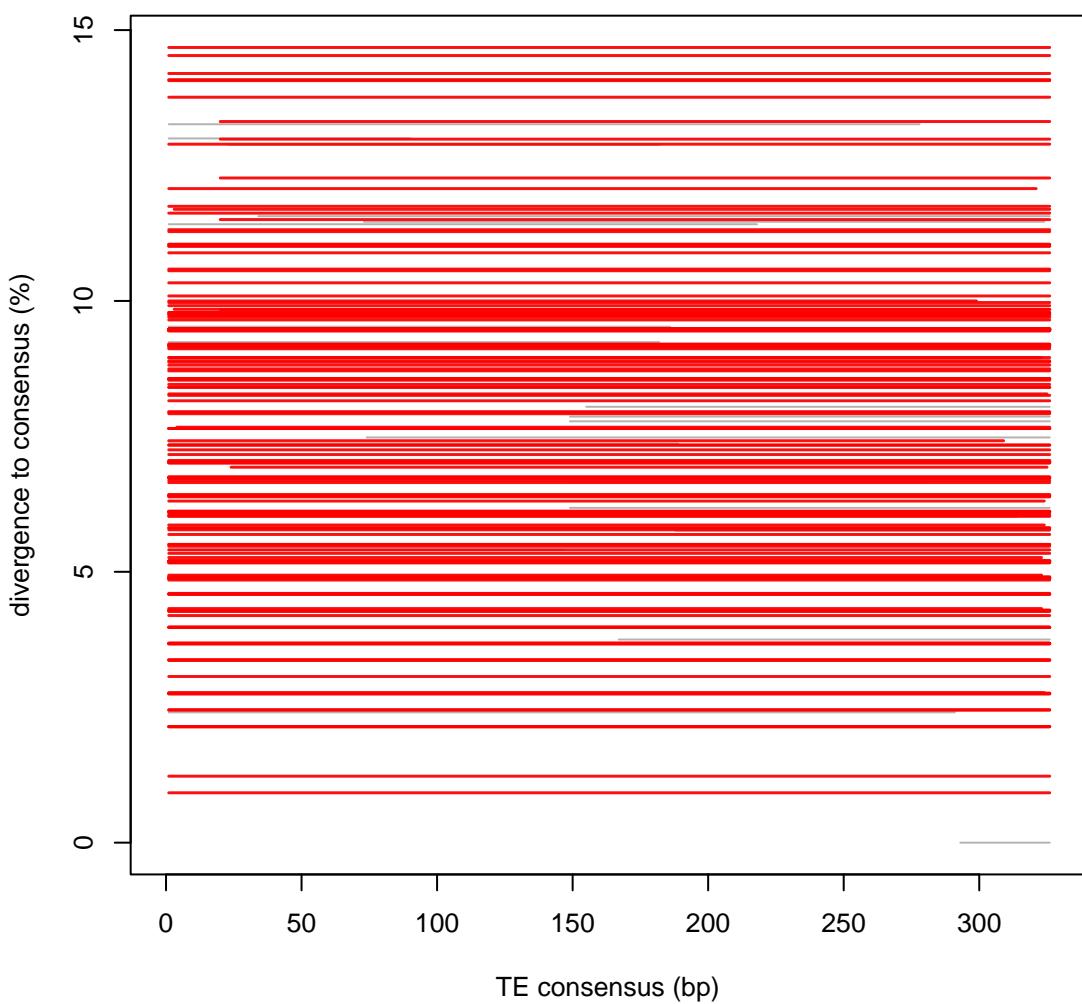


**TE: seq5**  
consensus size: 326bp; fragments: 204; full length: 182 ( $\geq 293.4$  bp)



**TE consensus genomic coverage**

