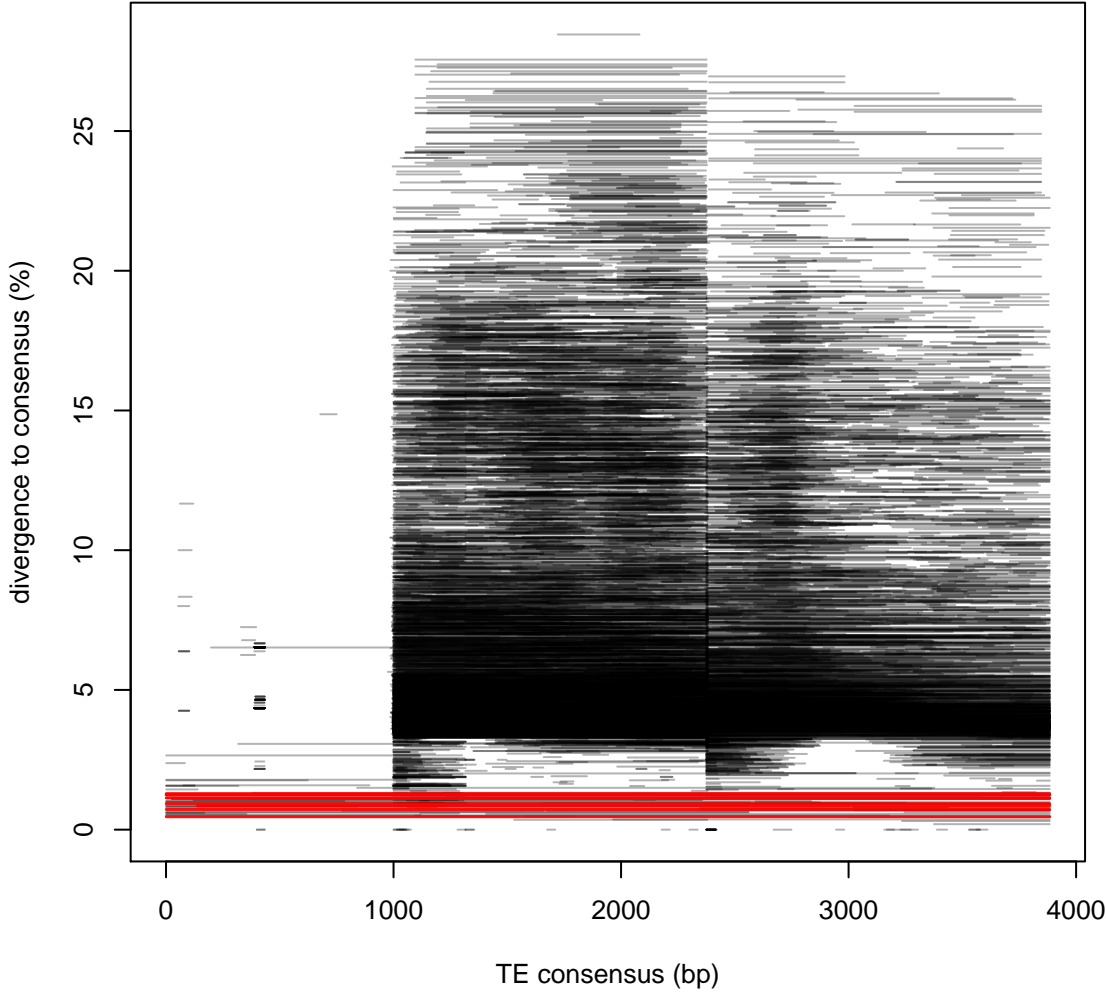
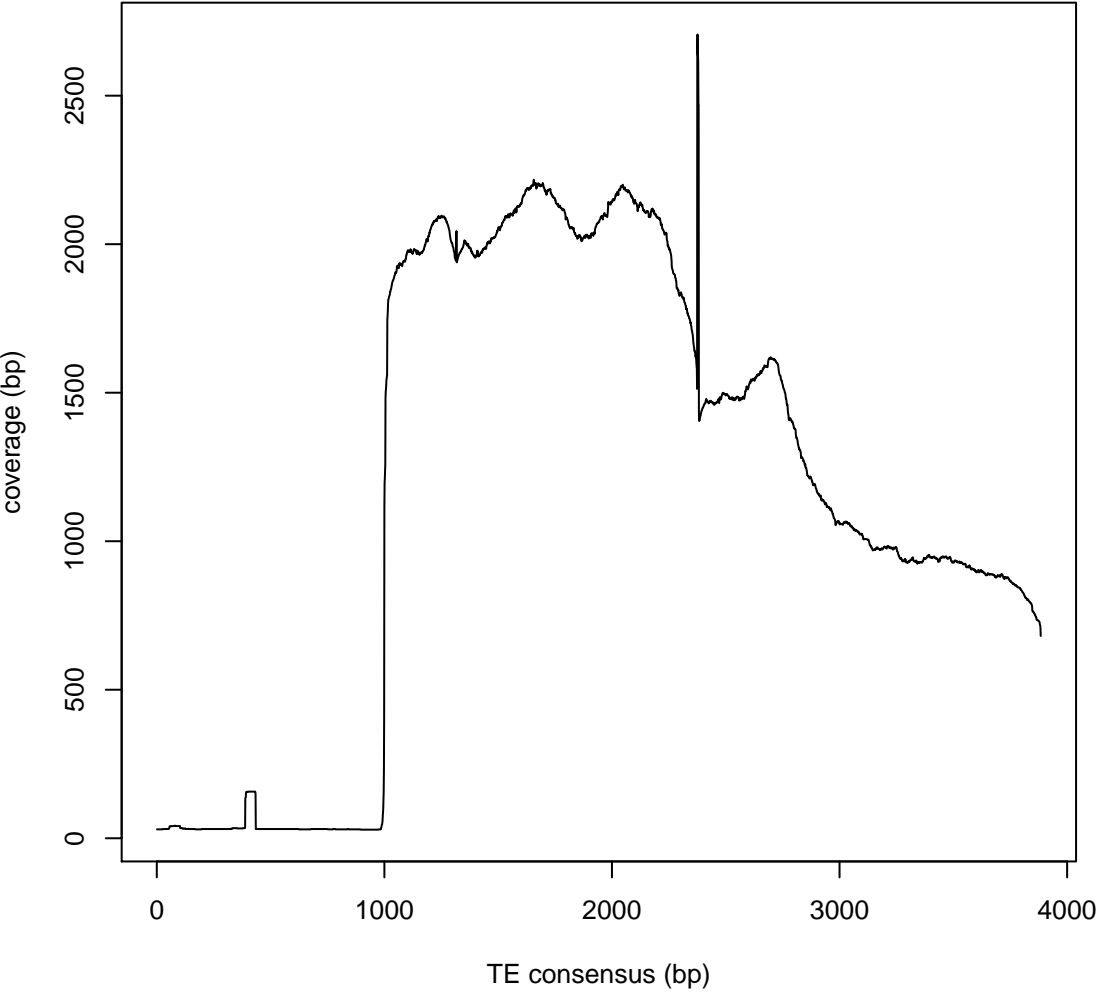


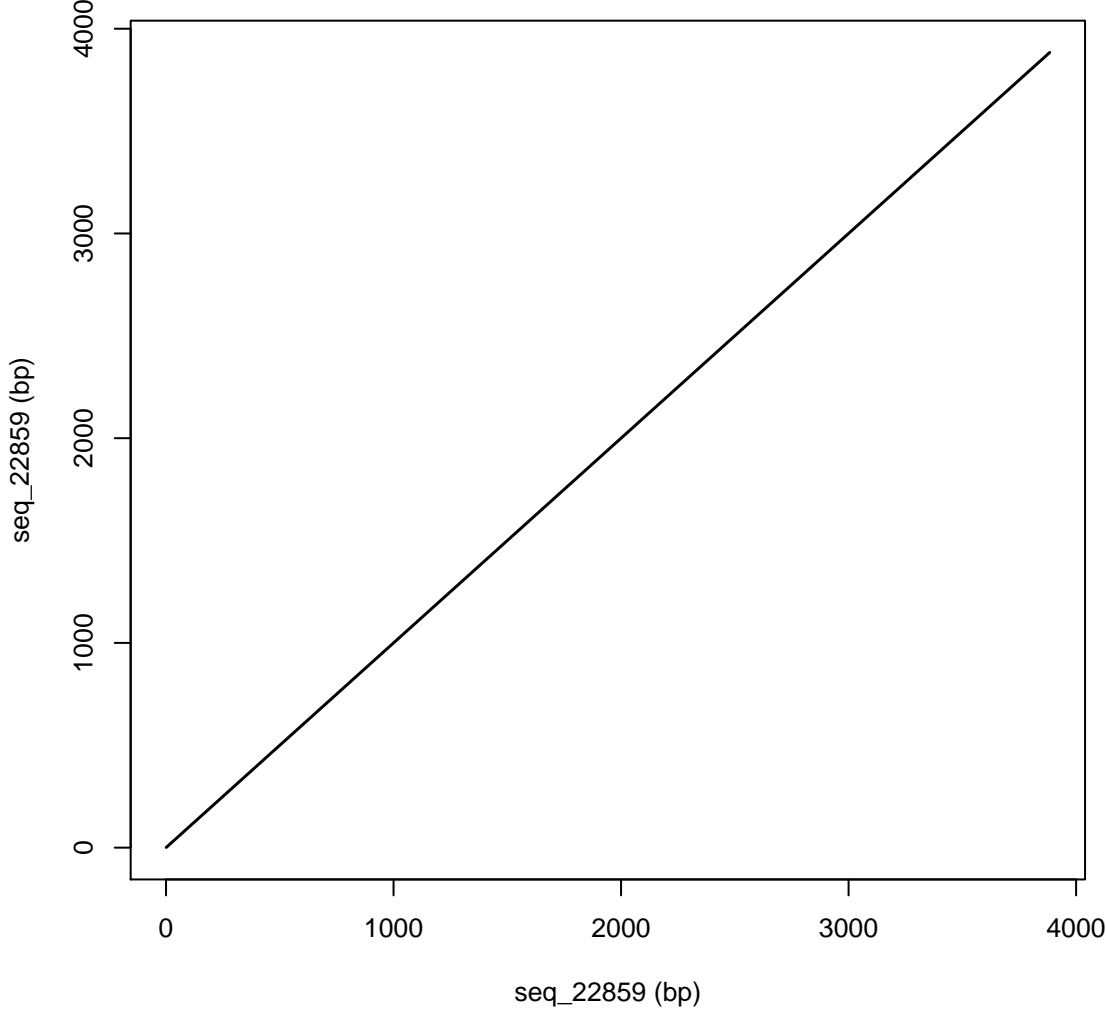
TE: seq_22859
consensus size: 3884bp; fragments: 8097; full length: 9 (>=3495.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

