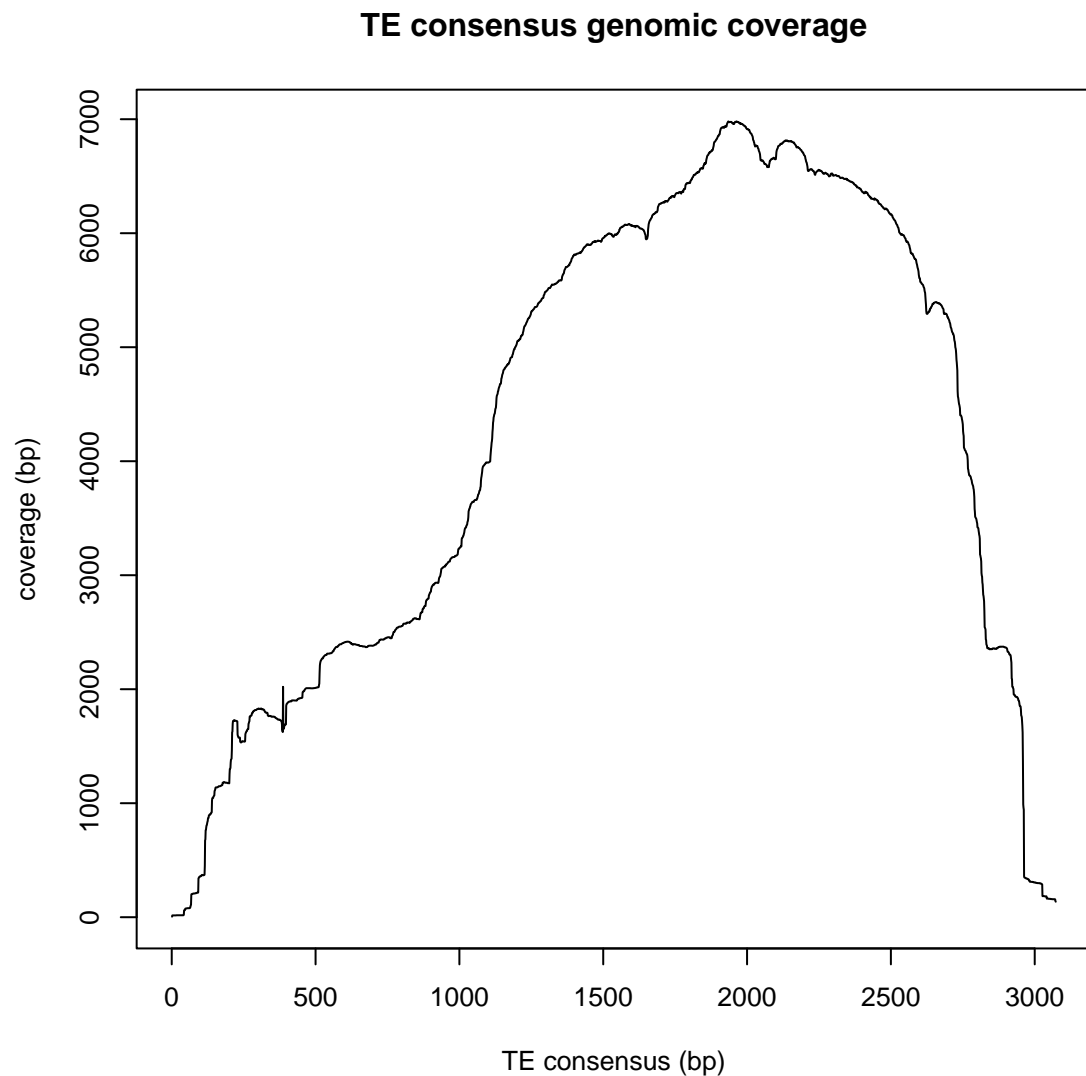
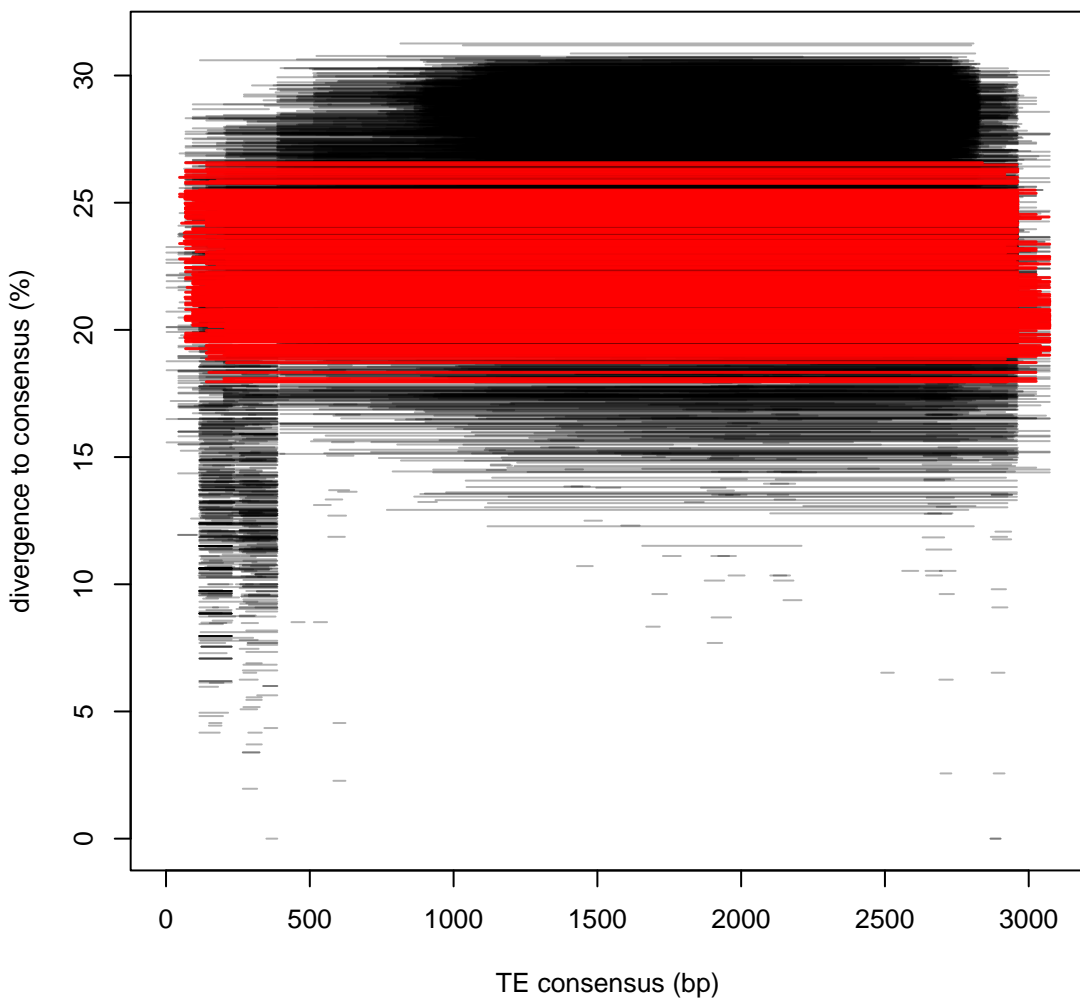
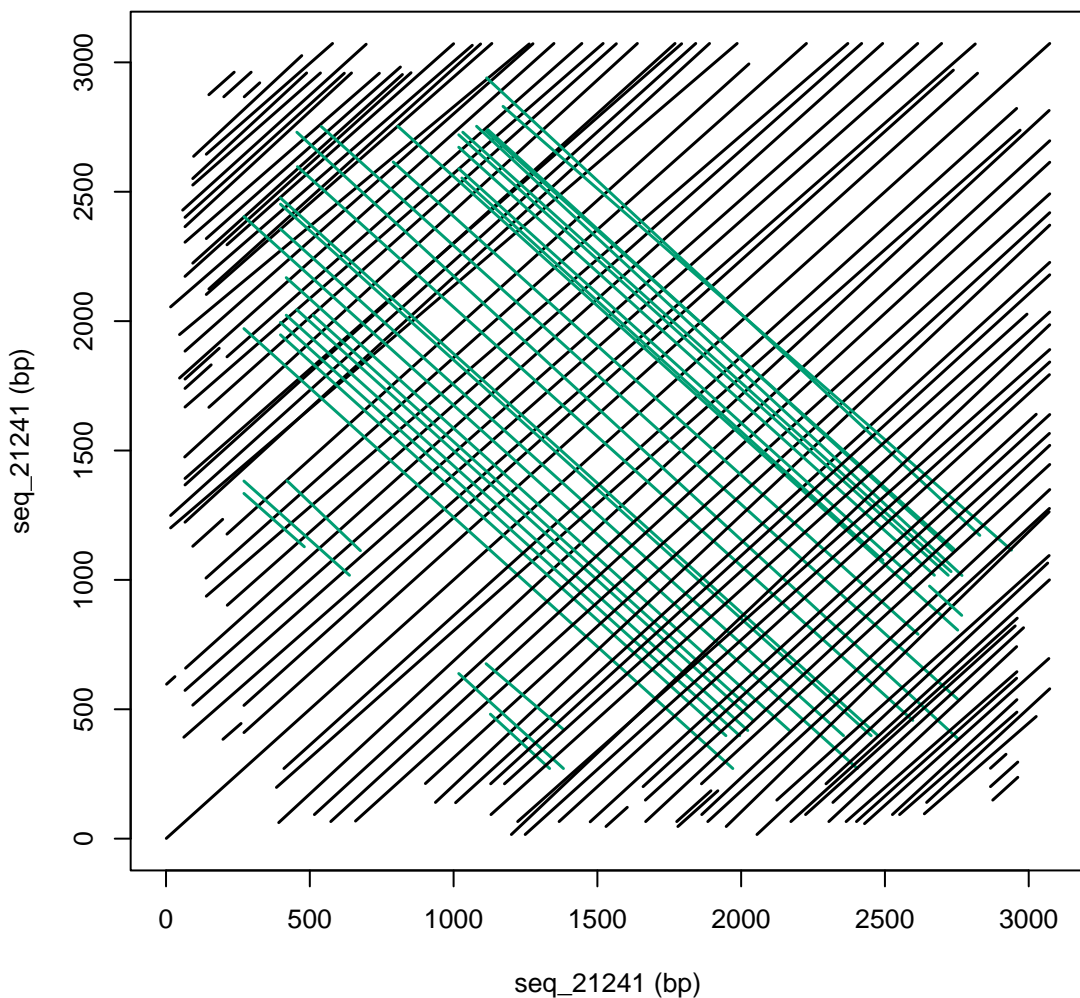


TE: seq_21241
consensus size: 3073bp; fragments: 9948; full length: 299 (>=2765.7bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

