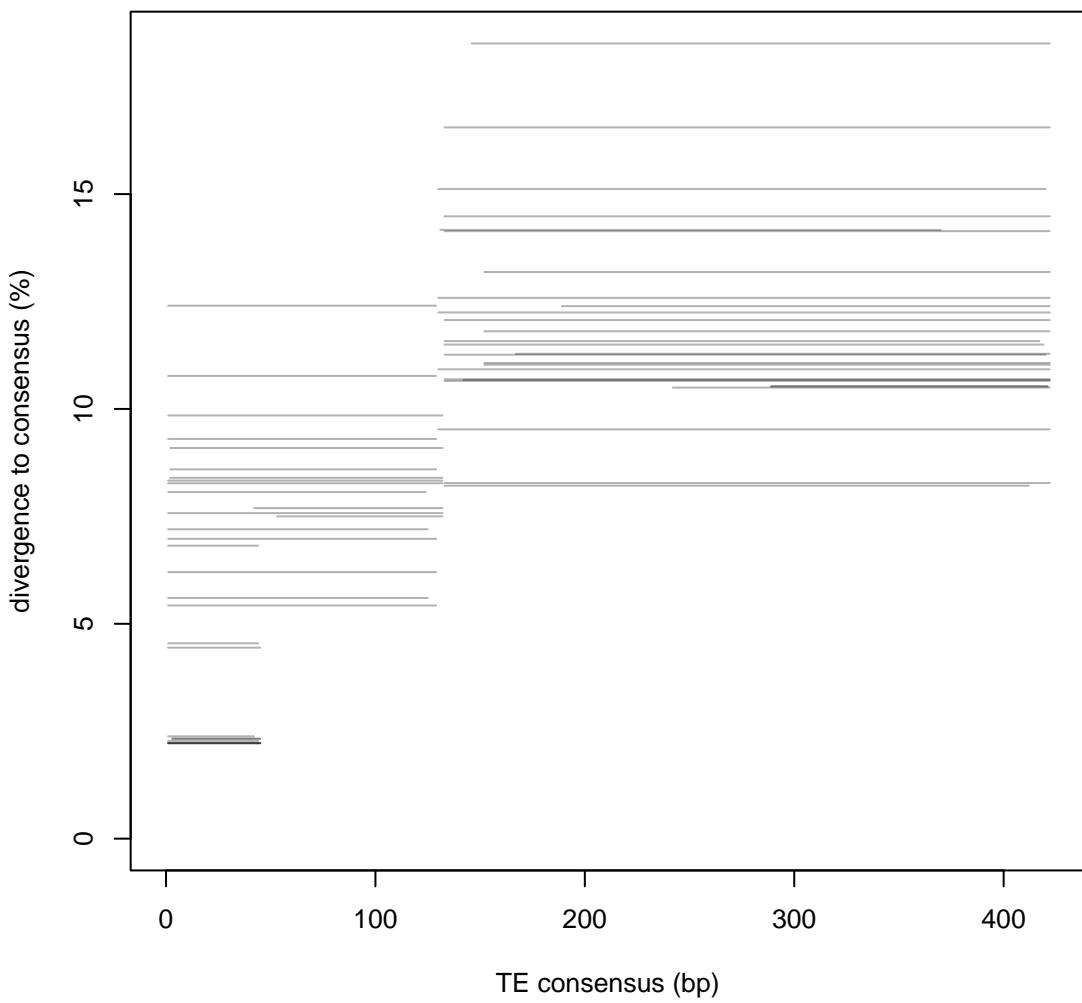


TE: seq6  
consensus size: 422bp; fragments: 57; full length: 0 (  $\geq$  379.8 bp )



### TE consensus genomic coverage

