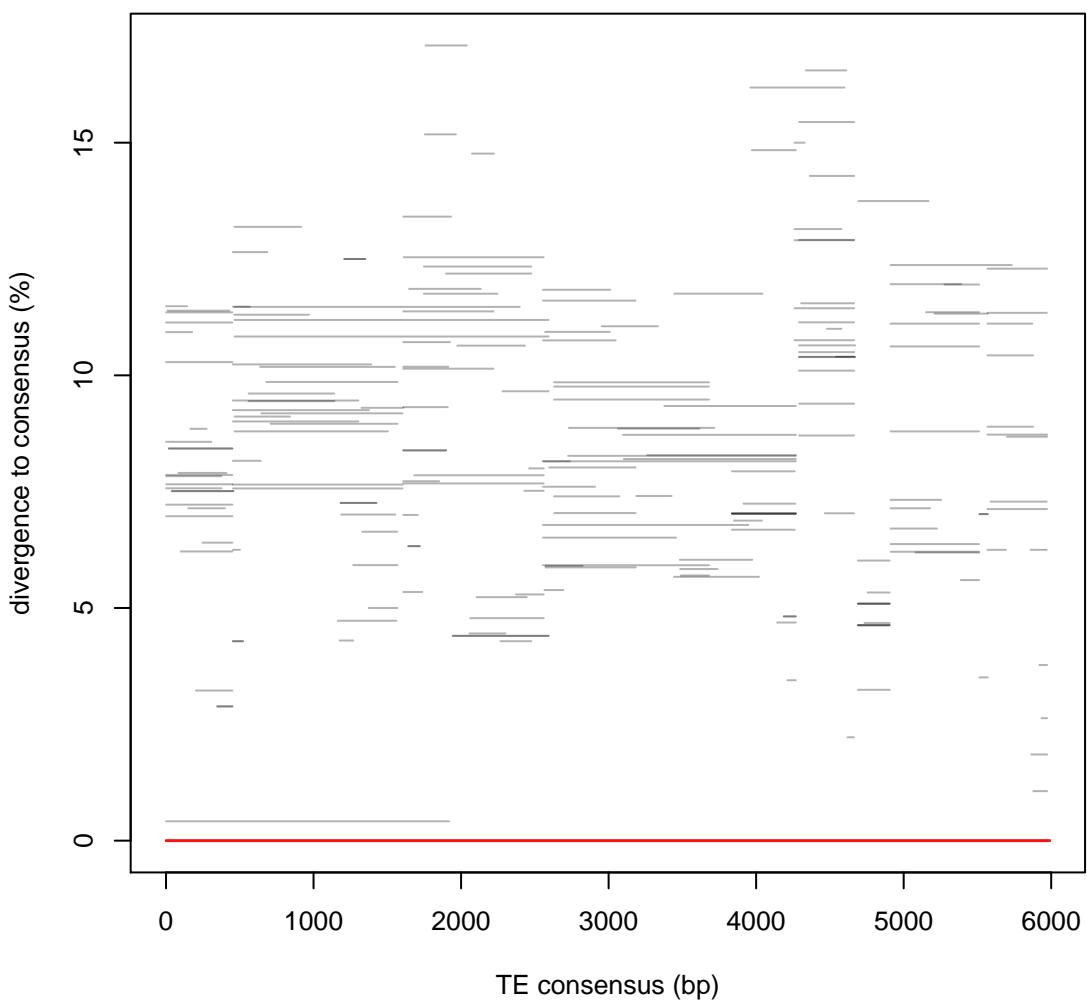
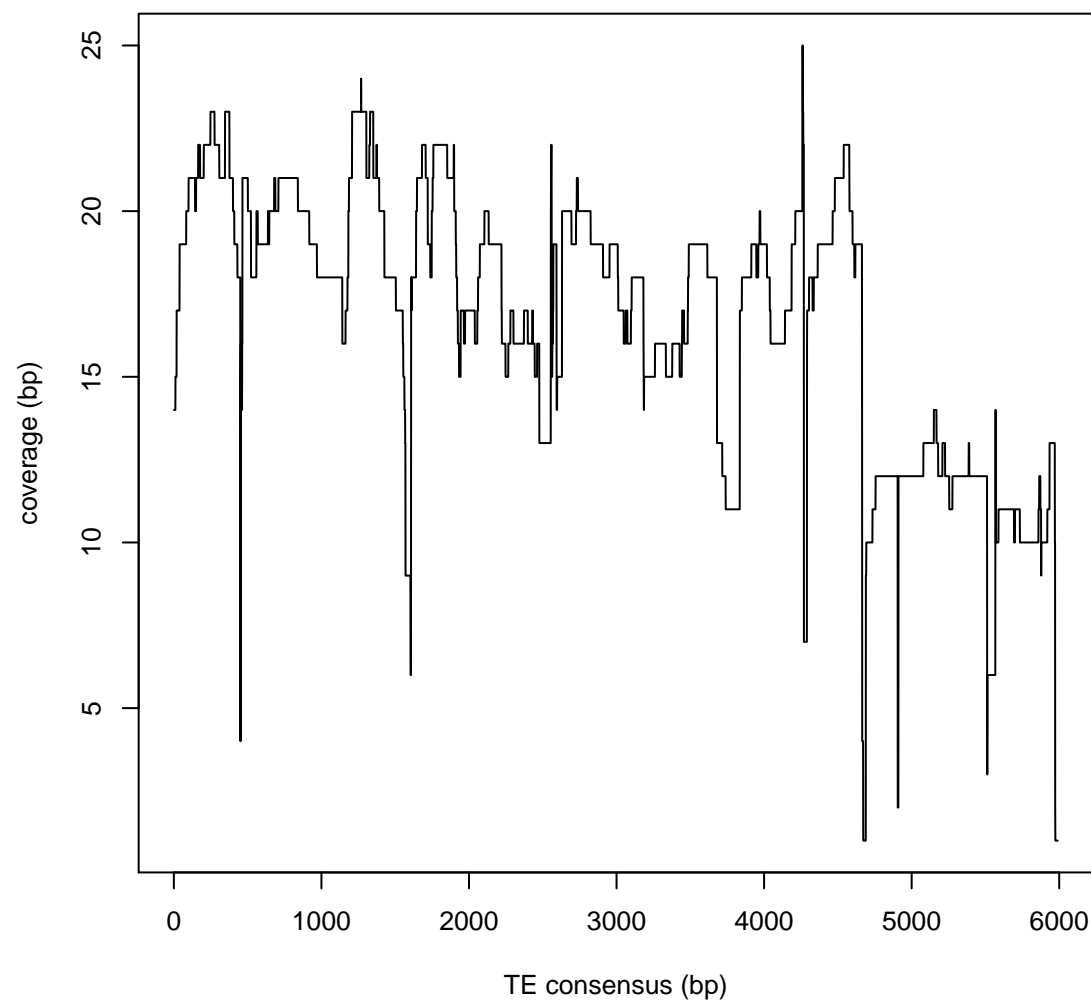


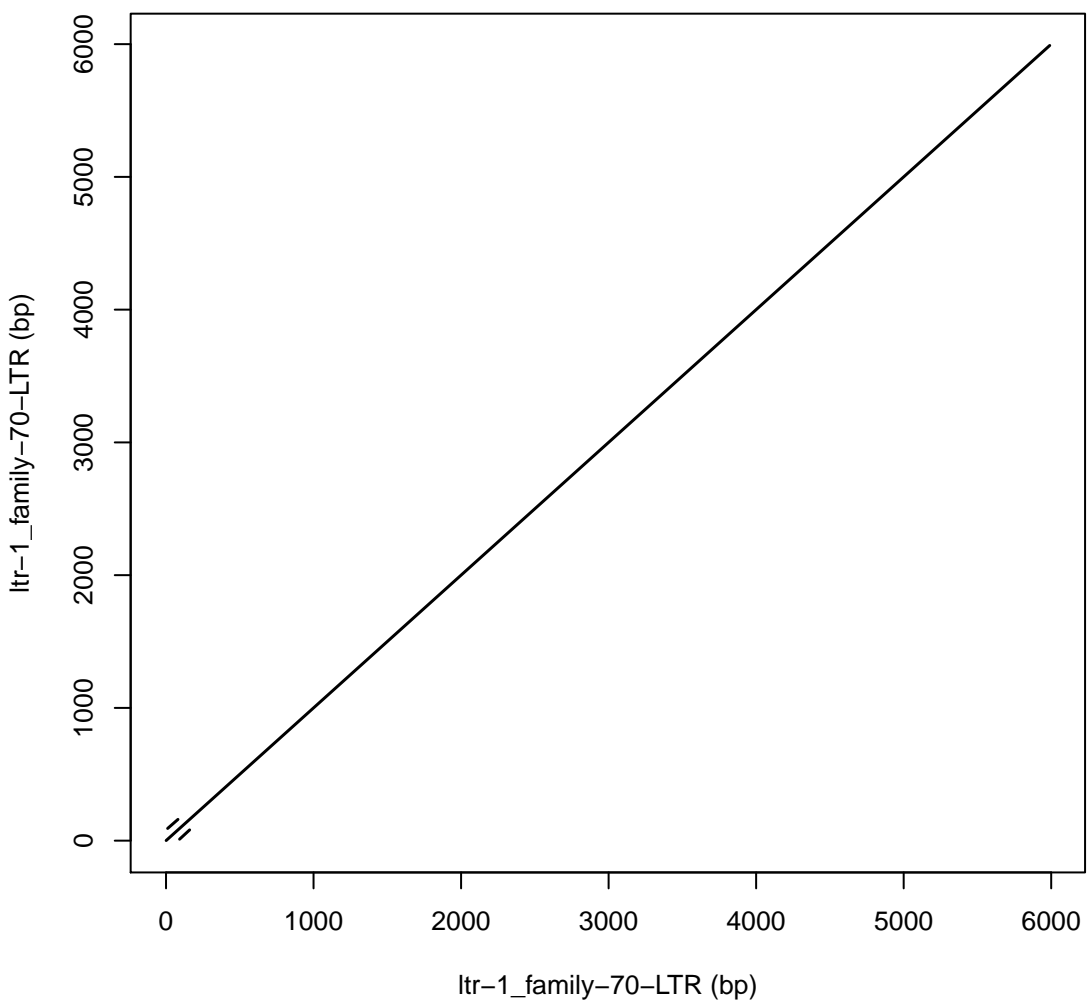
TE: ltr-1_family-70-LTR
consensus size: 5990bp; fragments: 211; full length: 1 (>=5391bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

