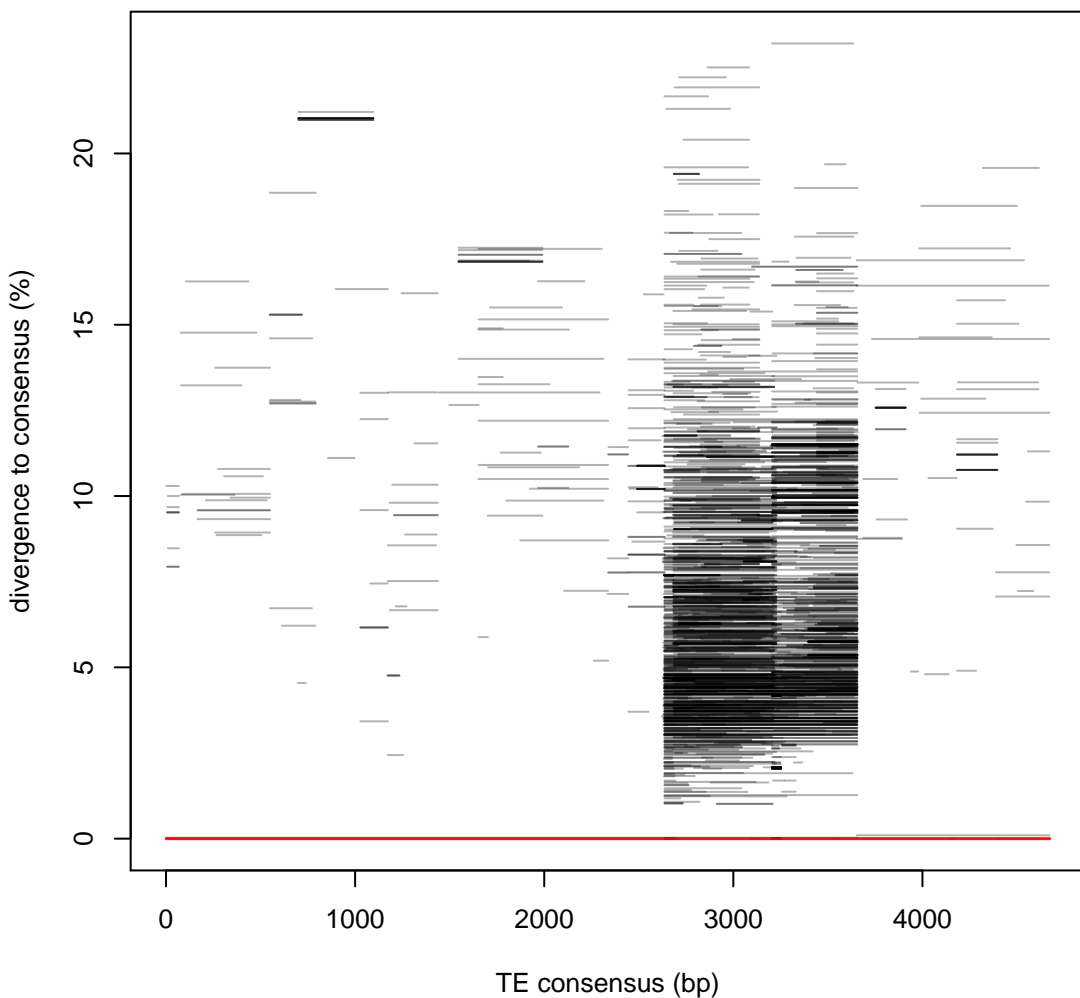
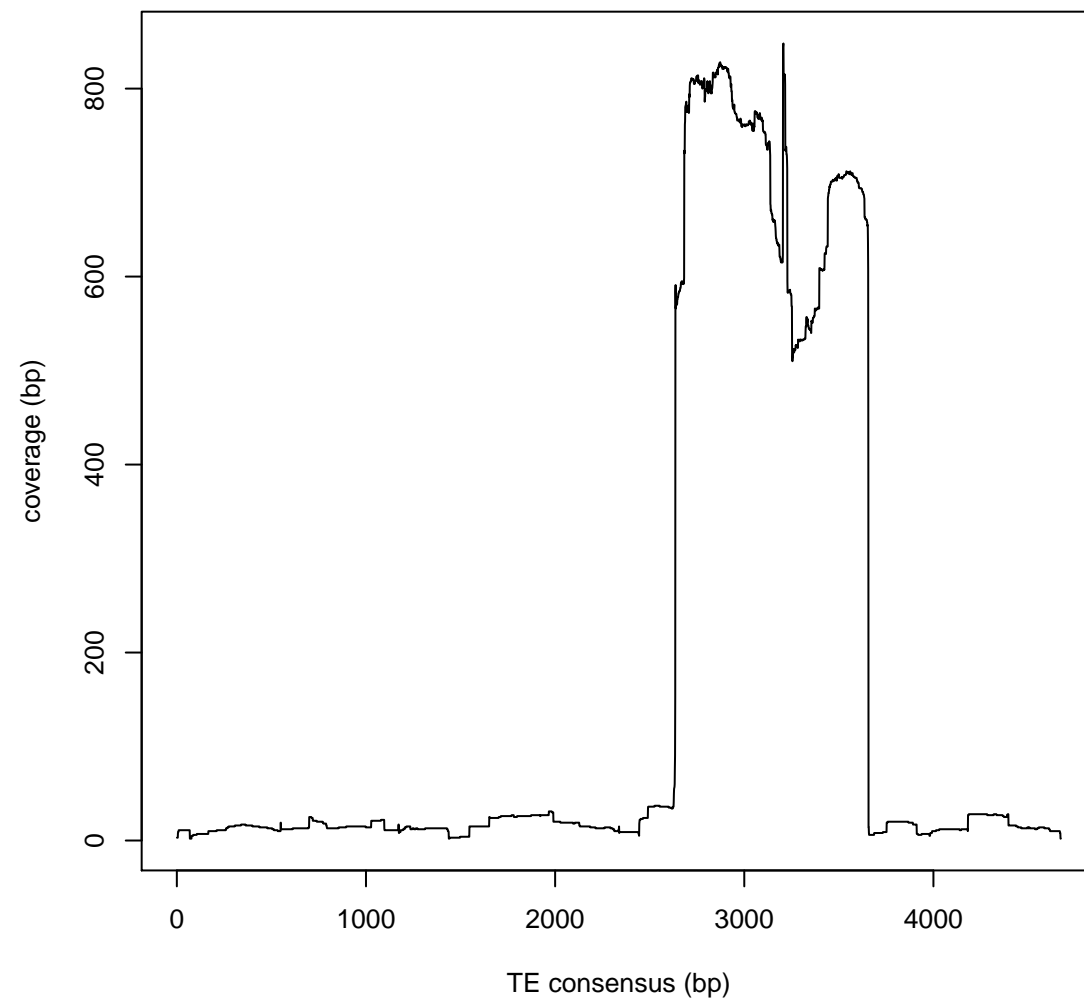


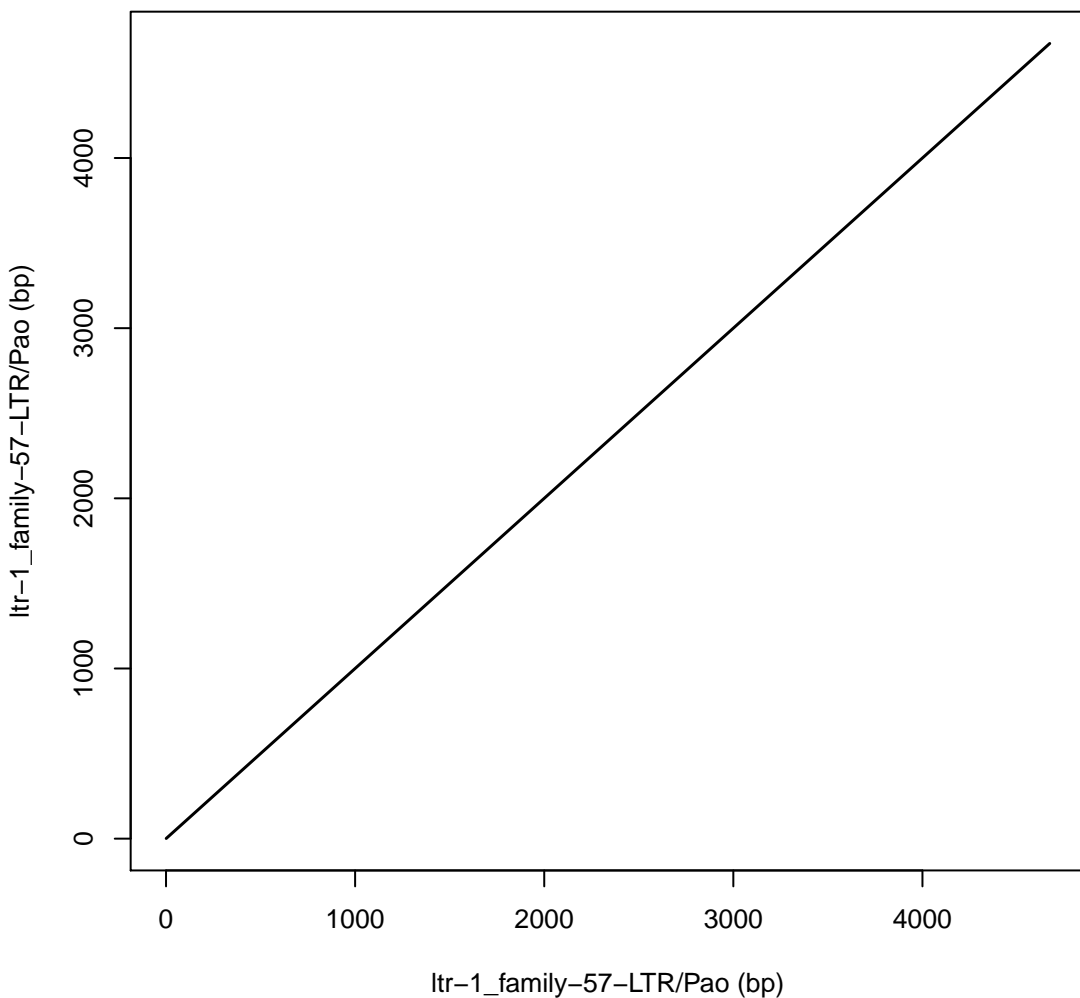
TE: ltr-1\_family-57-LTR/Pao  
consensus size: 4673bp; fragments: 1969; full length: 1 (>=4205.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

