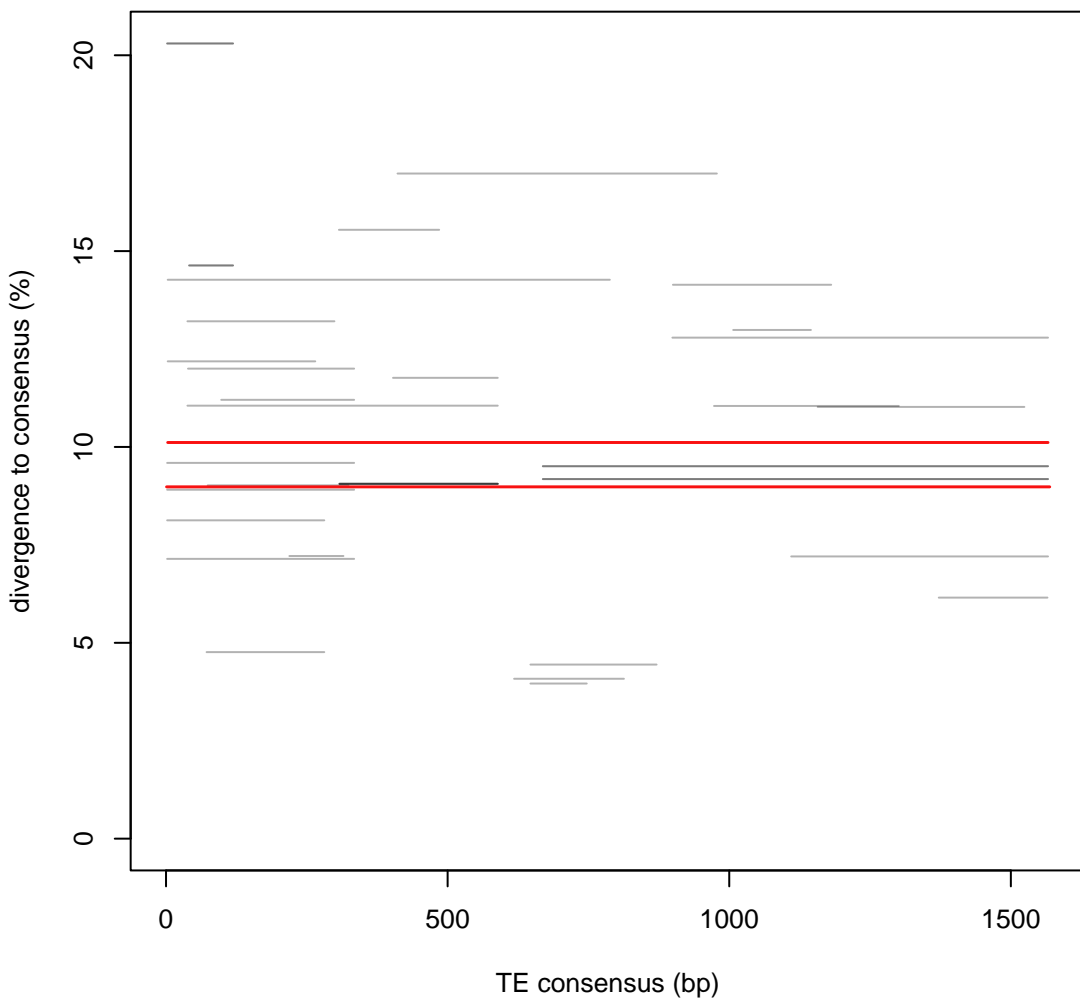
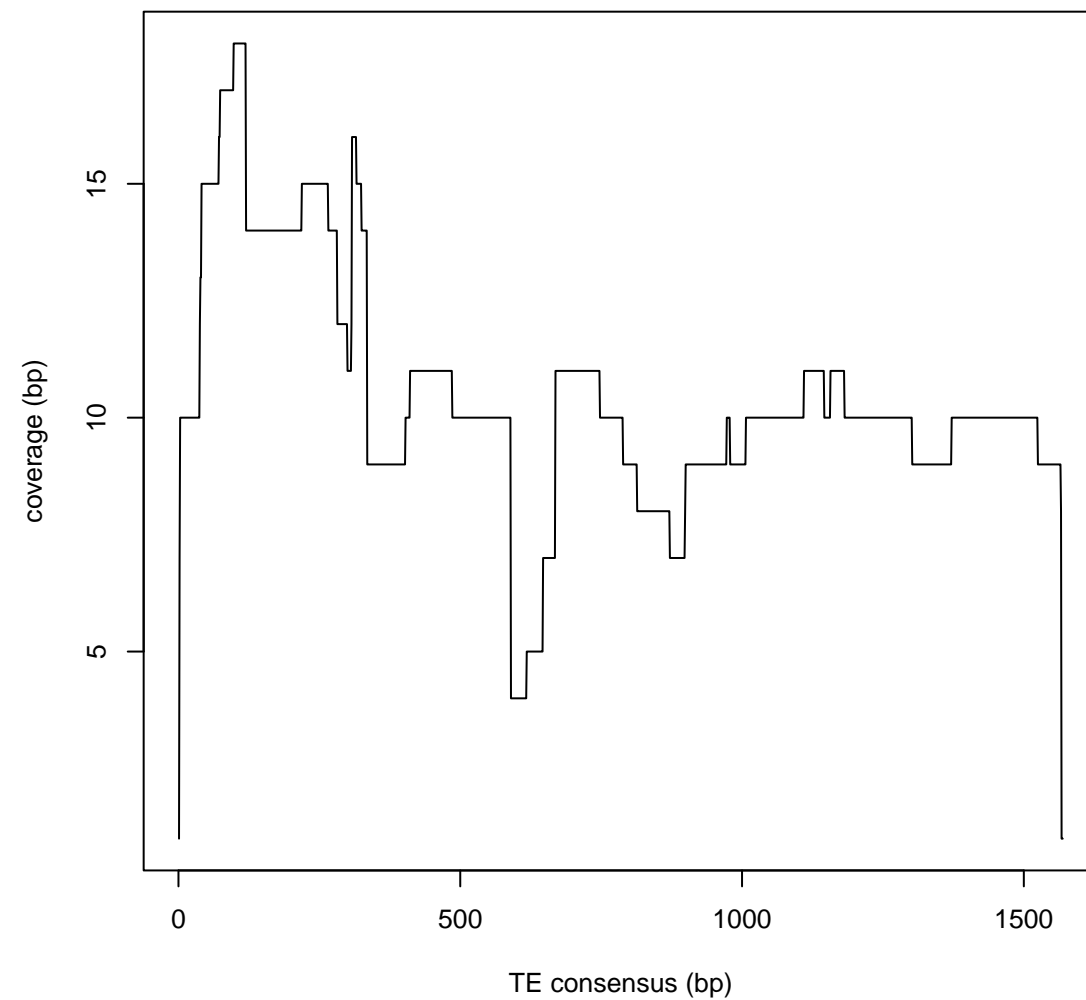


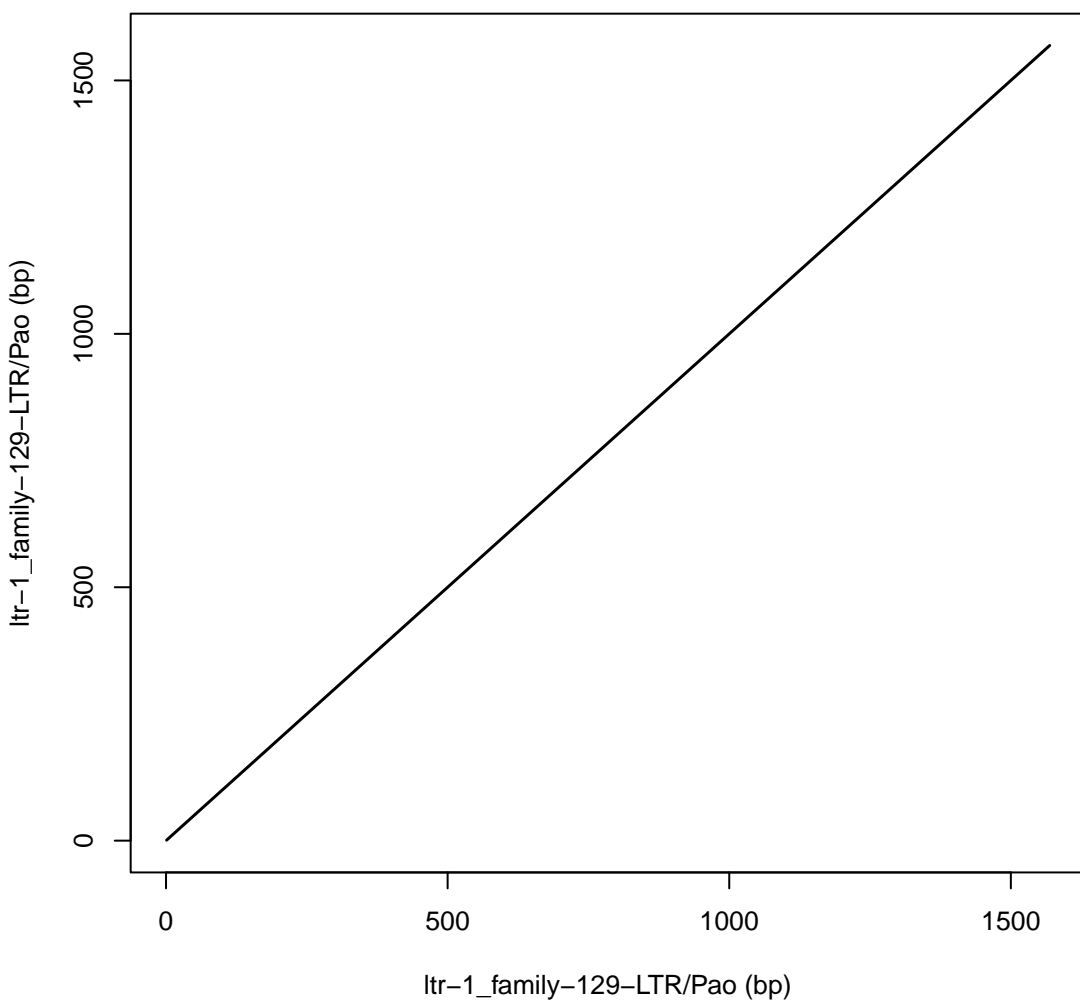
TE: ltr-1\_family-129-LTR/Pao  
consensus size: 1569bp; fragments: 40; full length: 2 ( $\geq 1412.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

