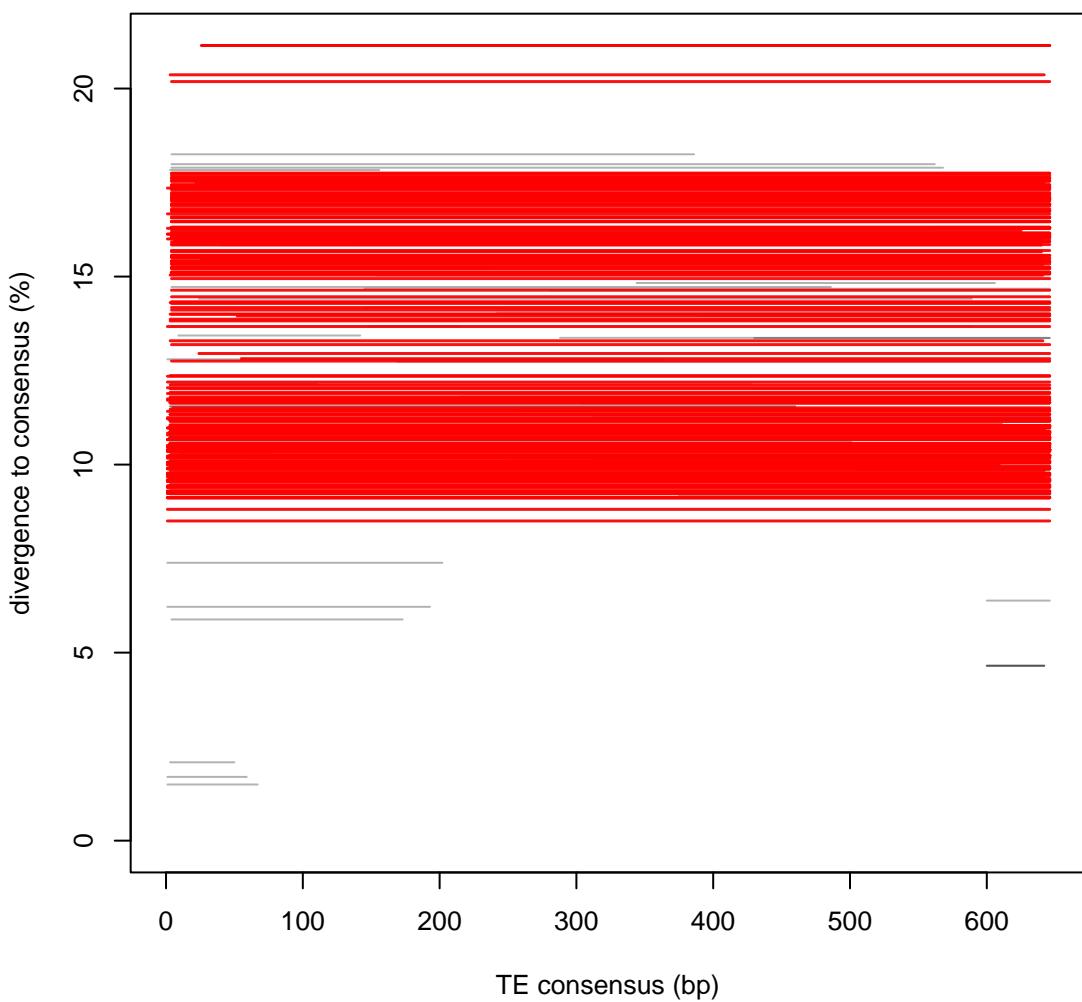
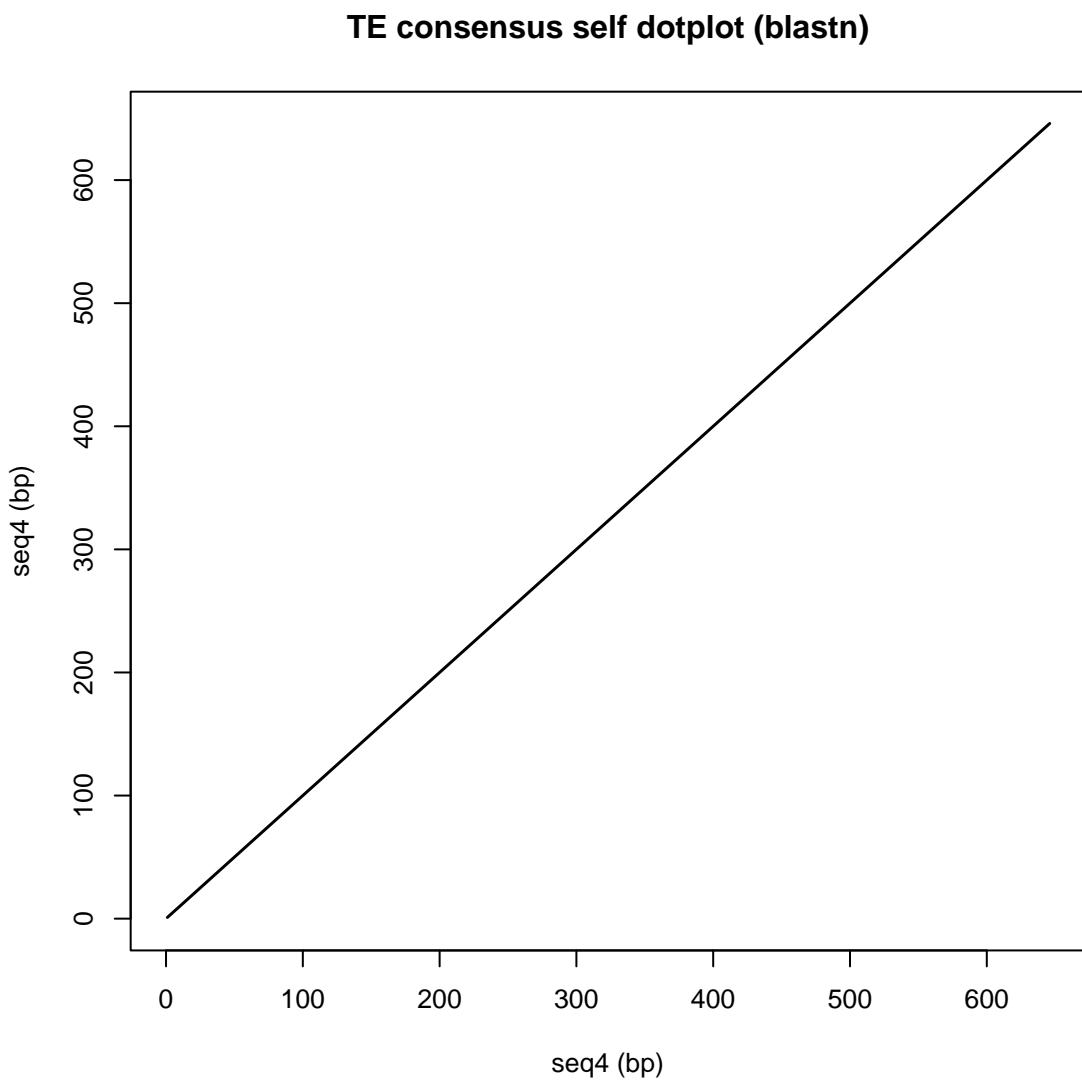
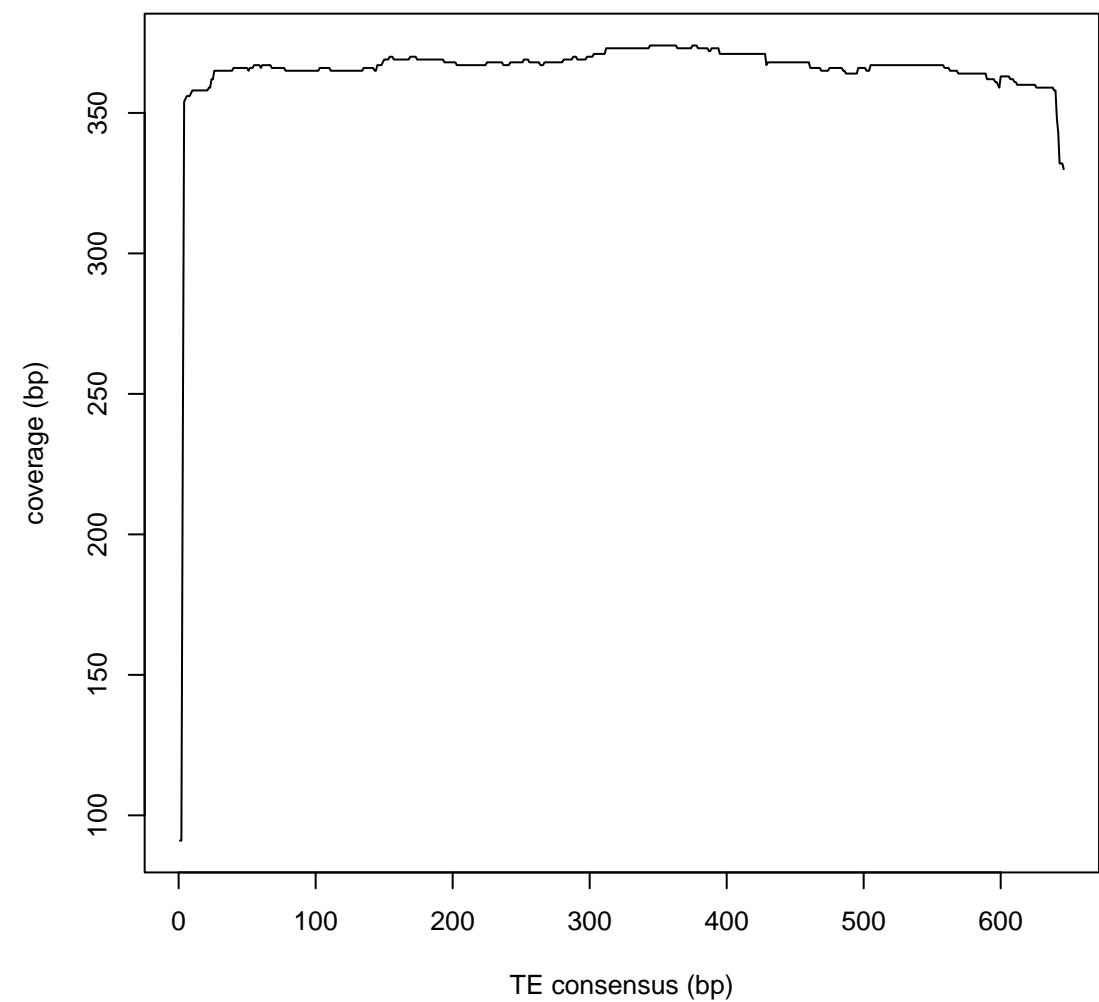


TE: seq4
consensus size: 646bp; fragments: 401; full length: 336 (≥ 581.4 bp)



TE consensus genomic coverage



no orf >400 bp detected