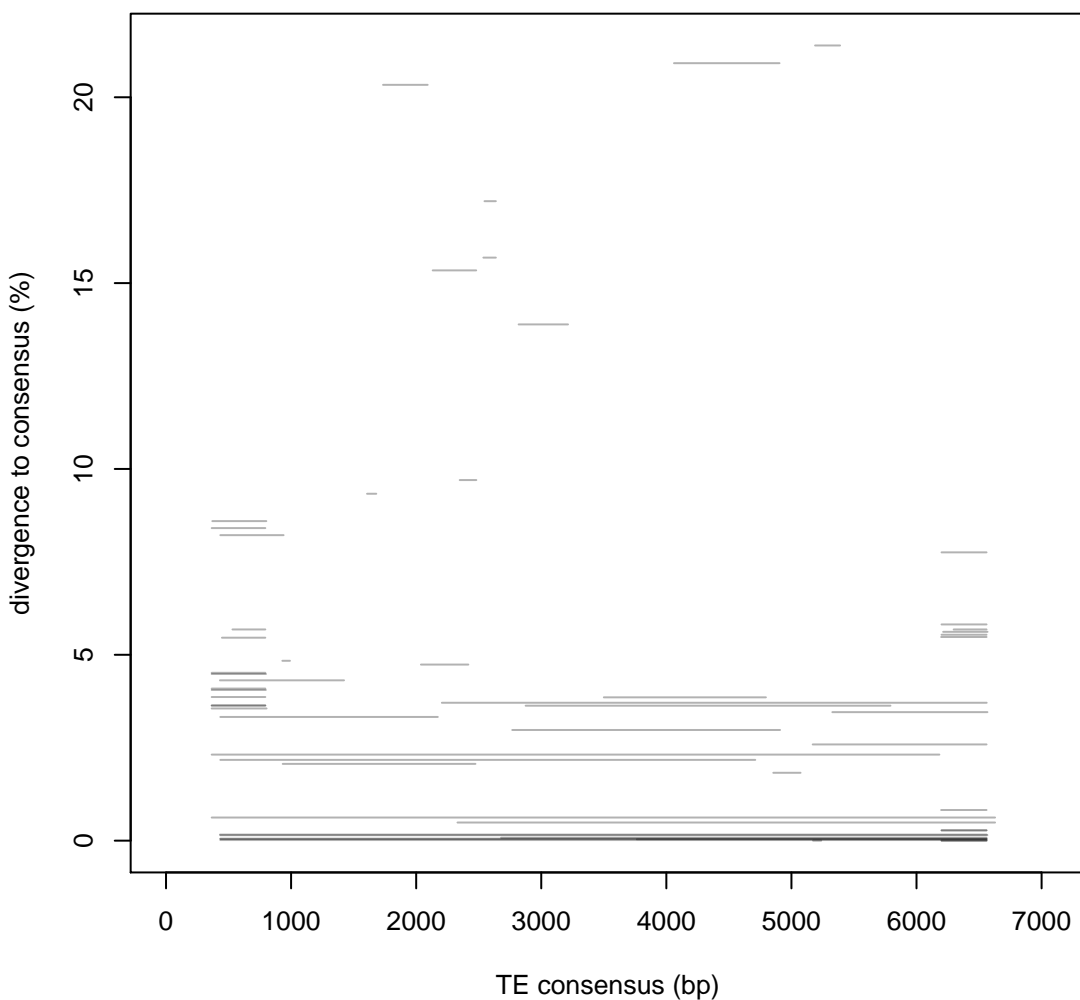
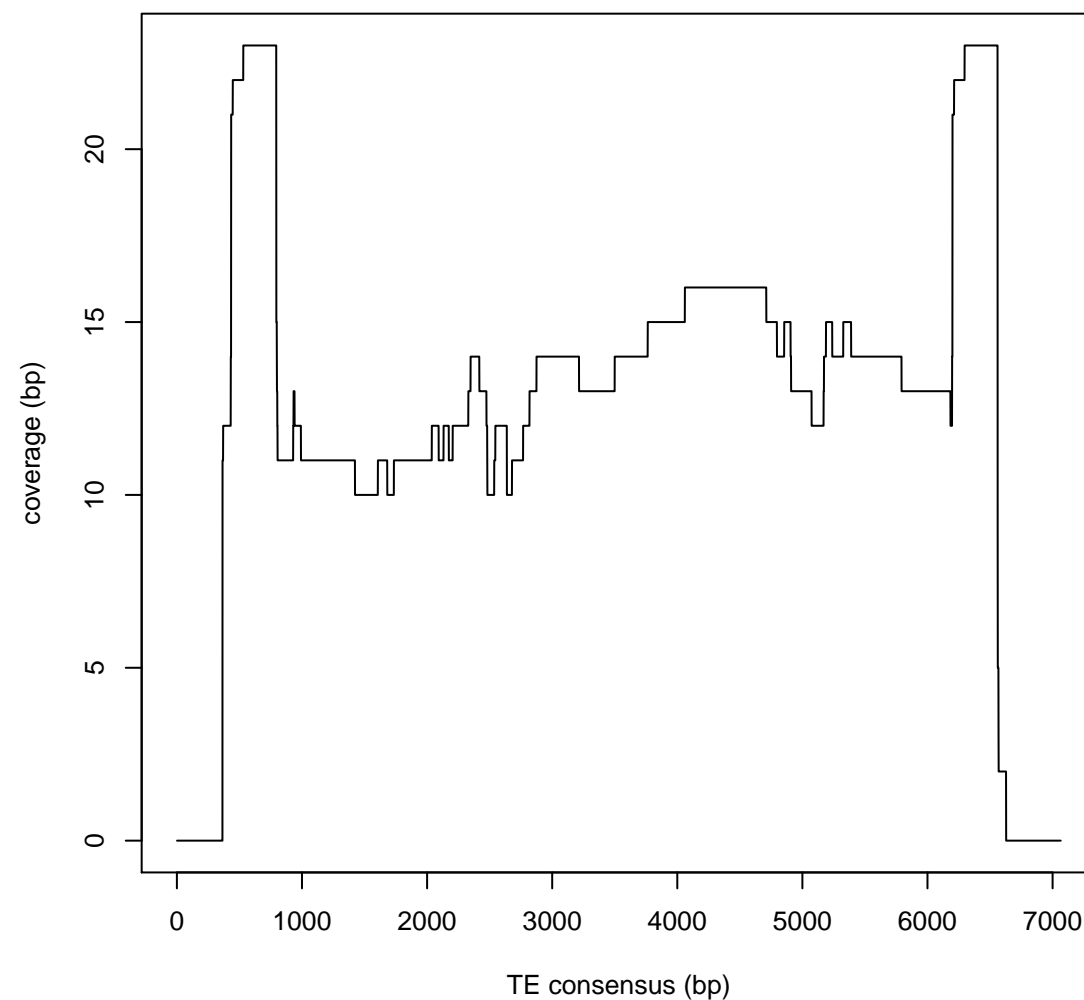


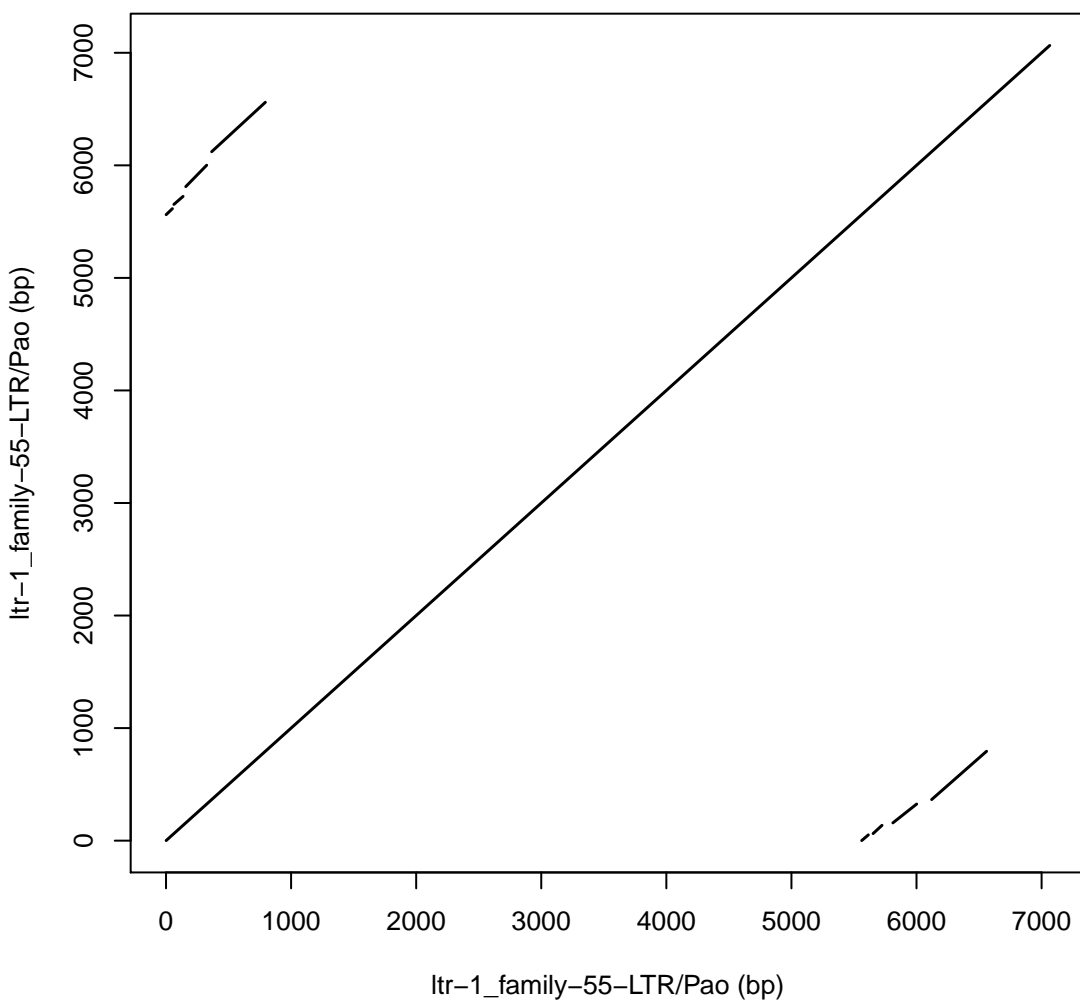
TE: ltr-1_family-55-LTR/Pao
consensus size: 7065bp; fragments: 57; full length: 0 (>=6358.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

