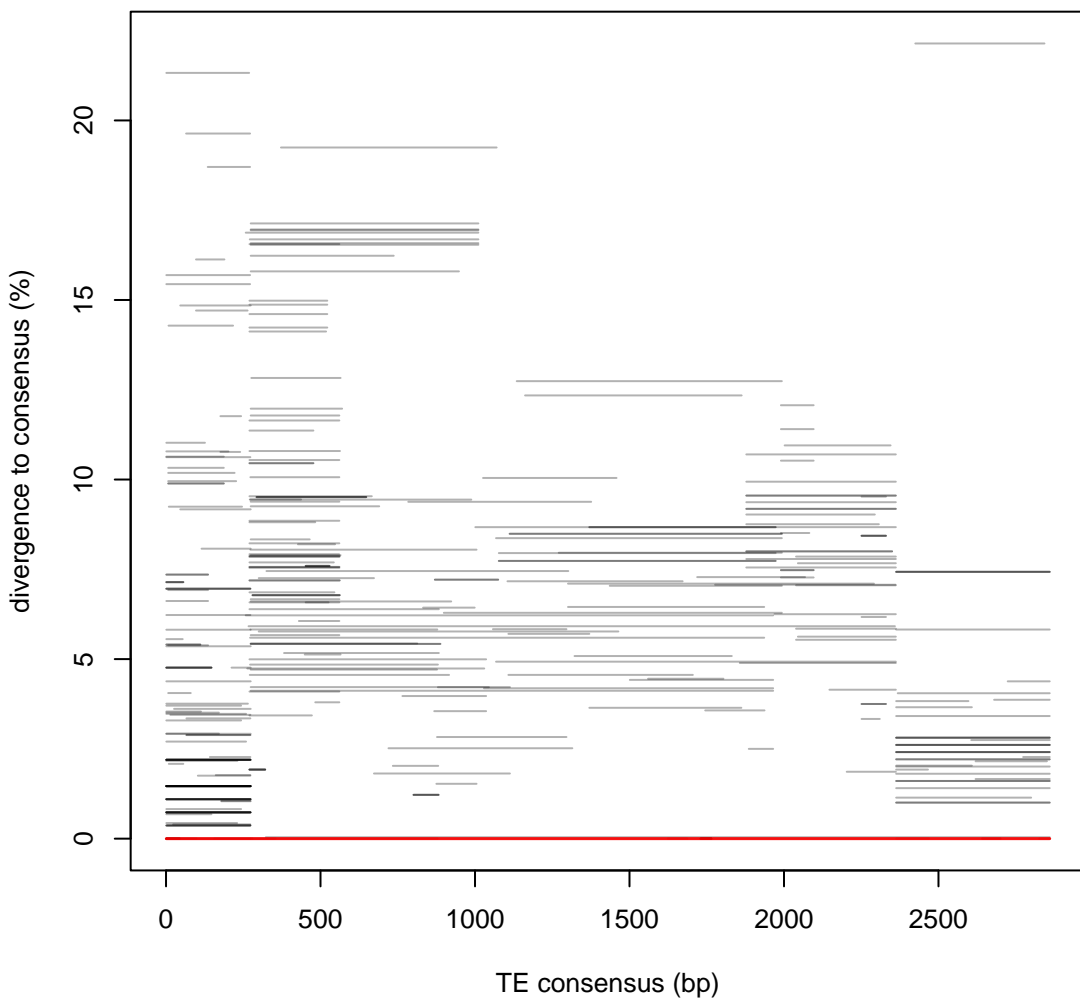
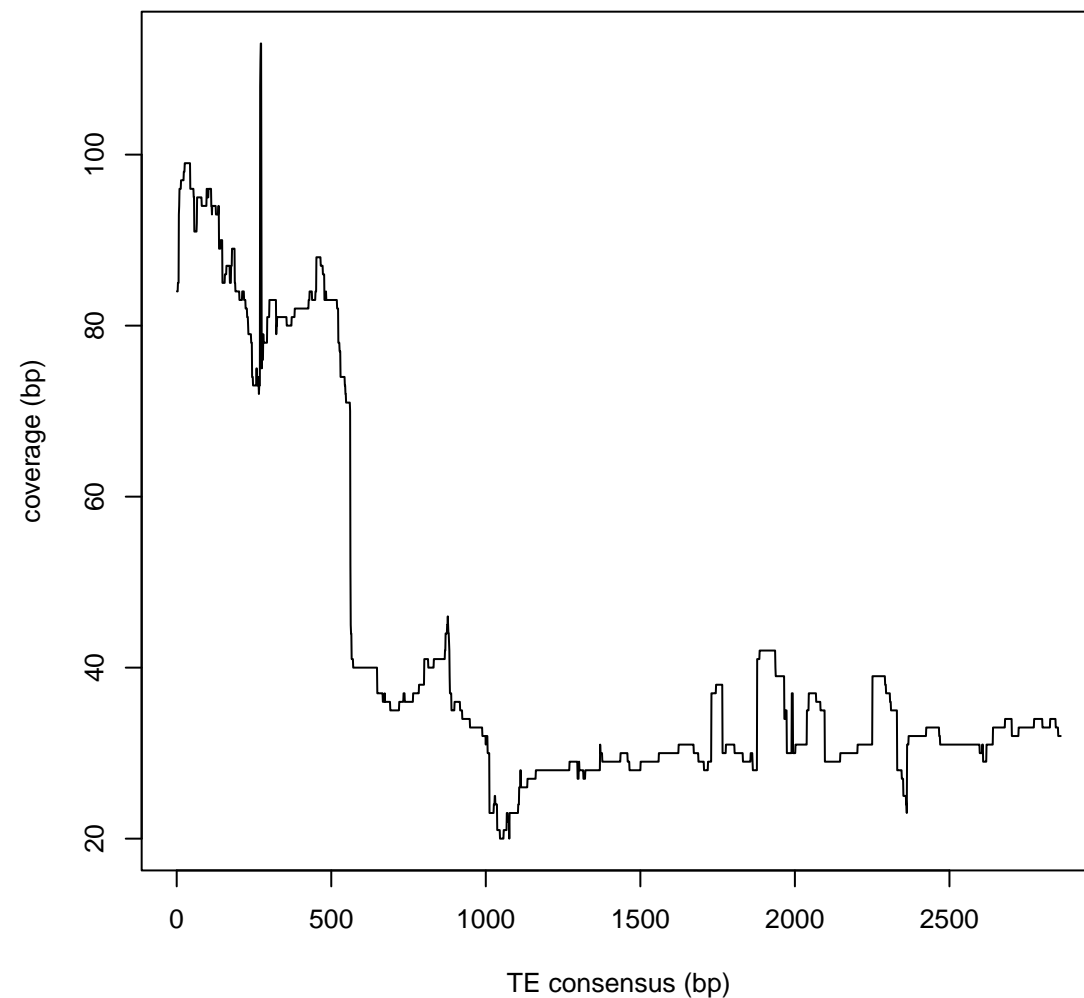


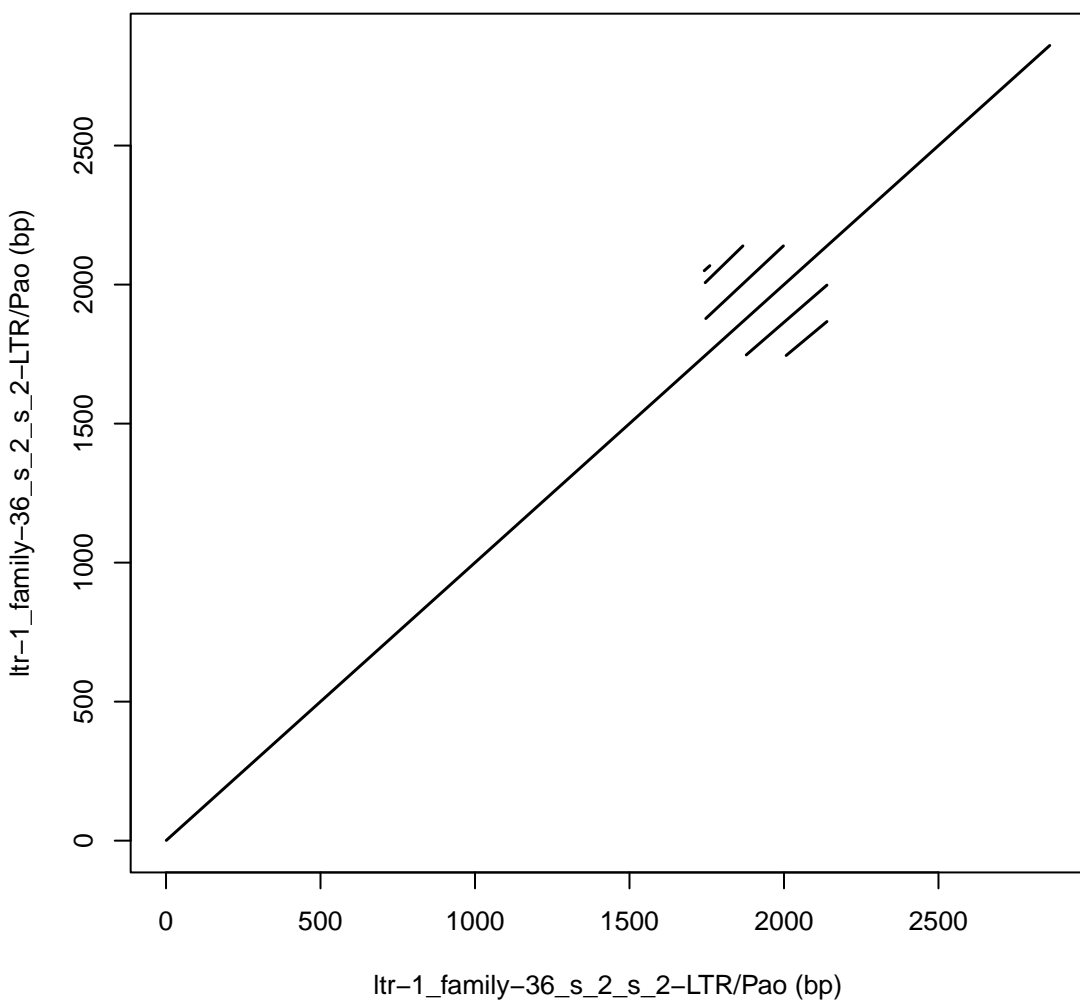
TE: ltr-1\_family-36\_s\_2\_s\_2-LTR/Pao  
consensus size: 2860bp; fragments: 345; full length: 1 (>=2574bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

