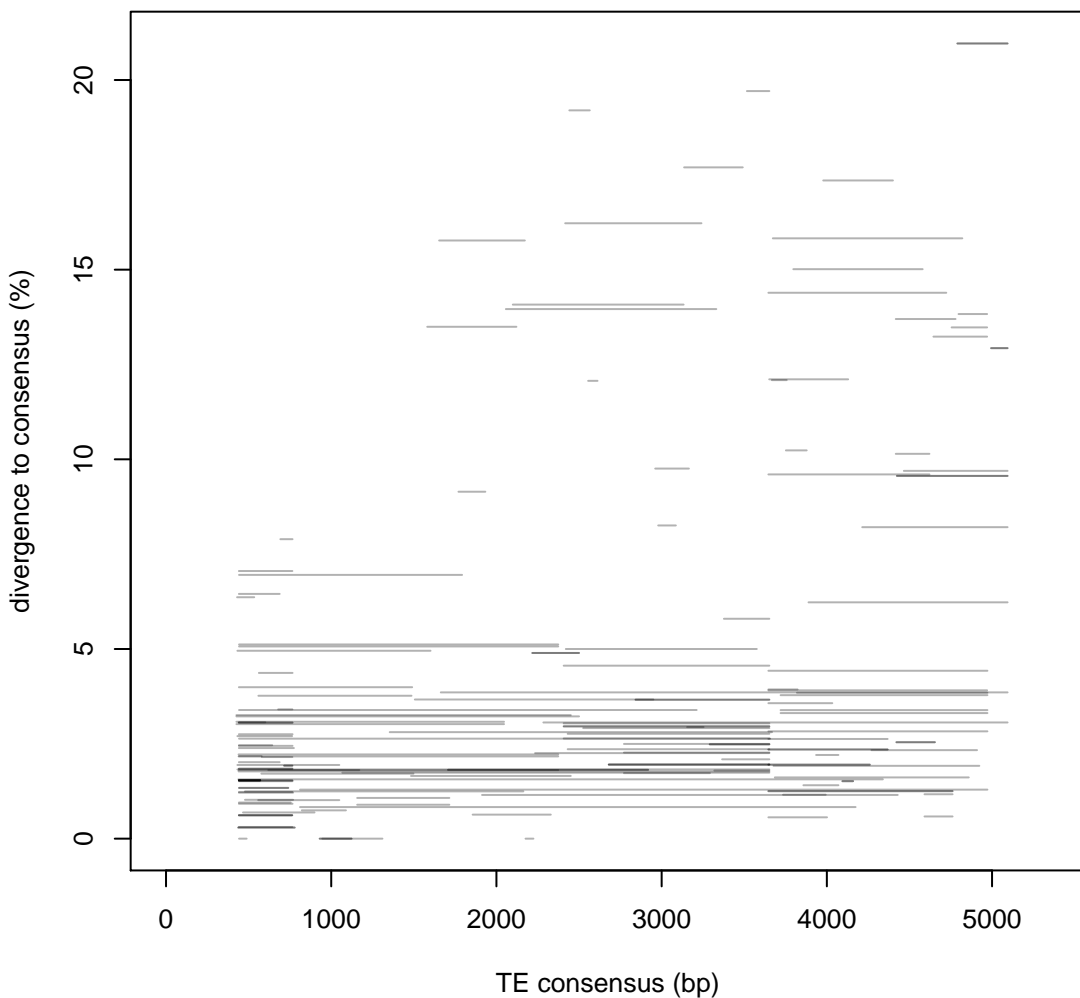
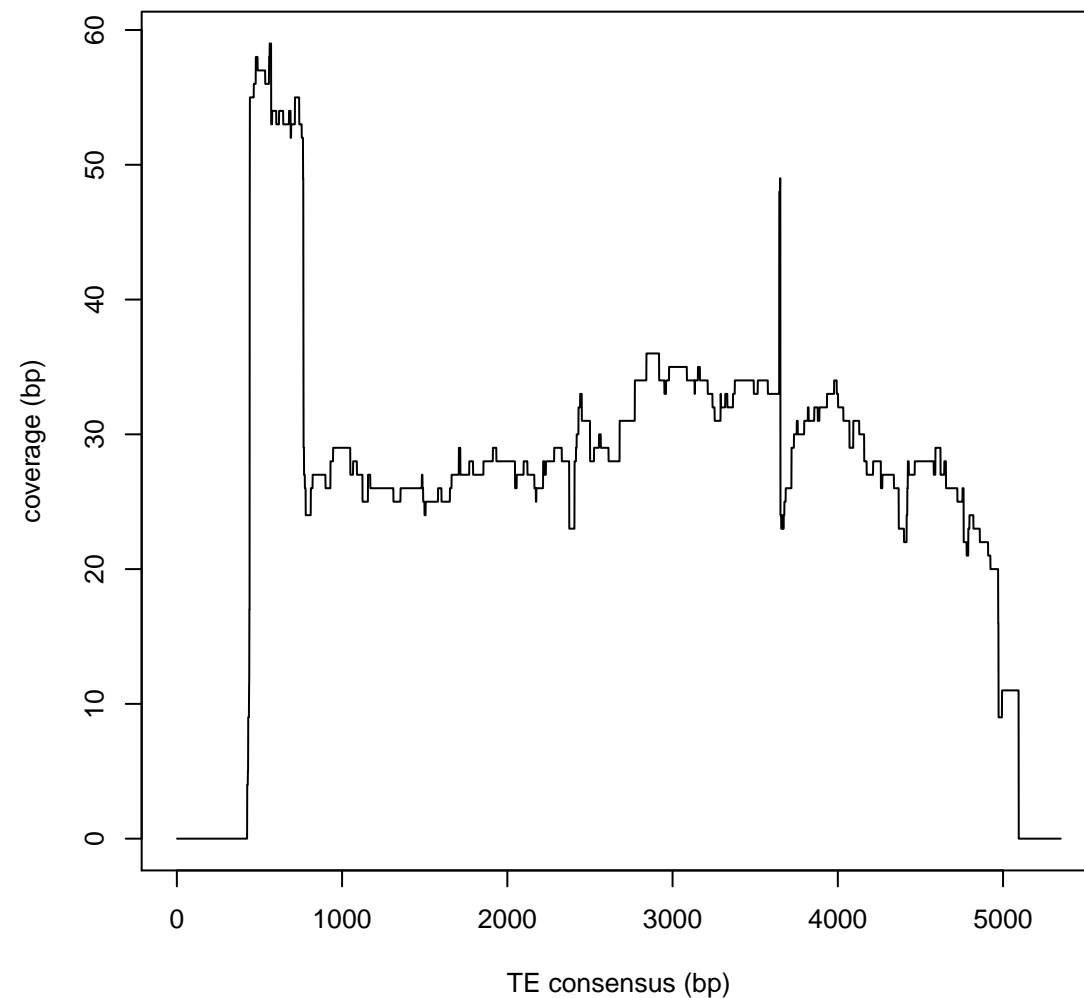


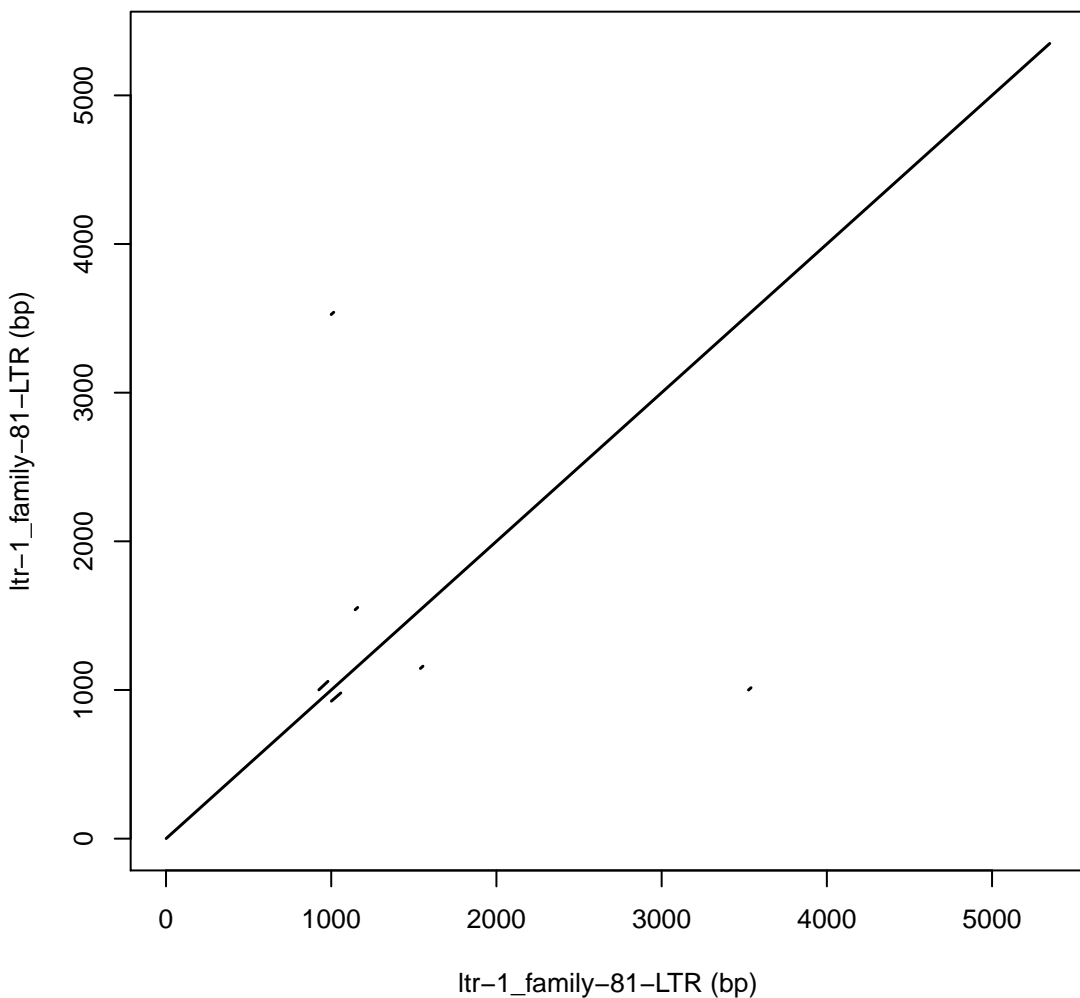
TE: ltr-1\_family-81-LTR  
consensus size: 5349bp; fragments: 176; full length: 0 (>=4814.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

