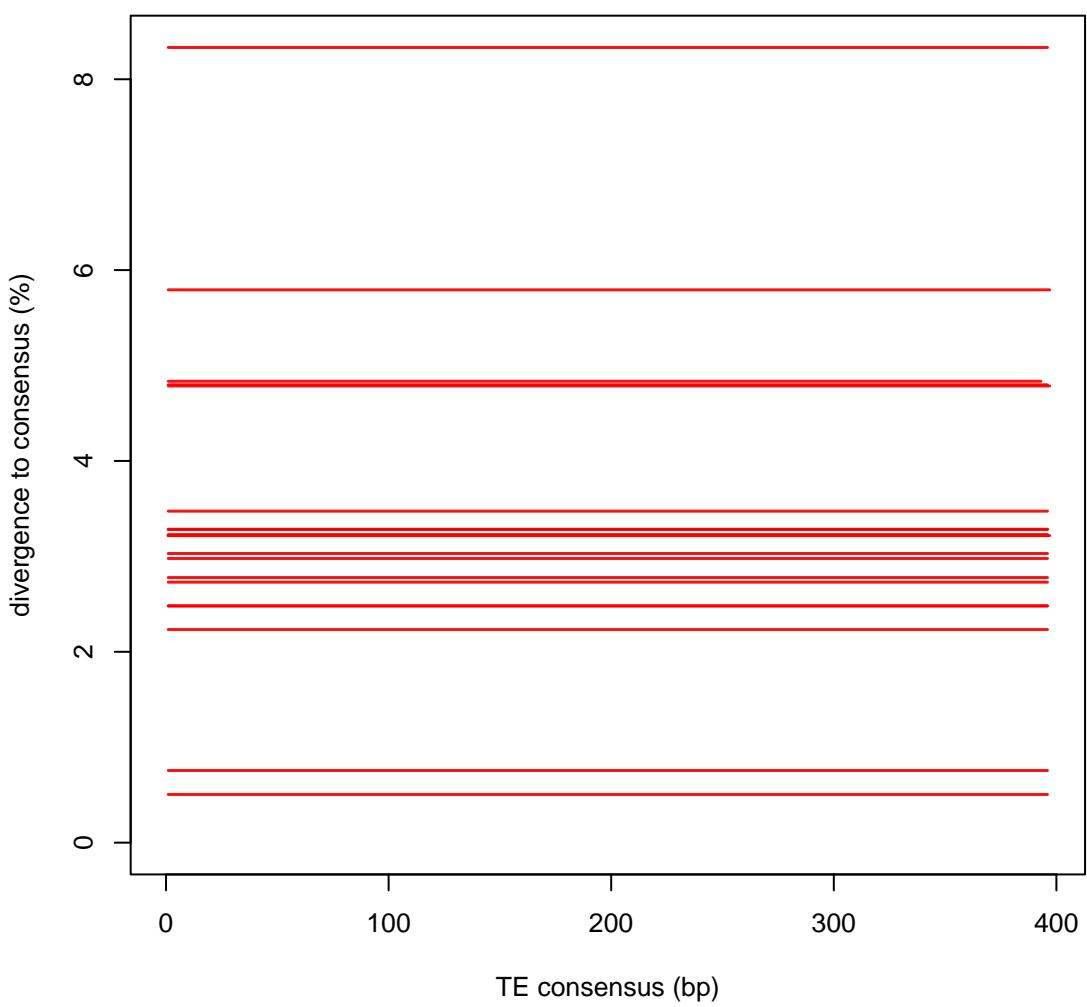


TE: oenMel1-36
consensus size: 397bp; fragments: 22; full length: 22 (≥ 357.3 bp)



TE consensus genomic coverage

