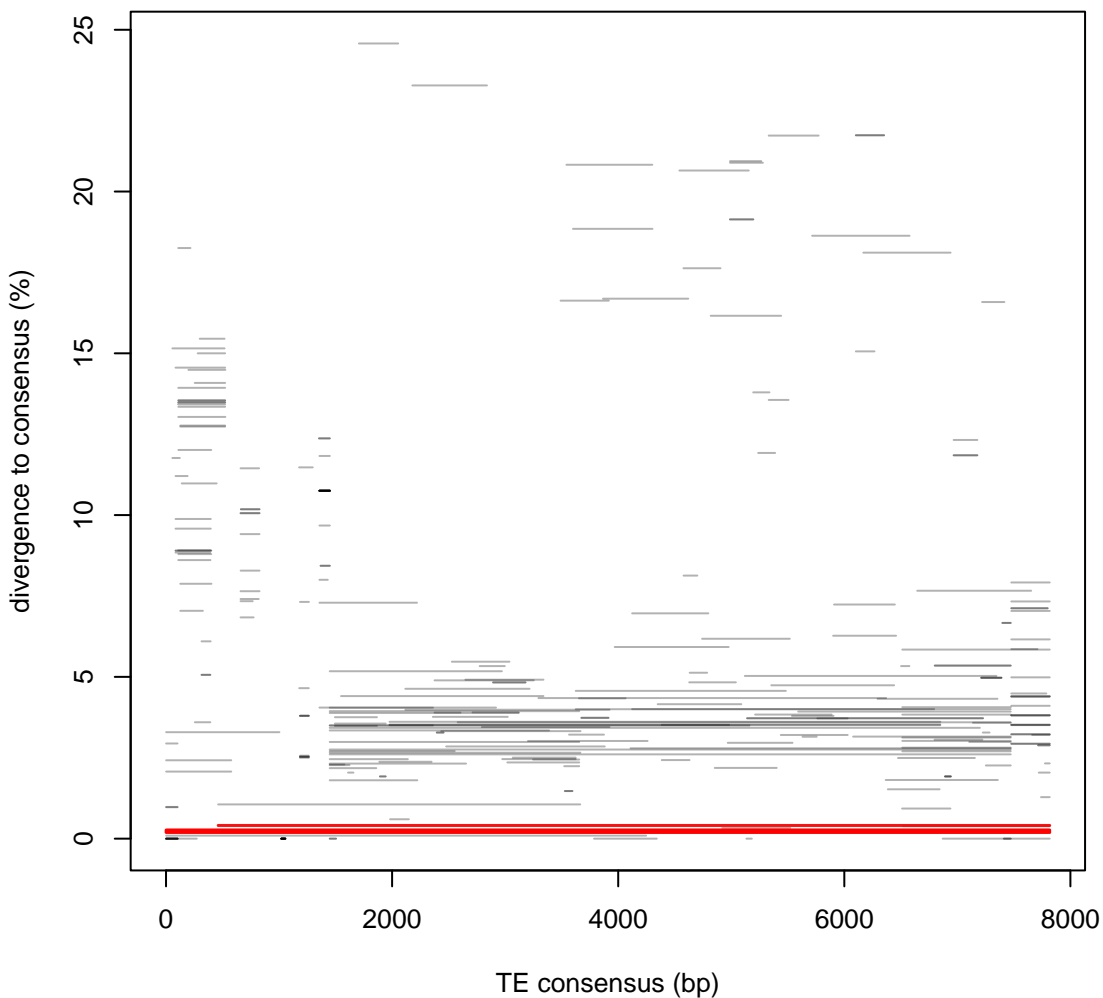
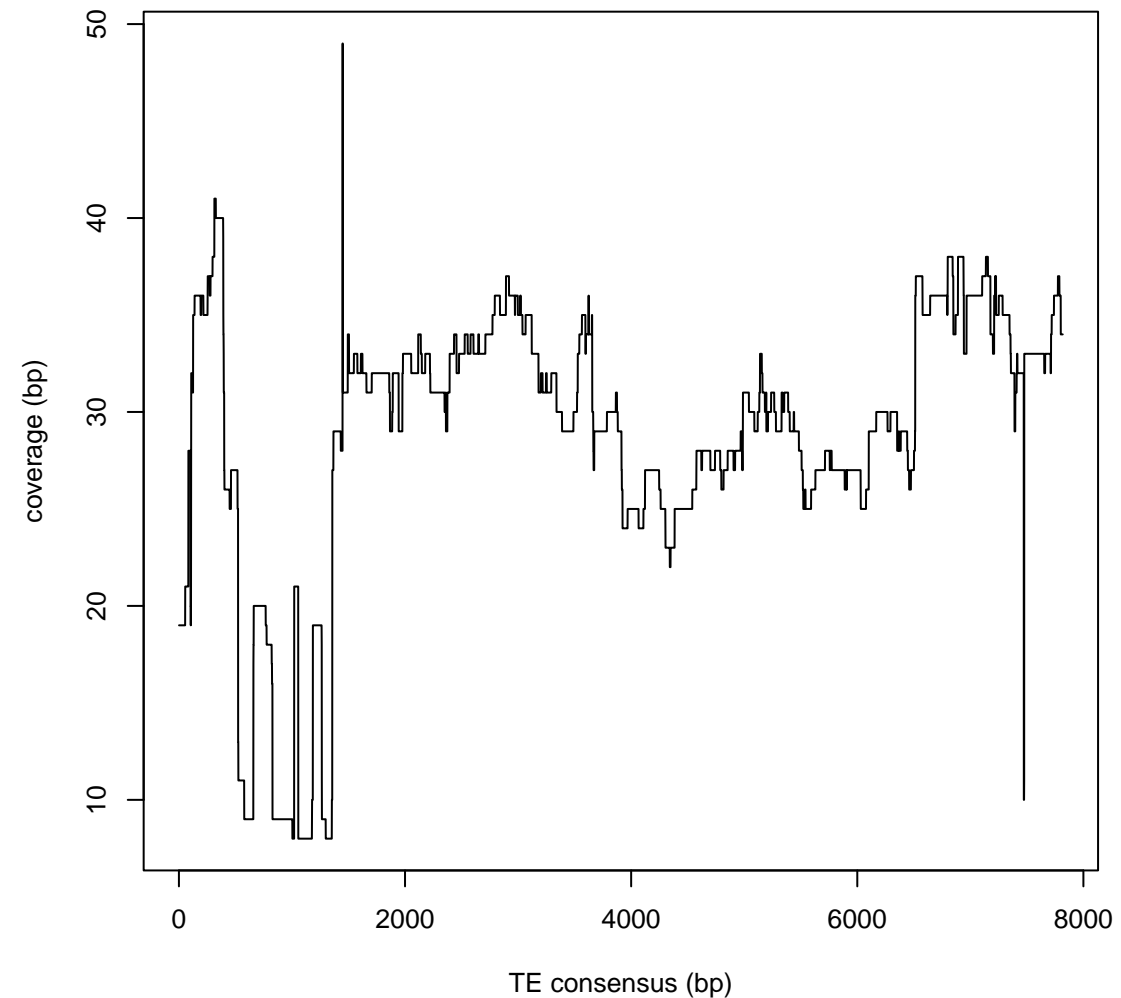


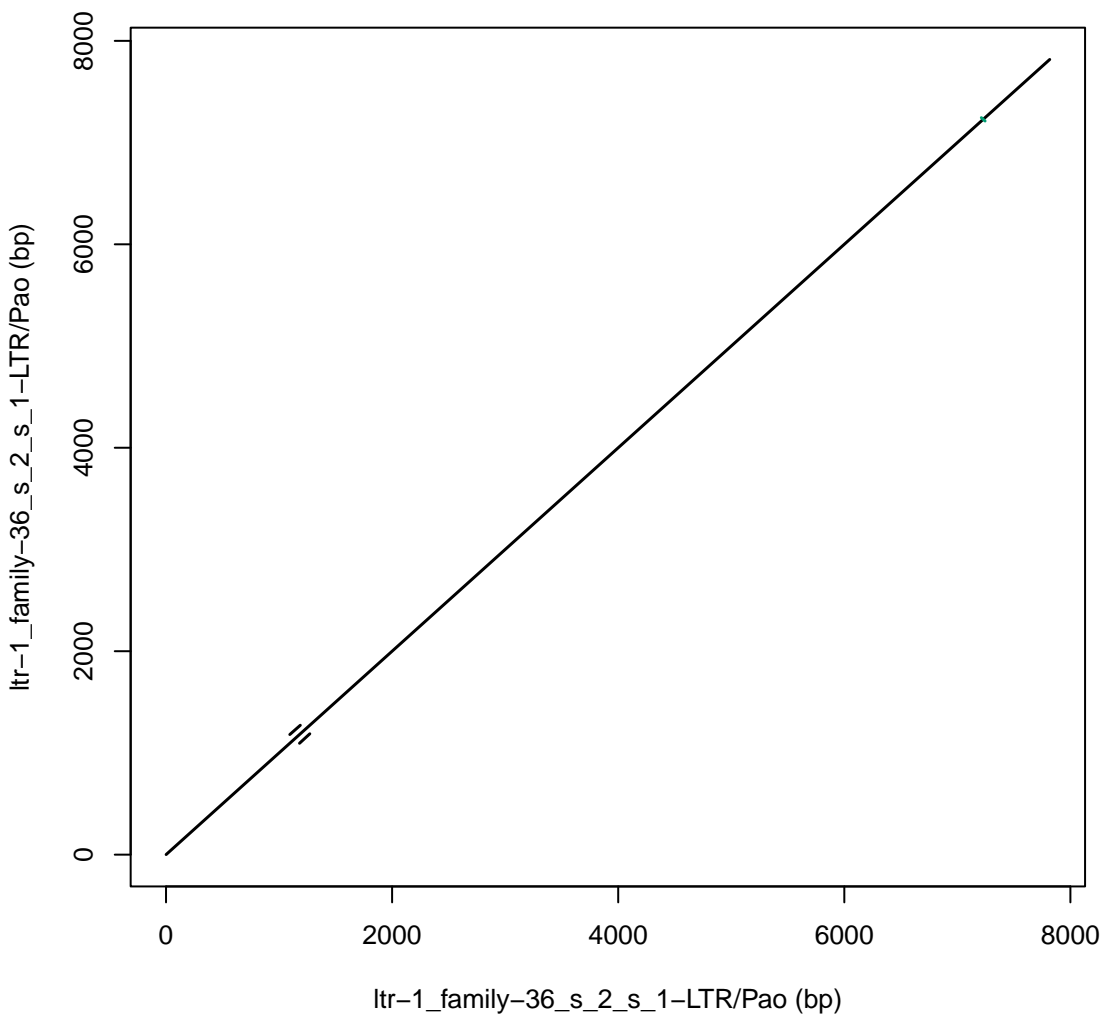
TE: ltr-1\_family-36\_s\_2\_s\_1-LTR/Pao  
consensus size: 7817bp; fragments: 294; full length: 6 ( $\geq 7035.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

