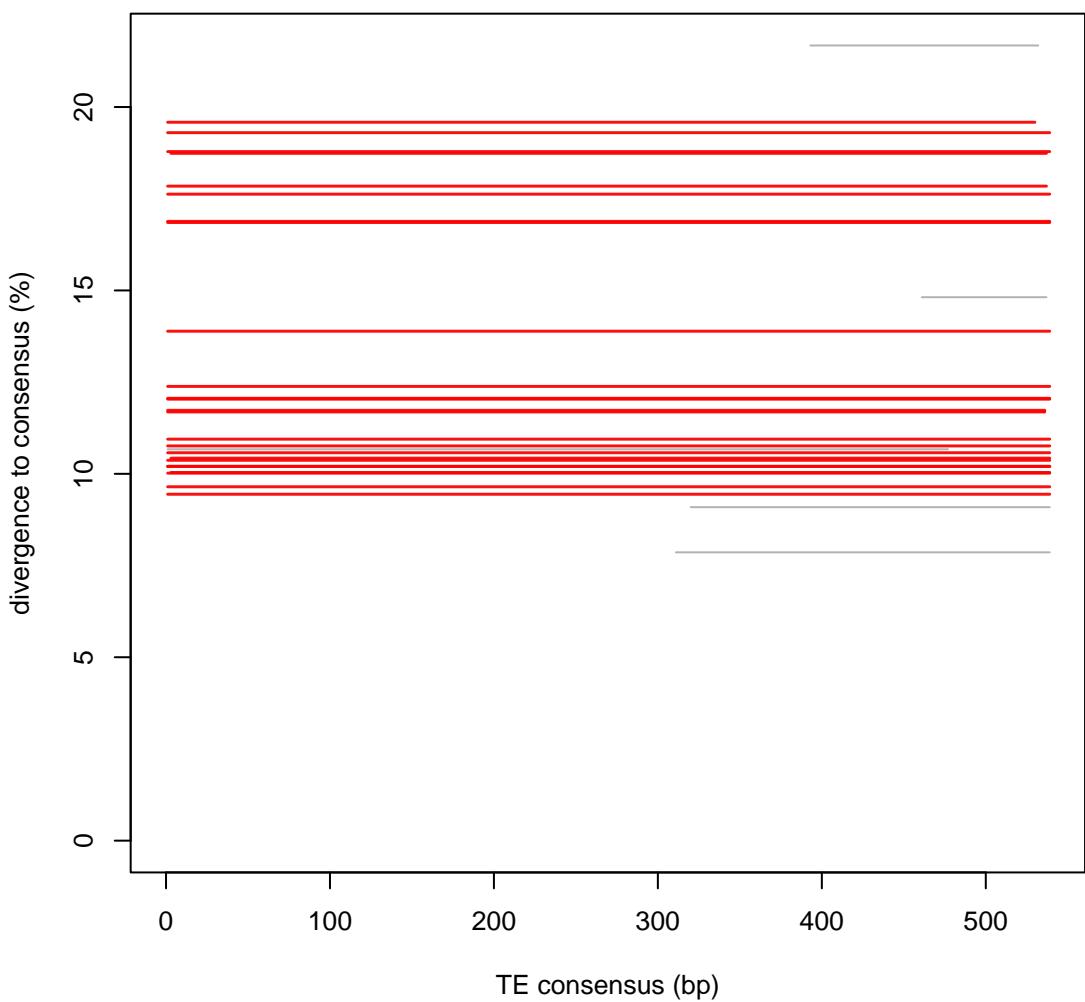
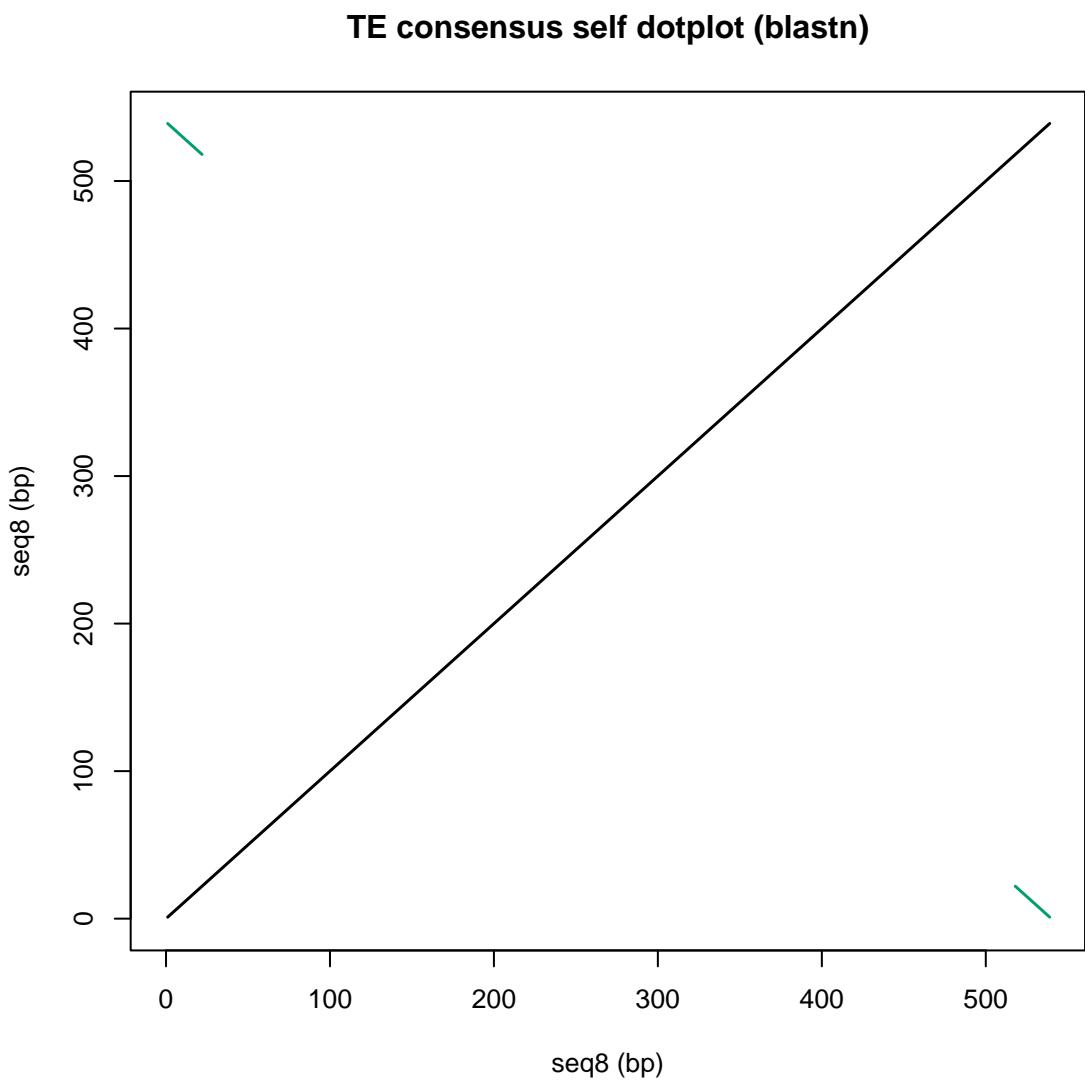
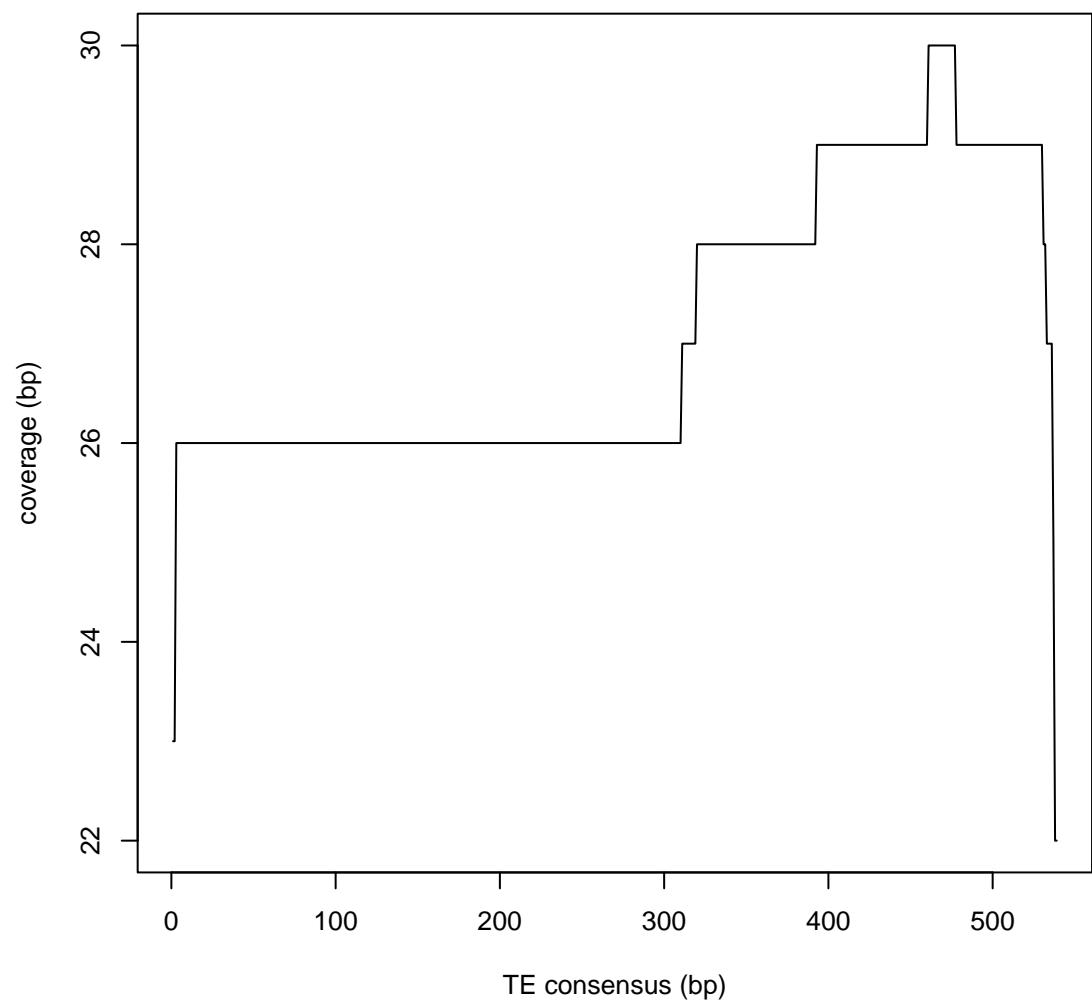


TE: seq8
consensus size: 539bp; fragments: 30; full length: 25 (≥ 485.1 bp)



TE consensus genomic coverage



no orf >400 bp detected

