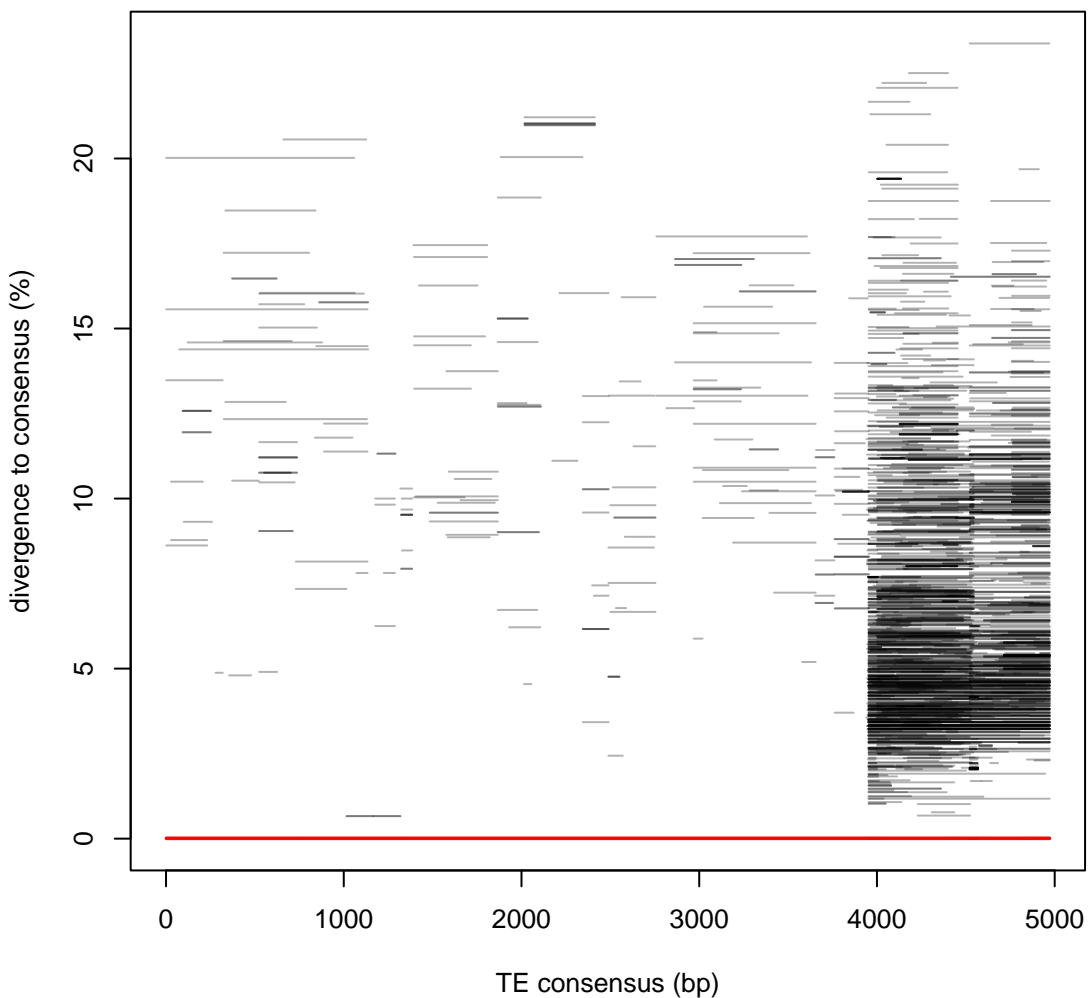
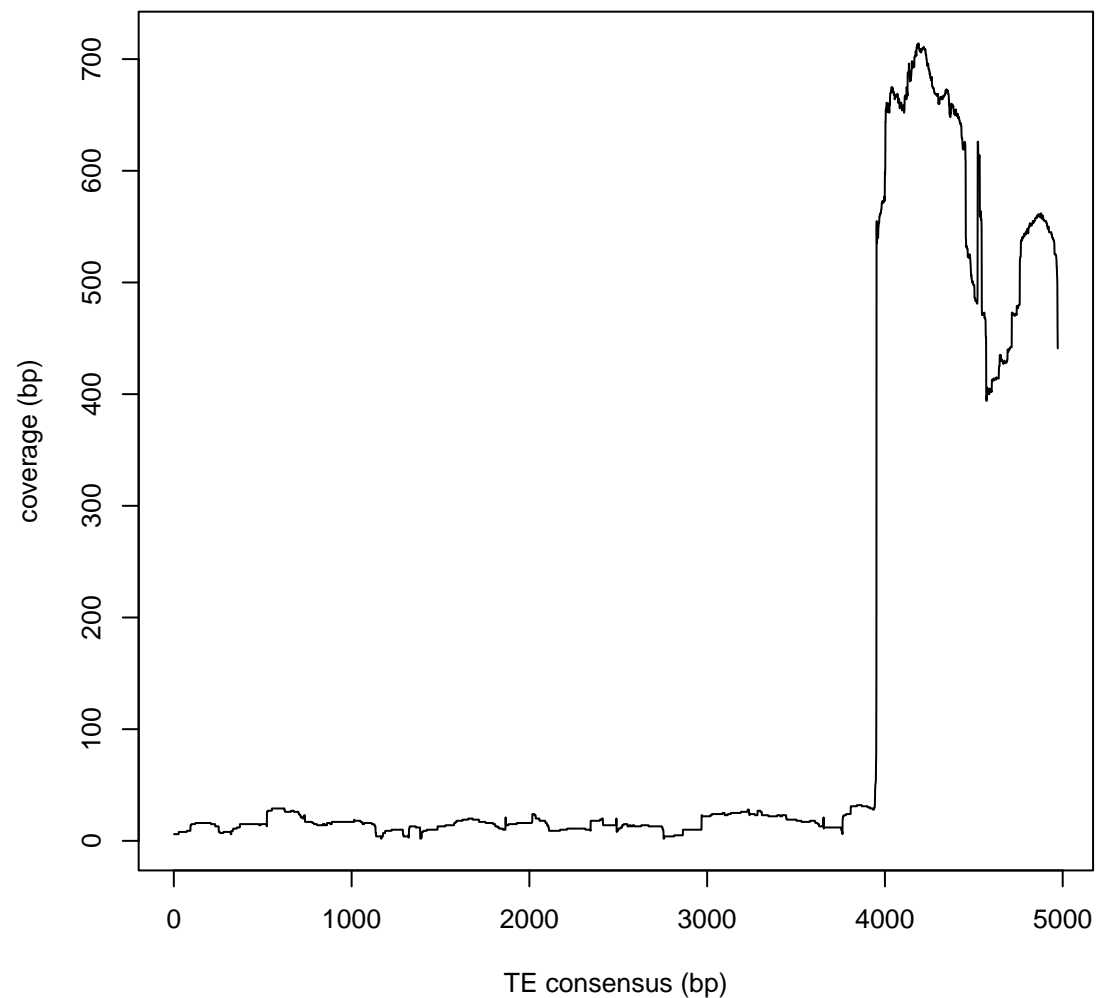


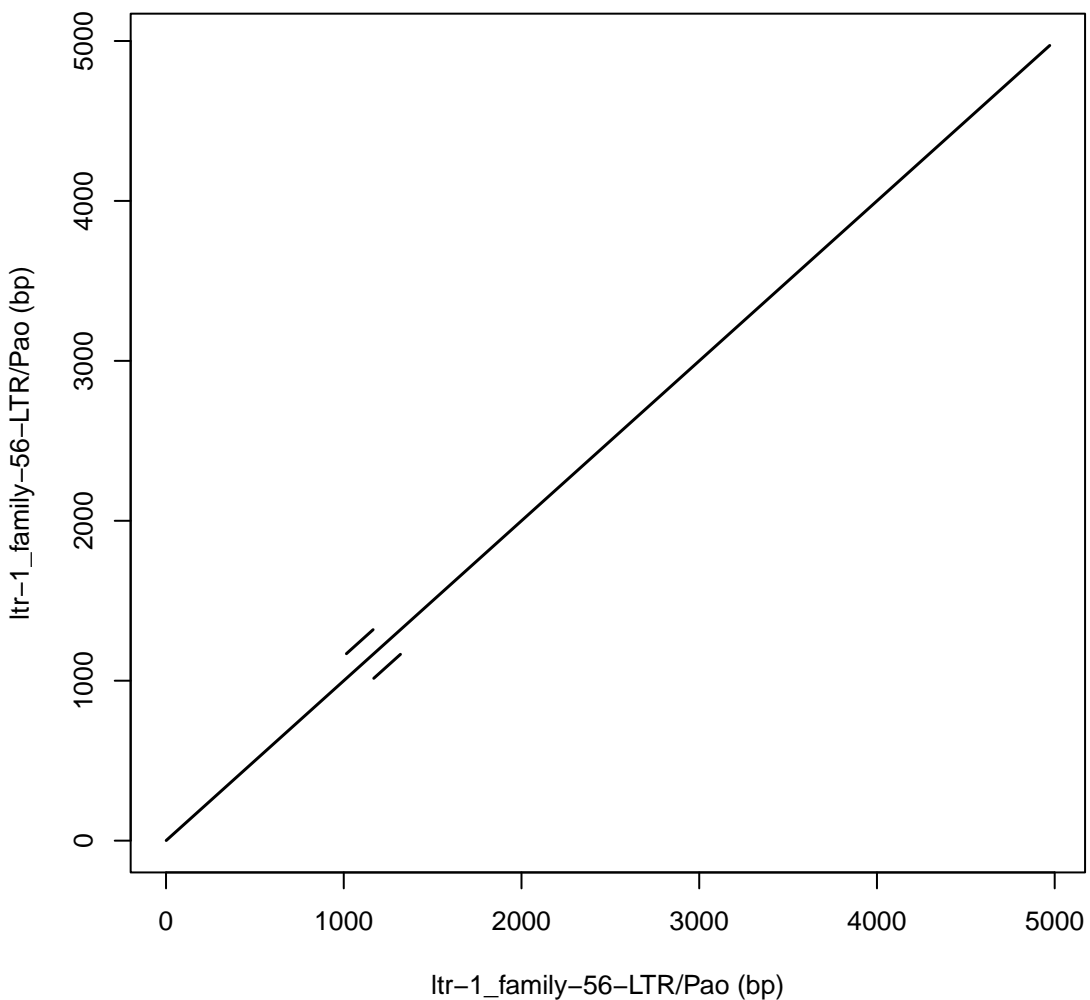
TE: ltr-1_family-56-LTR/Pao
consensus size: 4972bp; fragments: 1746; full length: 2 (>=4474.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

