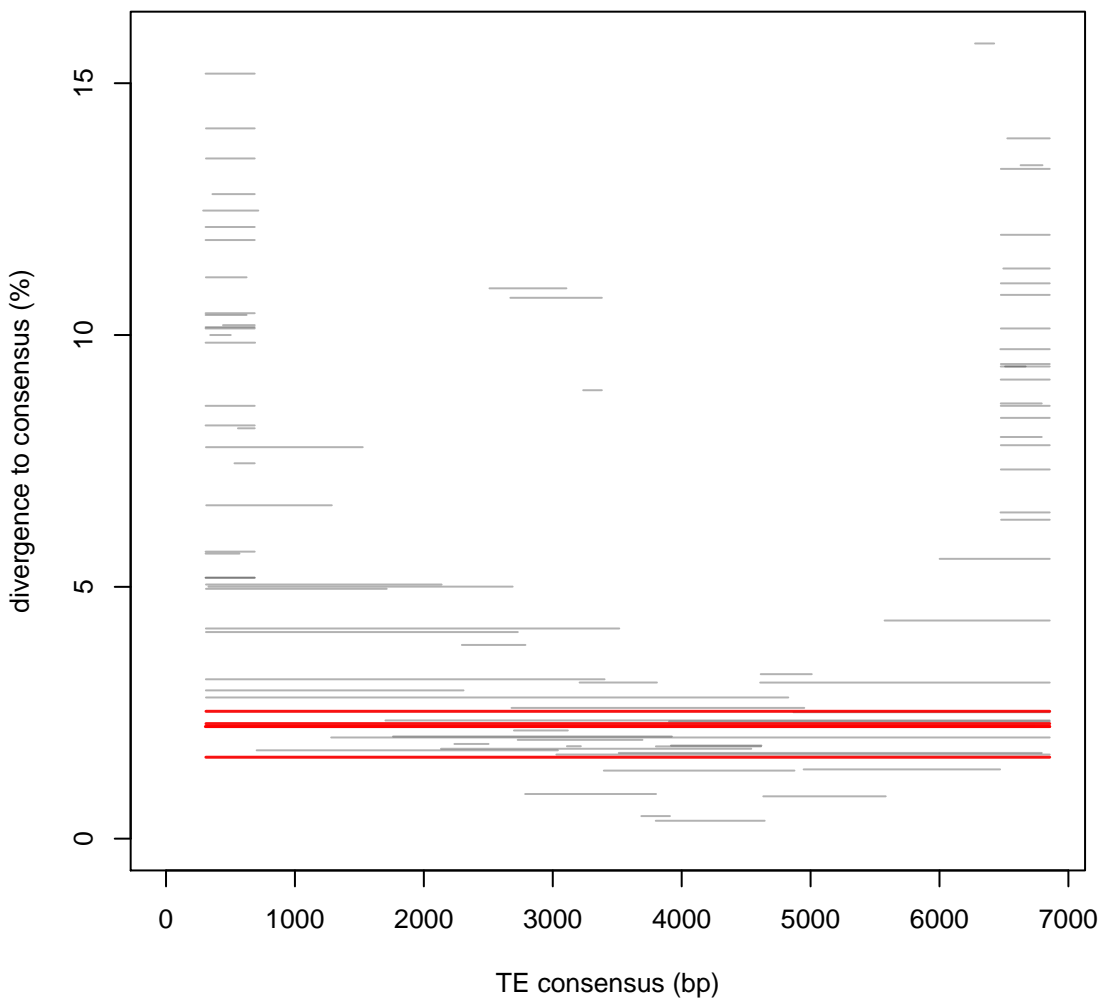
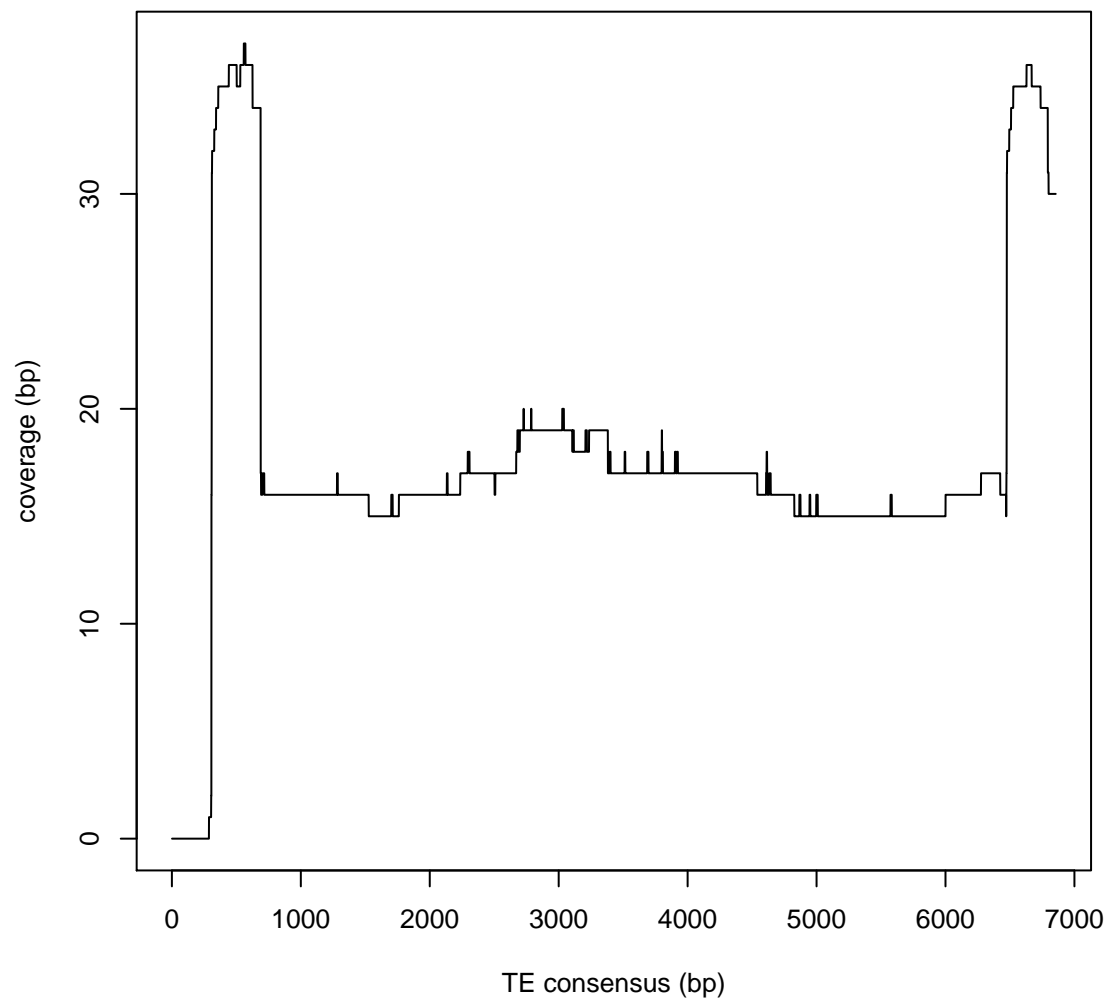


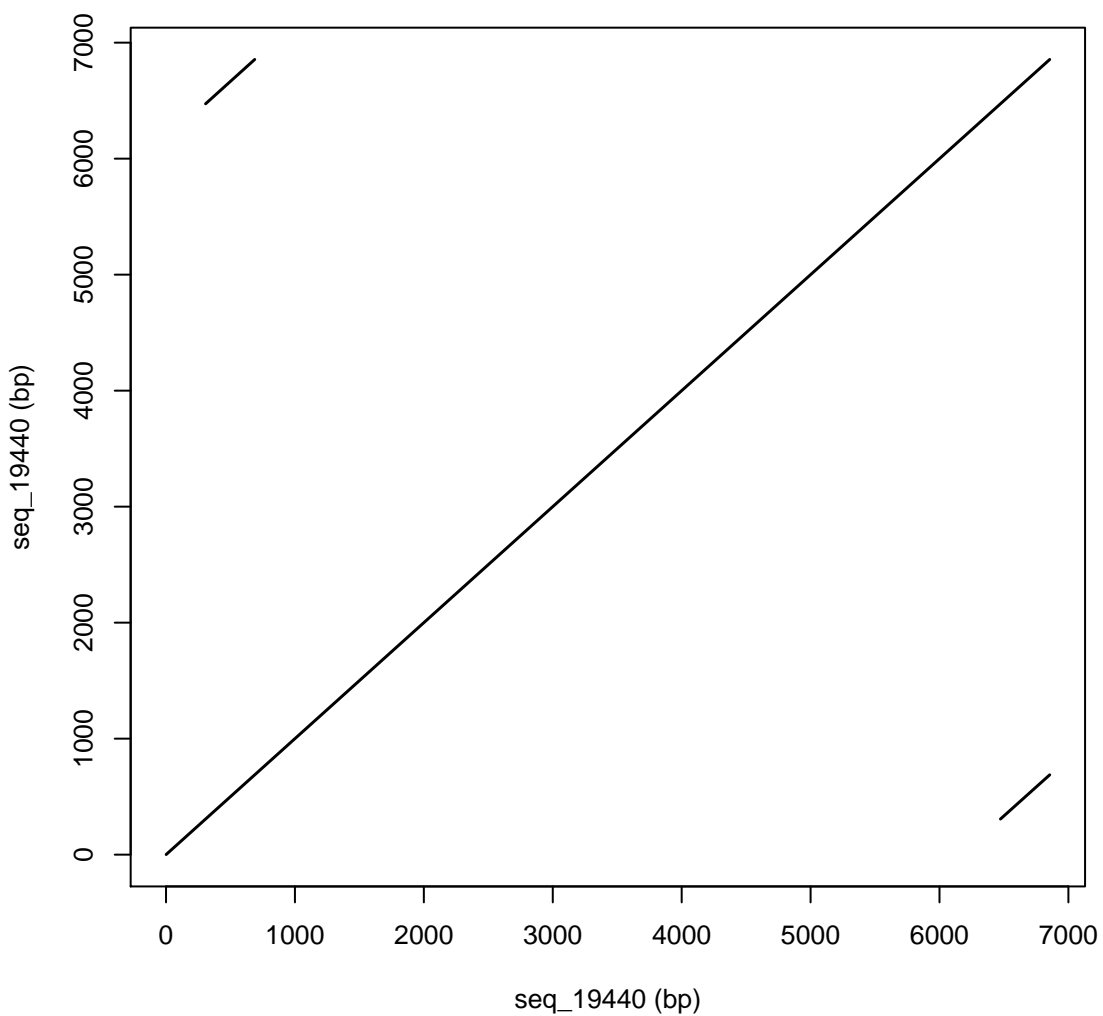
TE: seq_19440
consensus size: 6855bp; fragments: 92; full length: 5 ($\geq 6169.5\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

