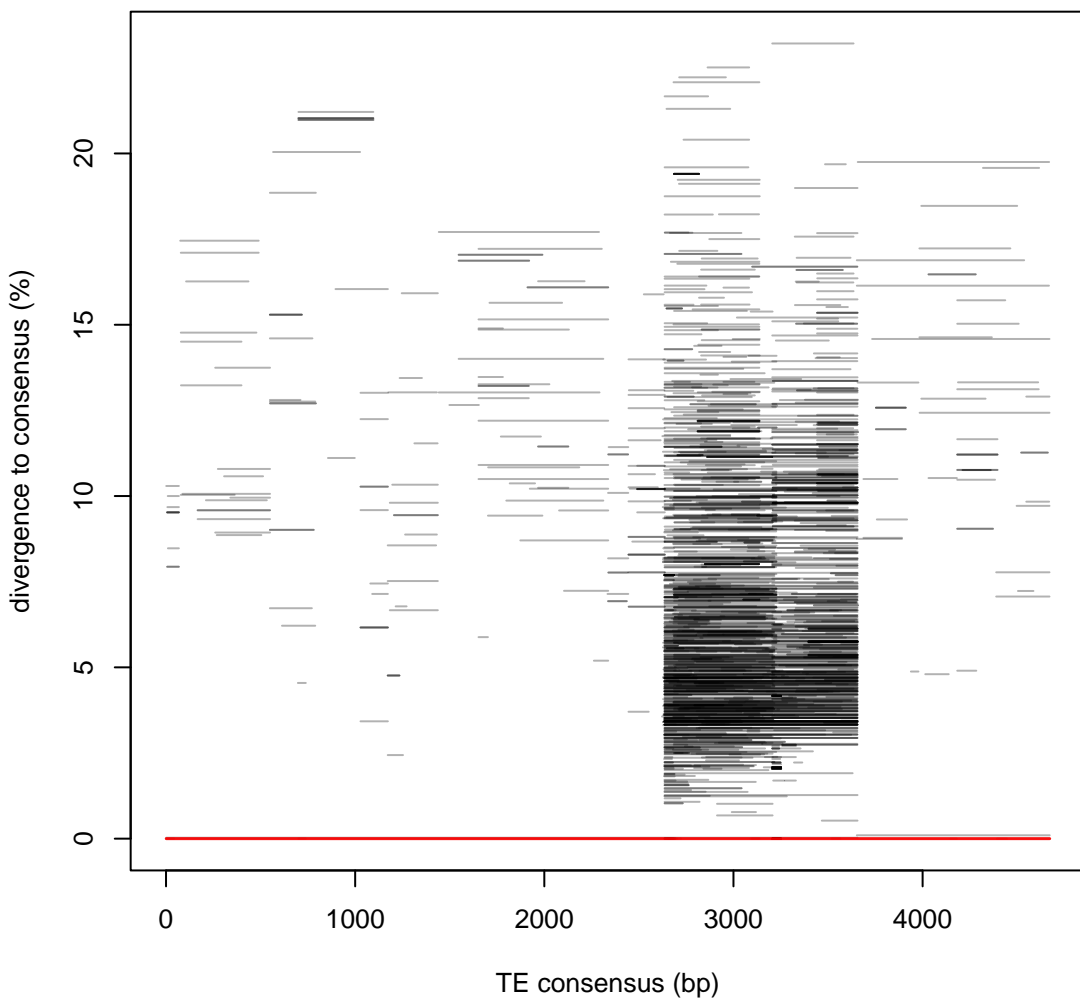
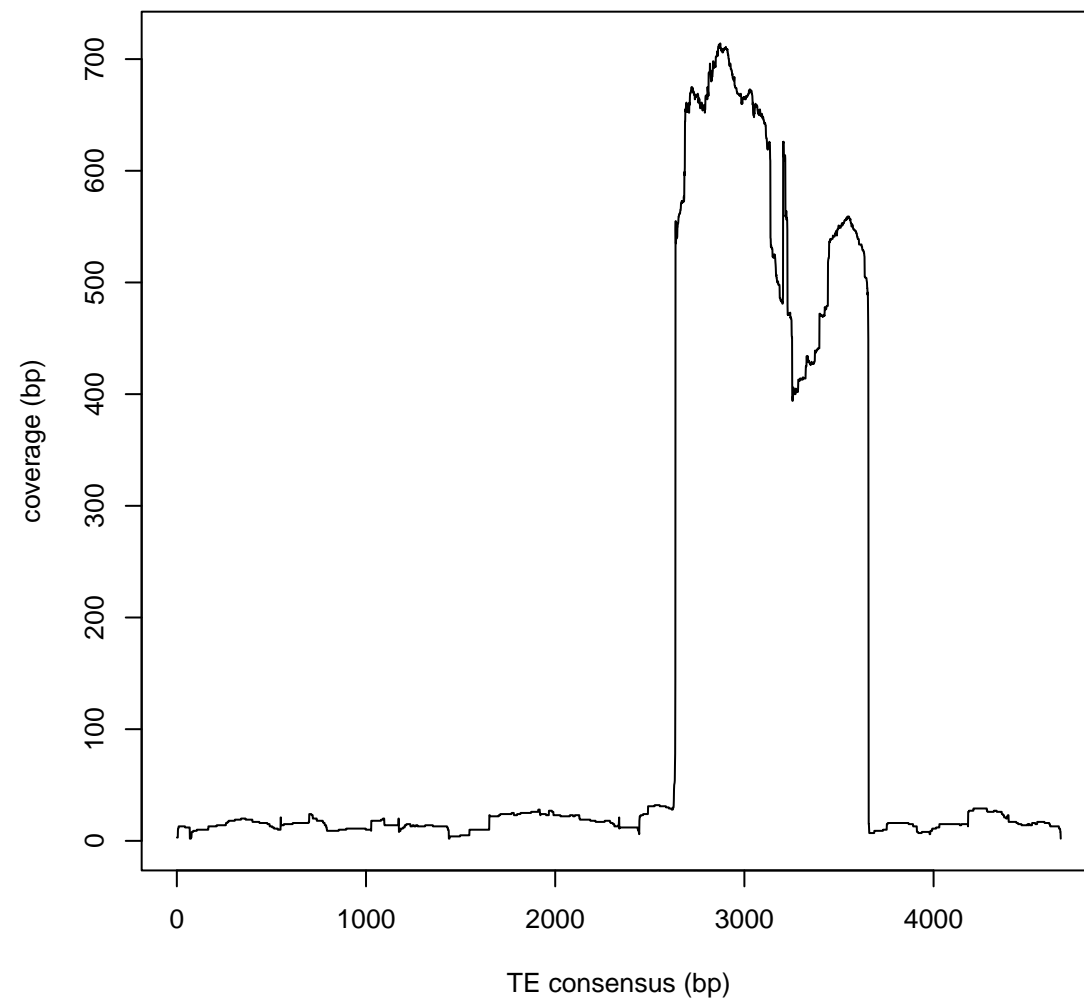


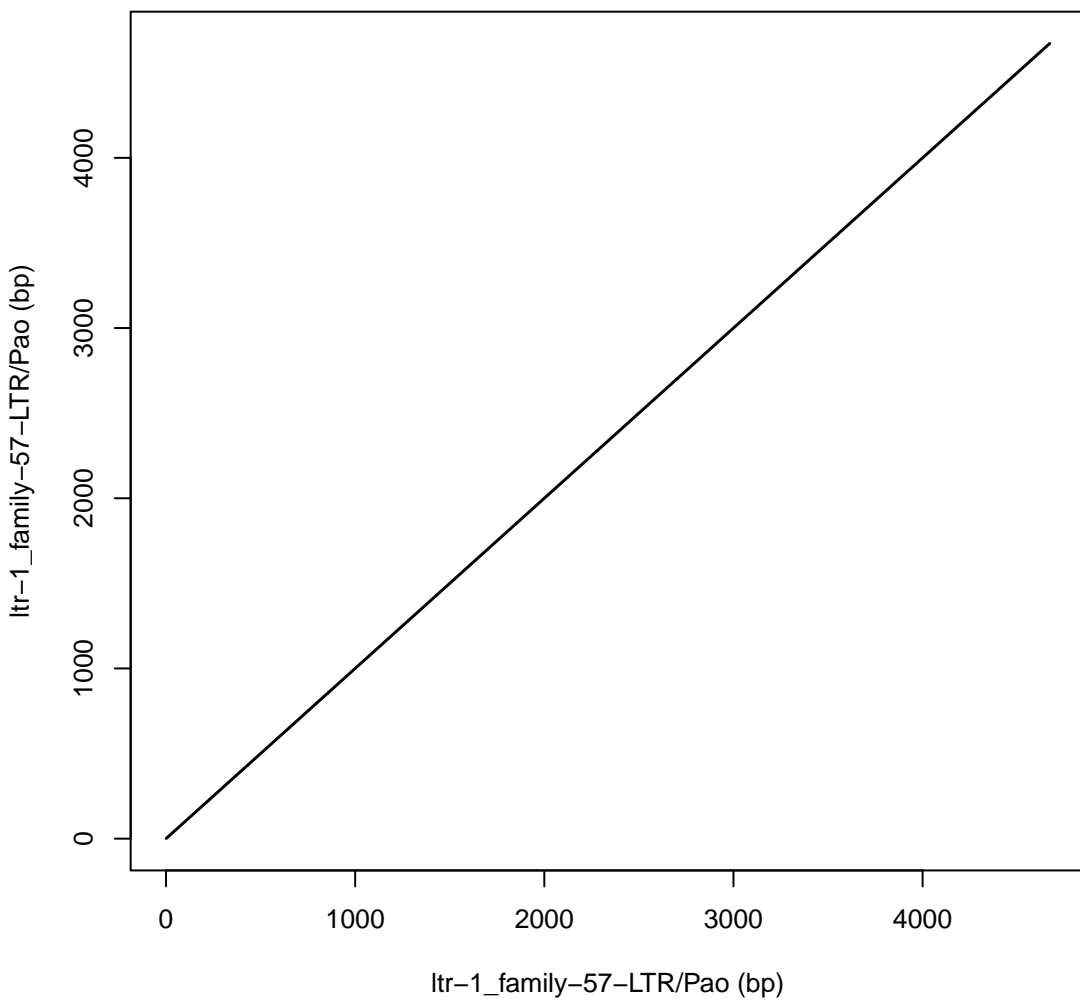
TE: ltr-1_family-57-LTR/Pao
consensus size: 4672bp; fragments: 1725; full length: 1 (>=4204.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

