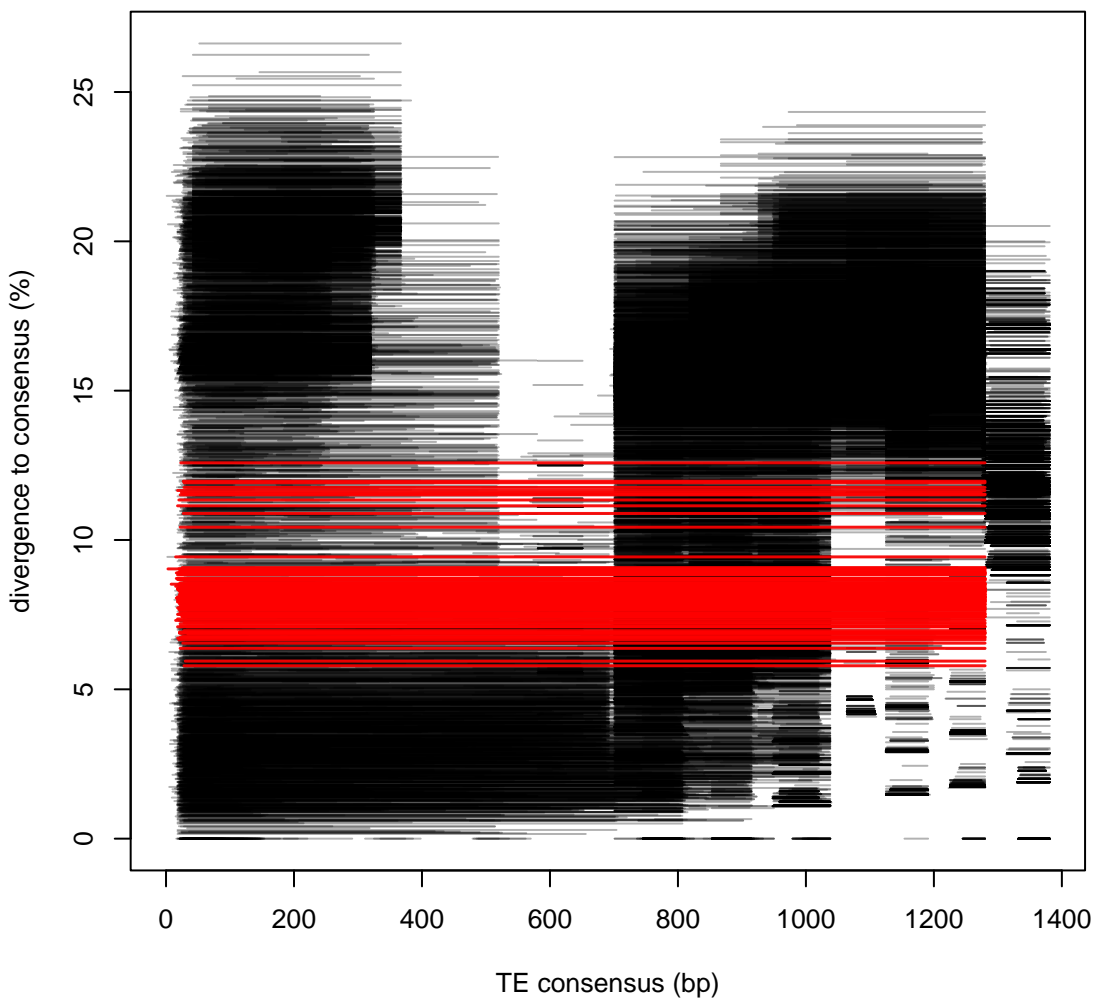
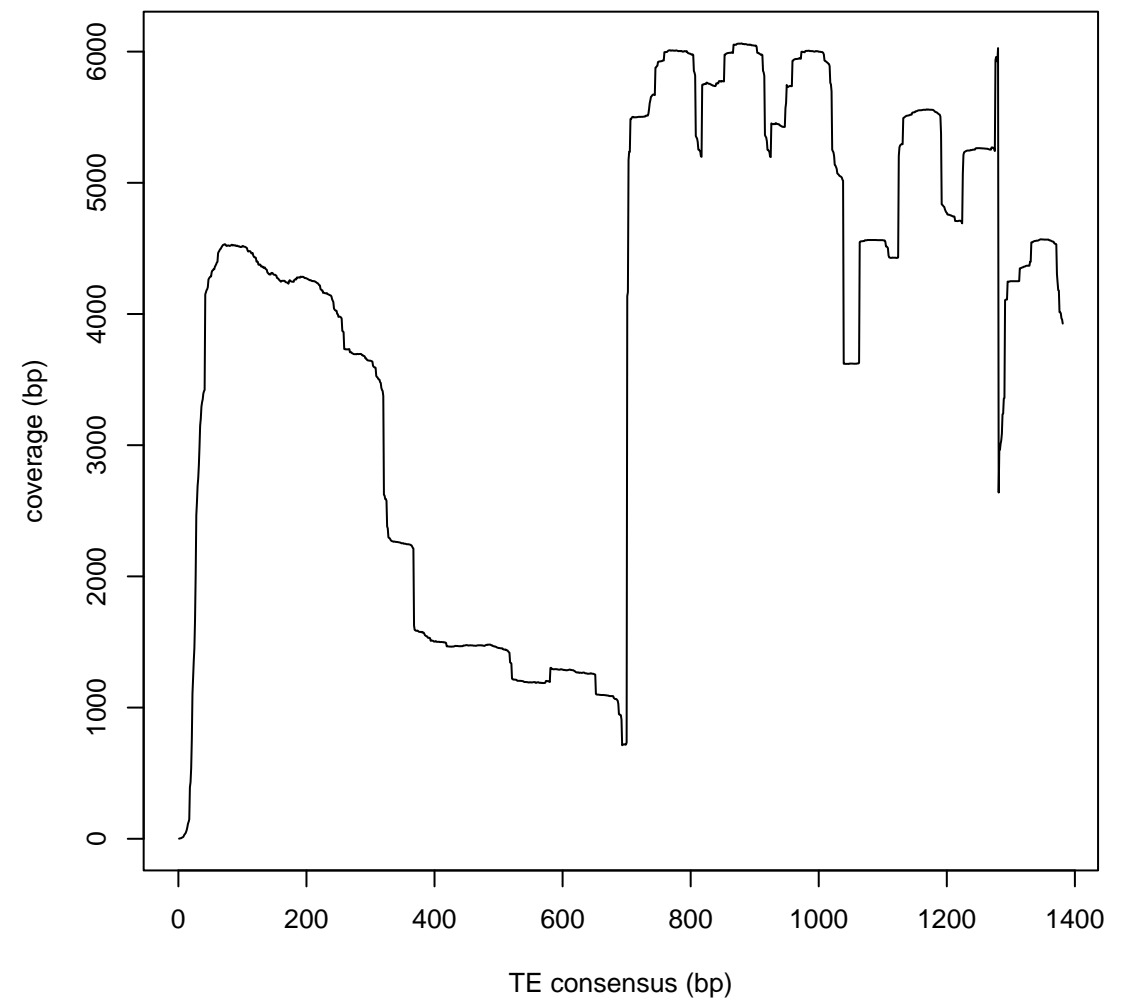


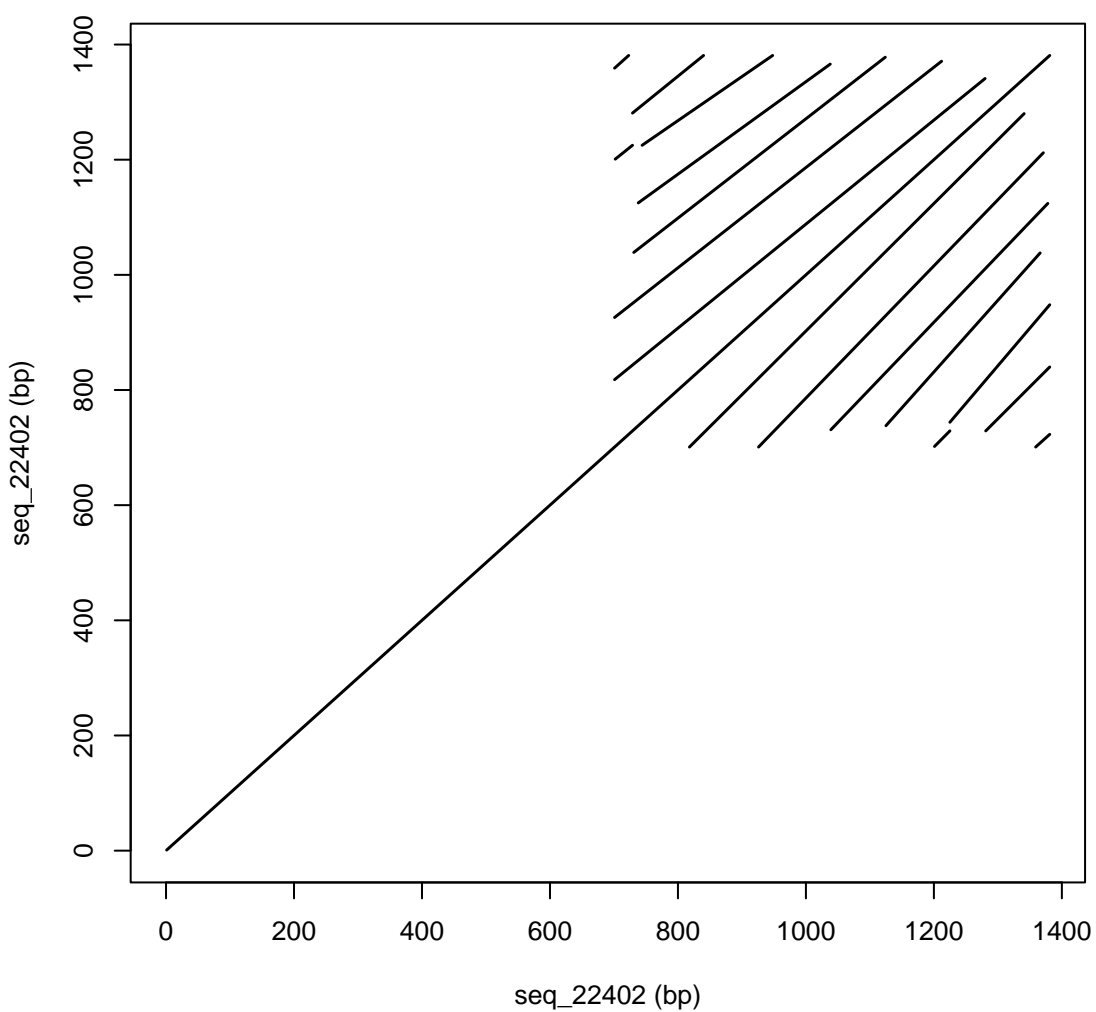
TE: seq_22402
consensus size: 1381bp; fragments: 20108; full length: 120 (≥ 1242.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

