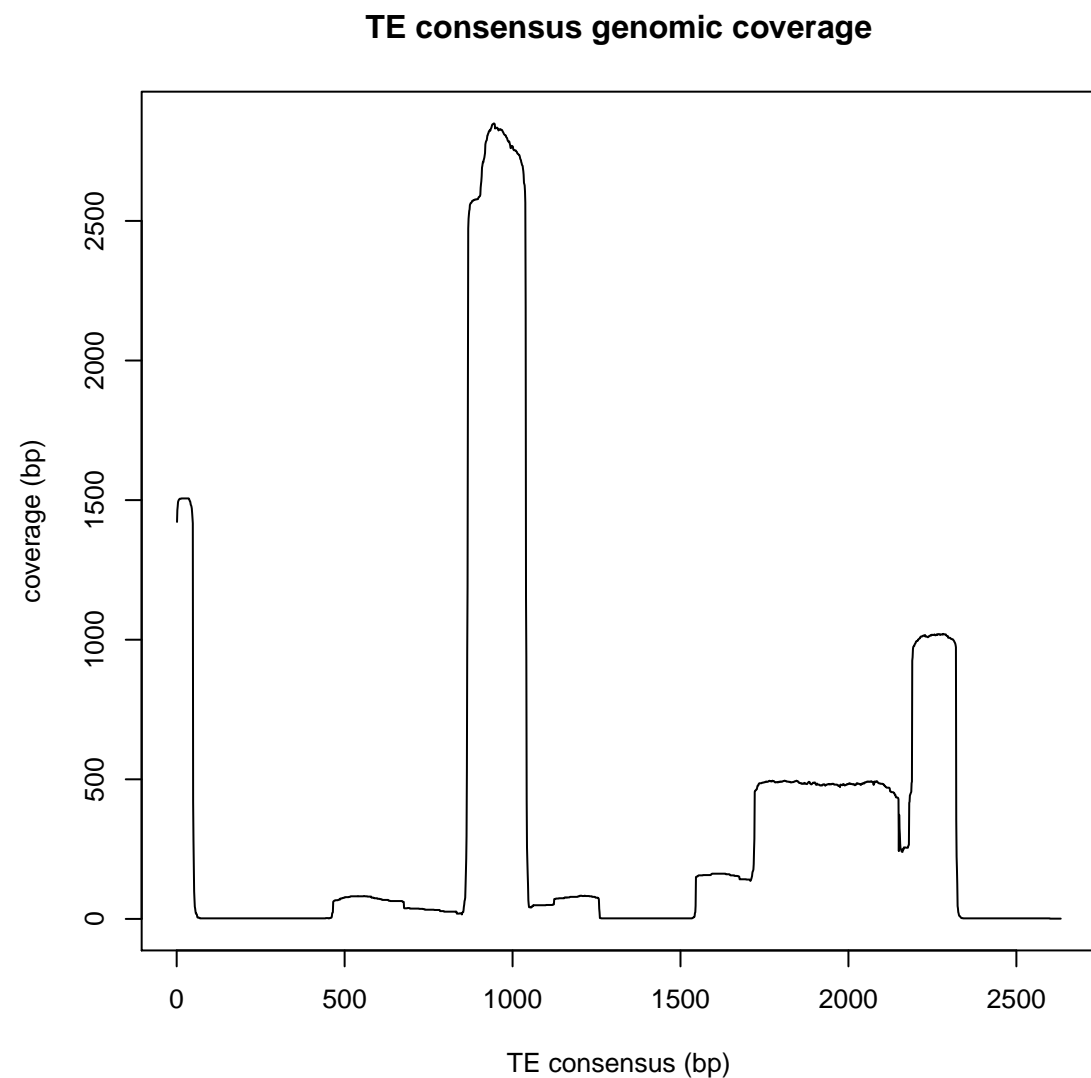
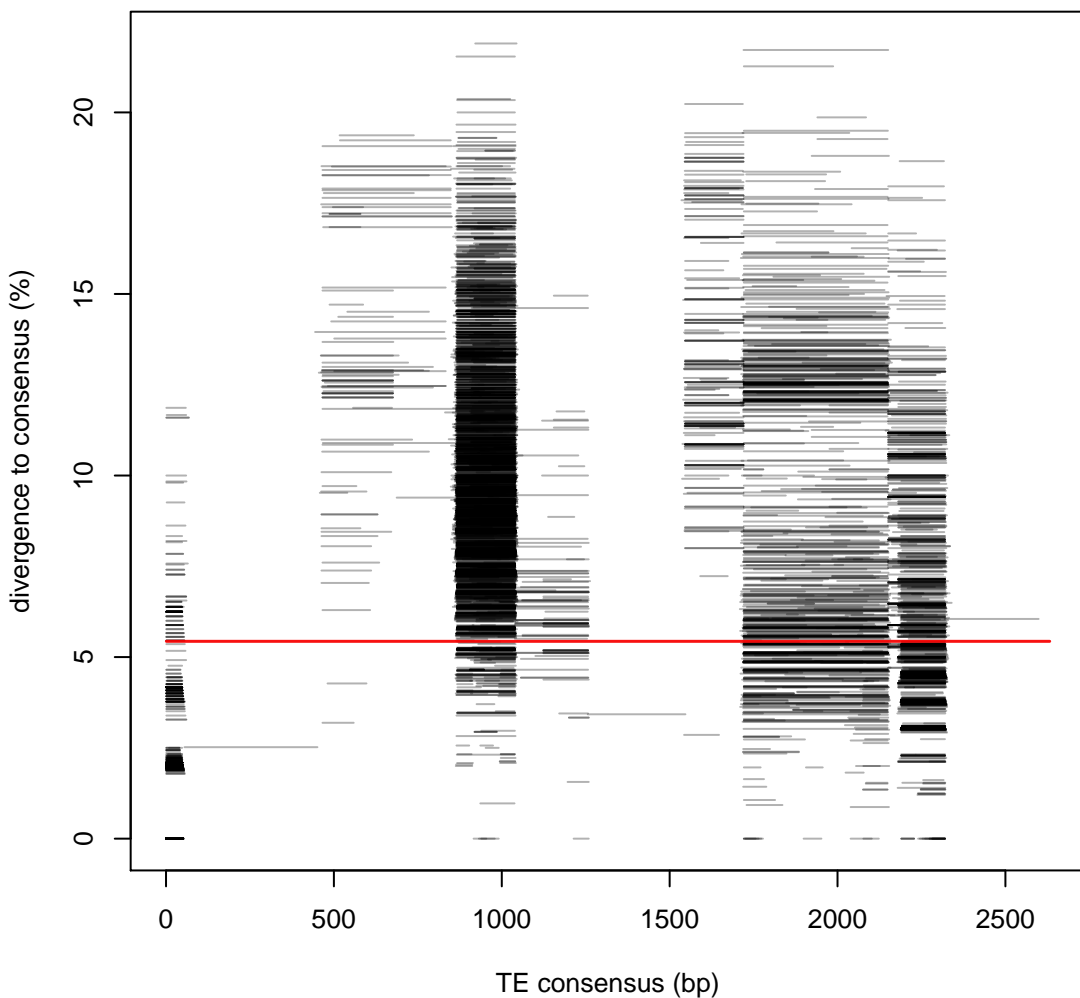
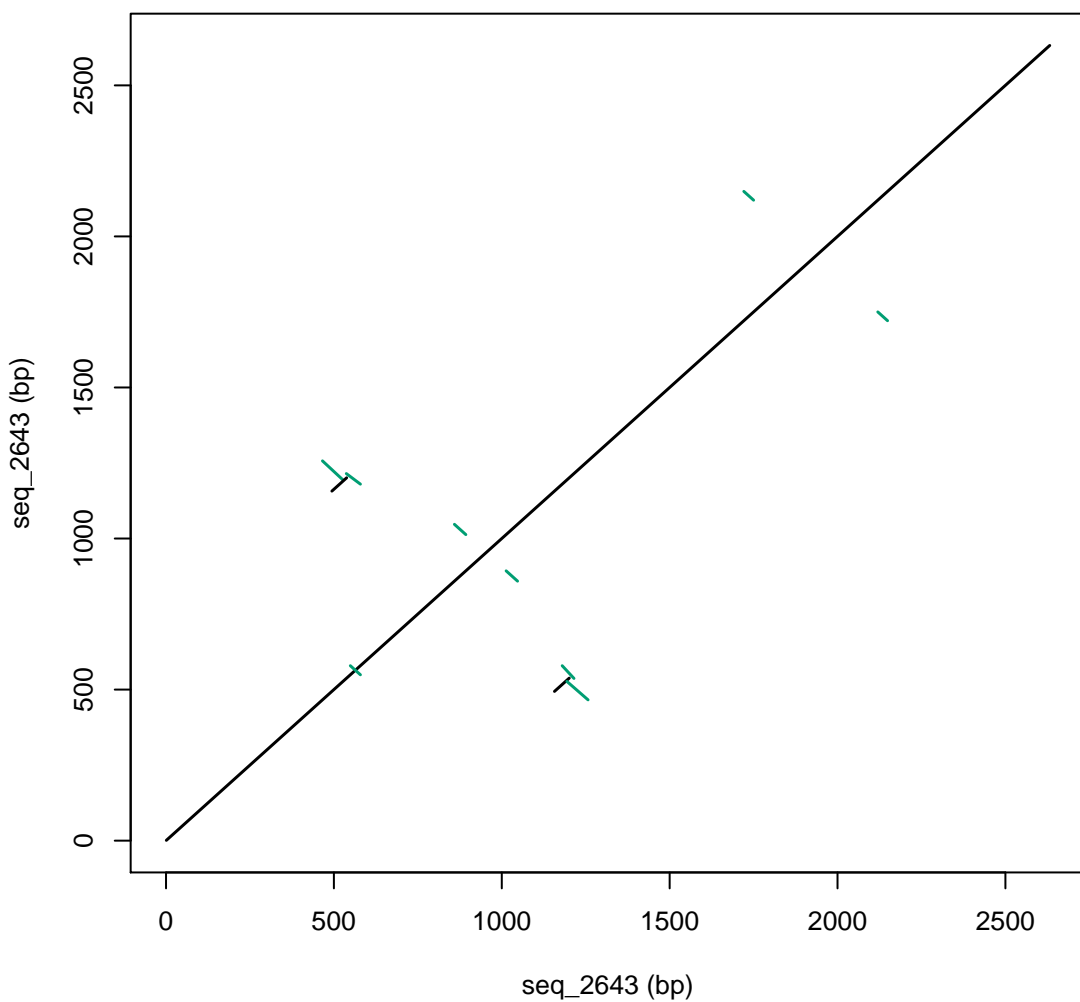


TE: seq_2643
consensus size: 2632bp; fragments: 6501; full length: 1 (>=2368.8bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

