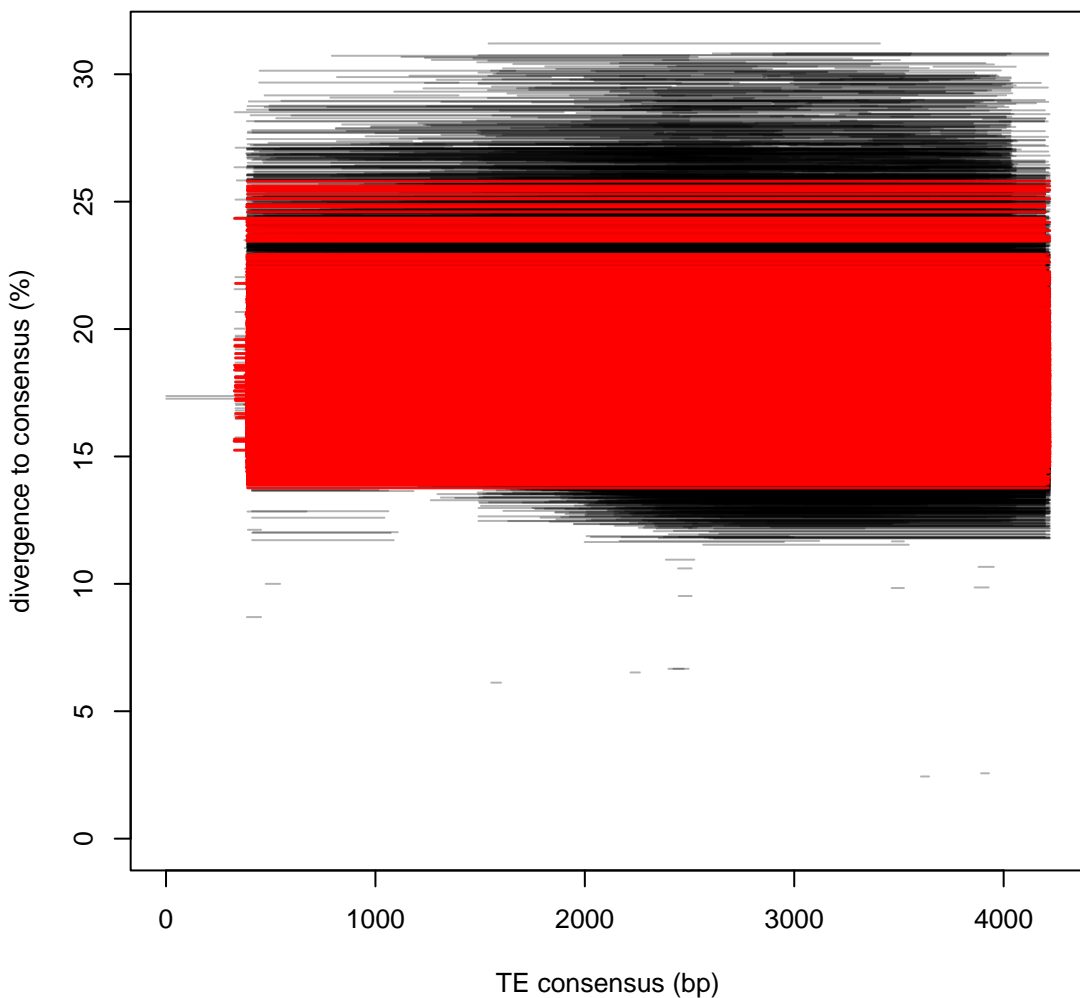
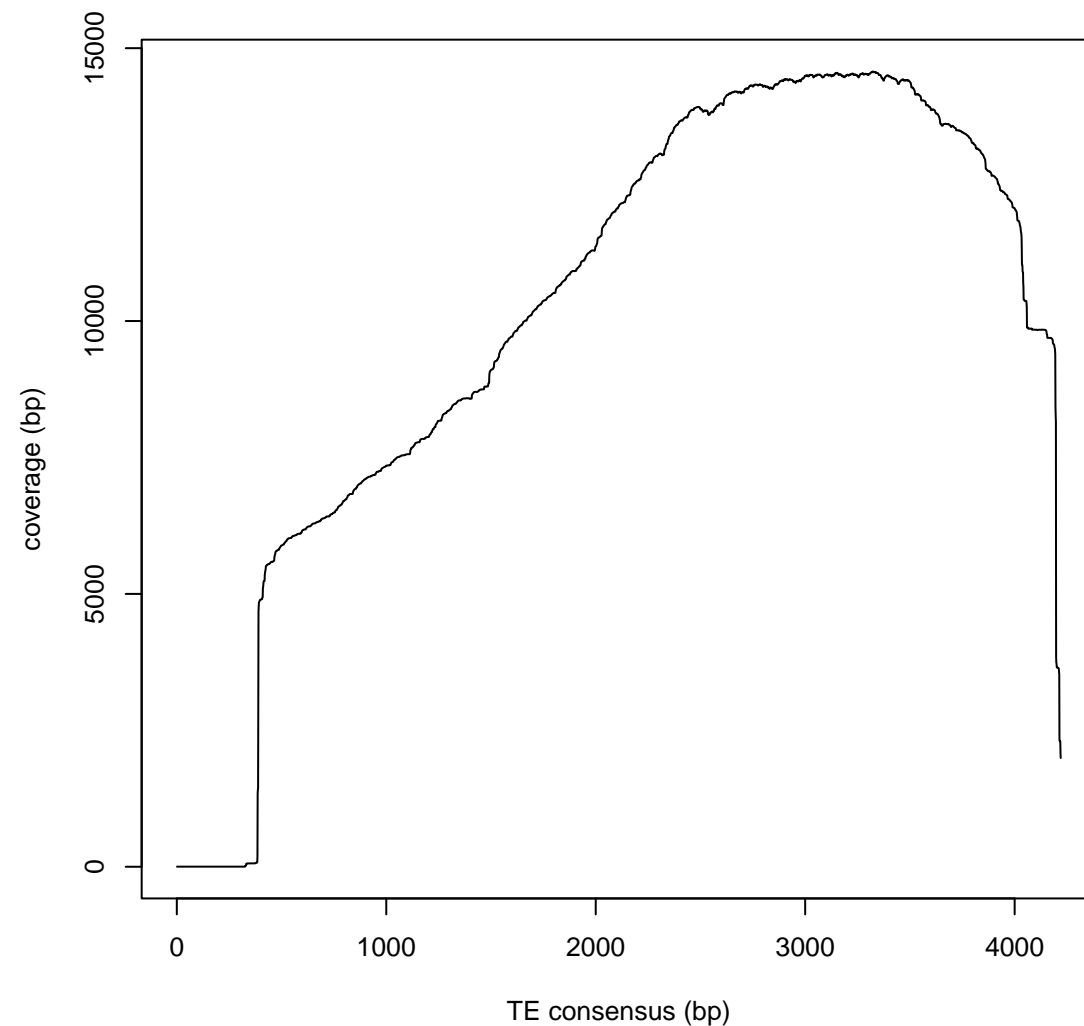


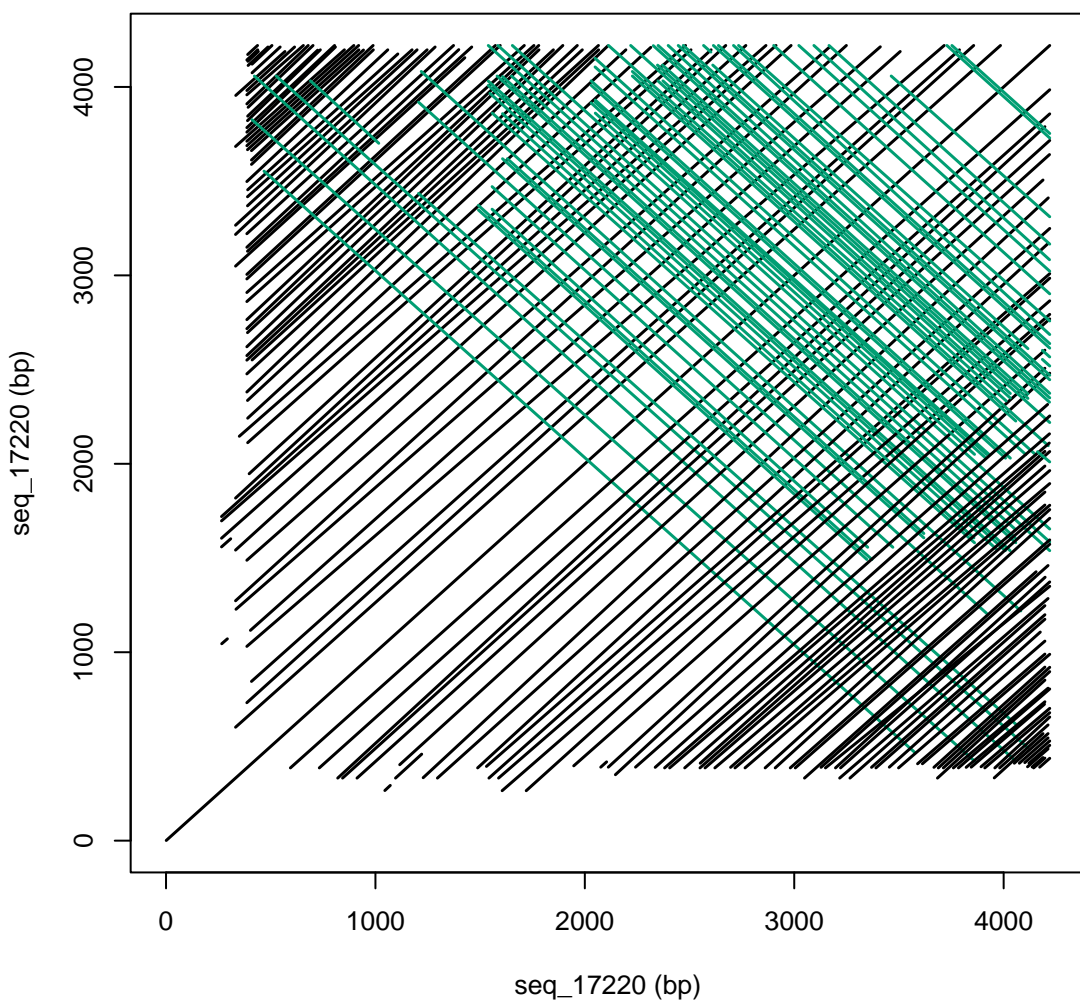
TE: seq_17220
consensus size: 4220bp; fragments: 18203; full length: 2507 (>=3798bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

