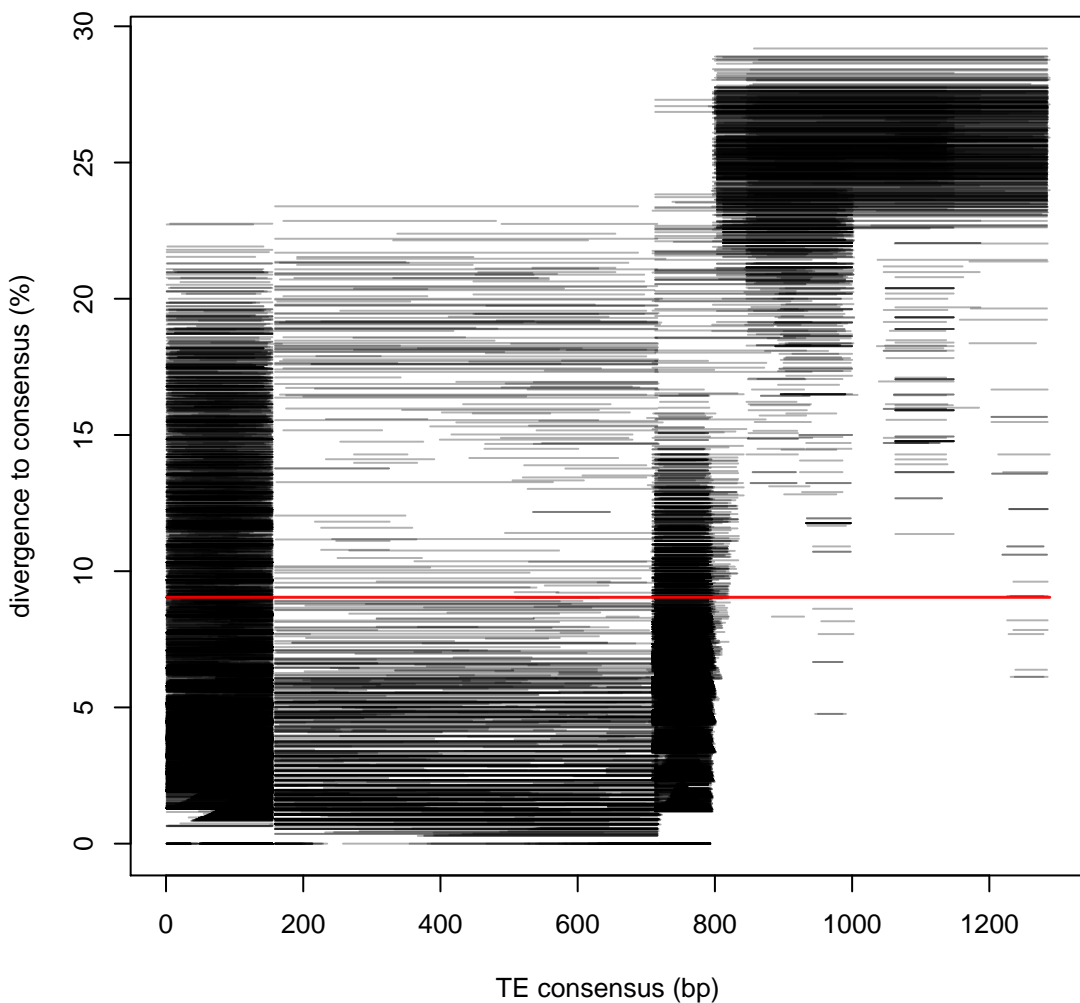
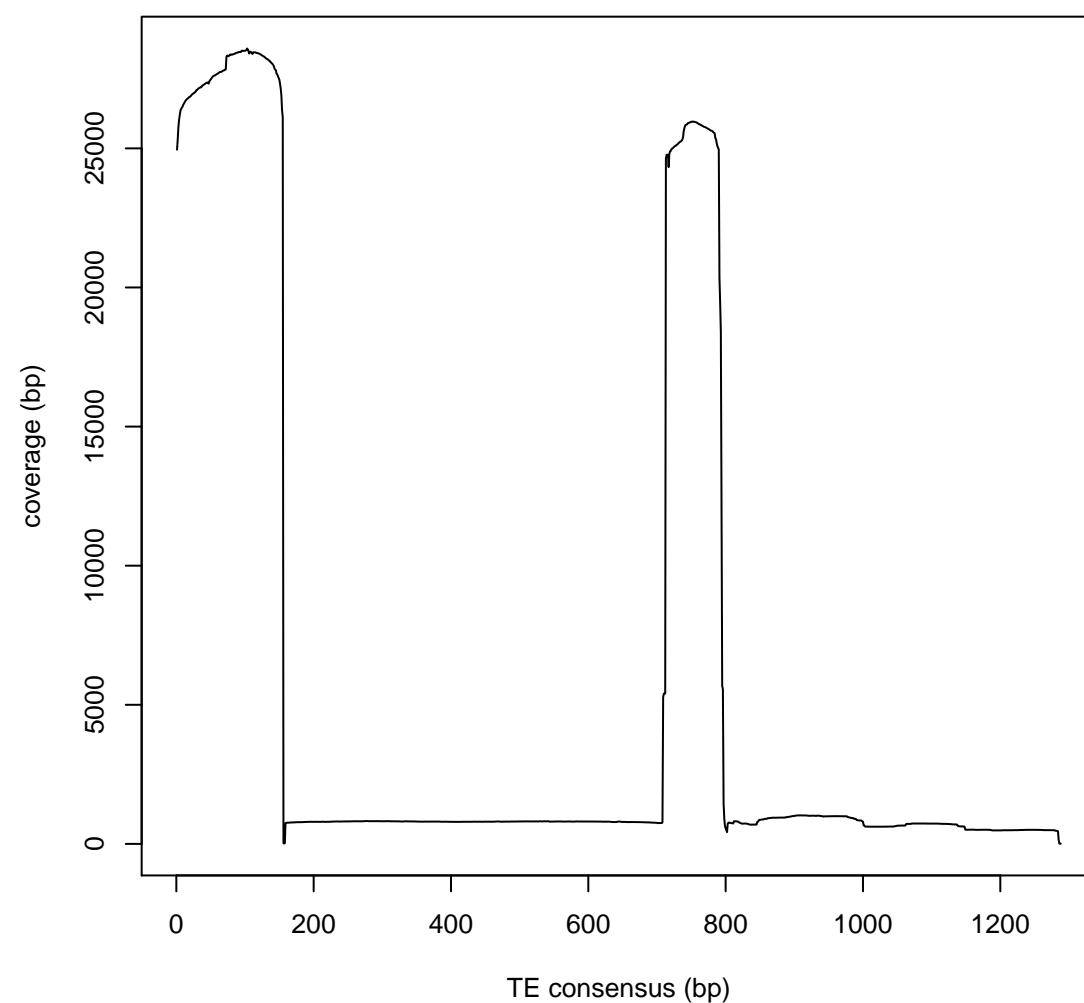


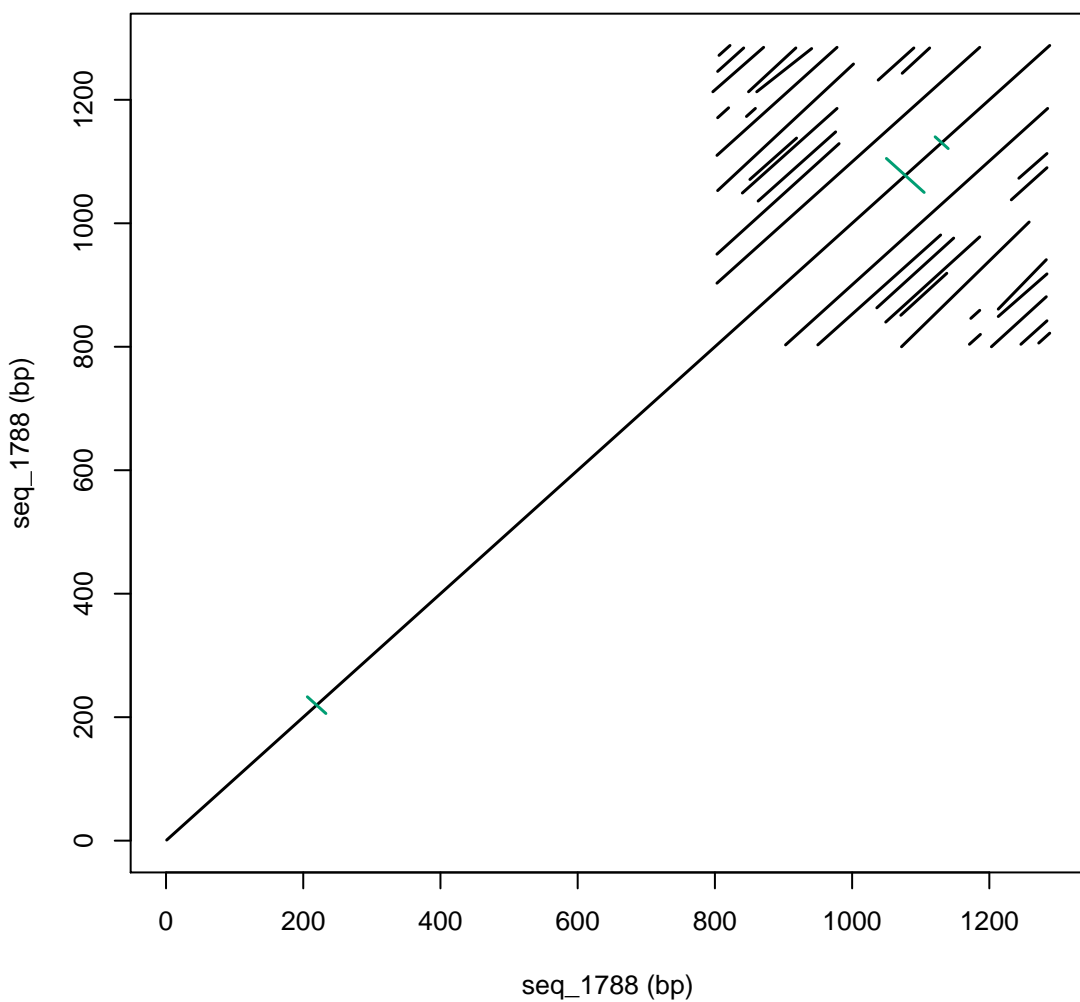
TE: seq_1788
consensus size: 1288bp; fragments: 57882; full length: 1 (>=1159.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

