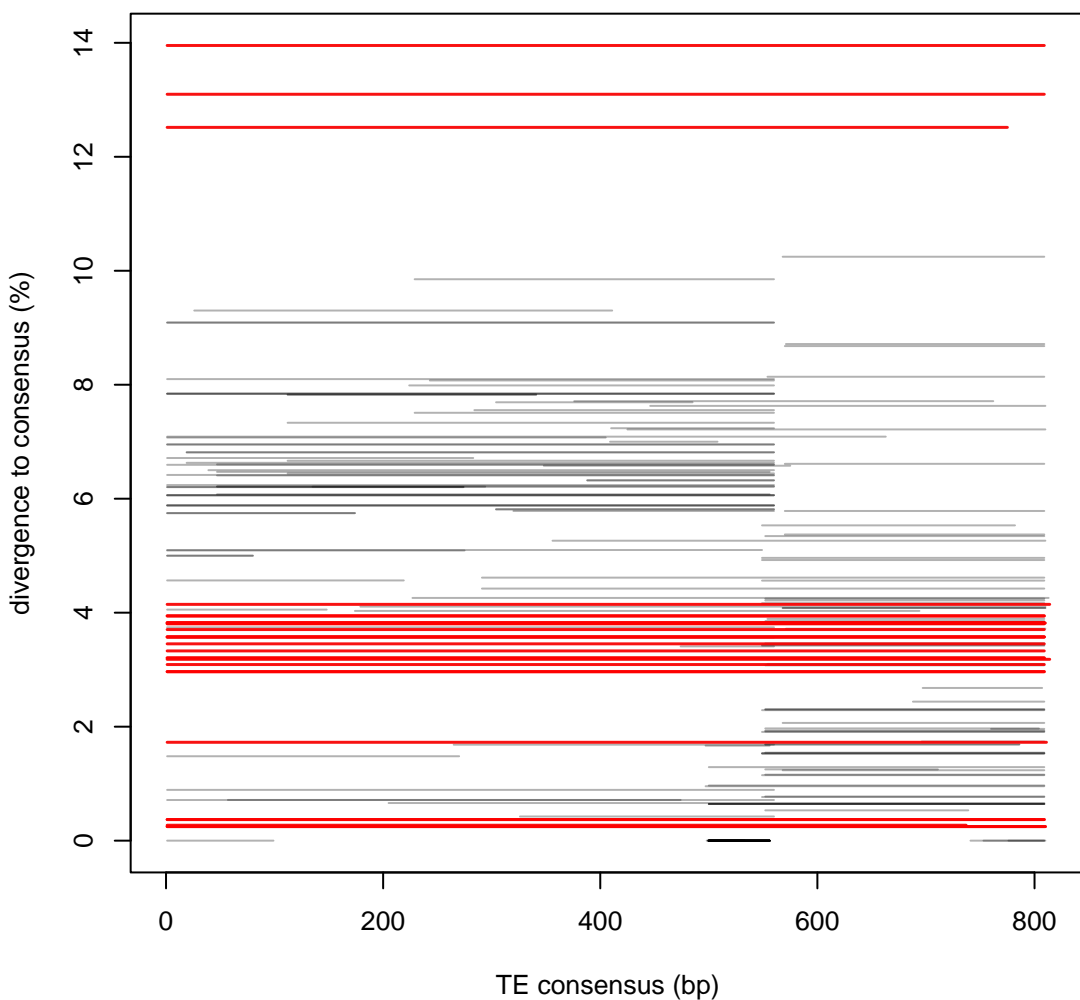
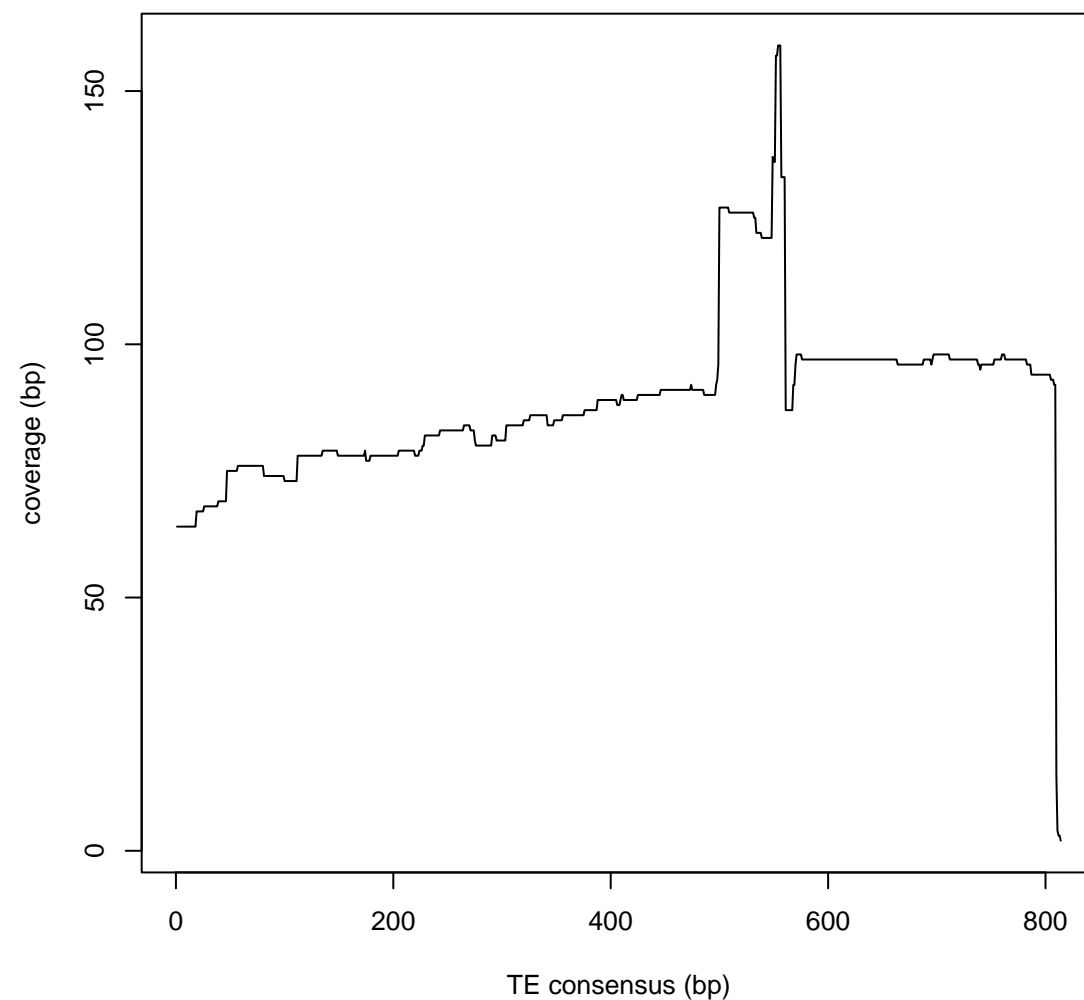


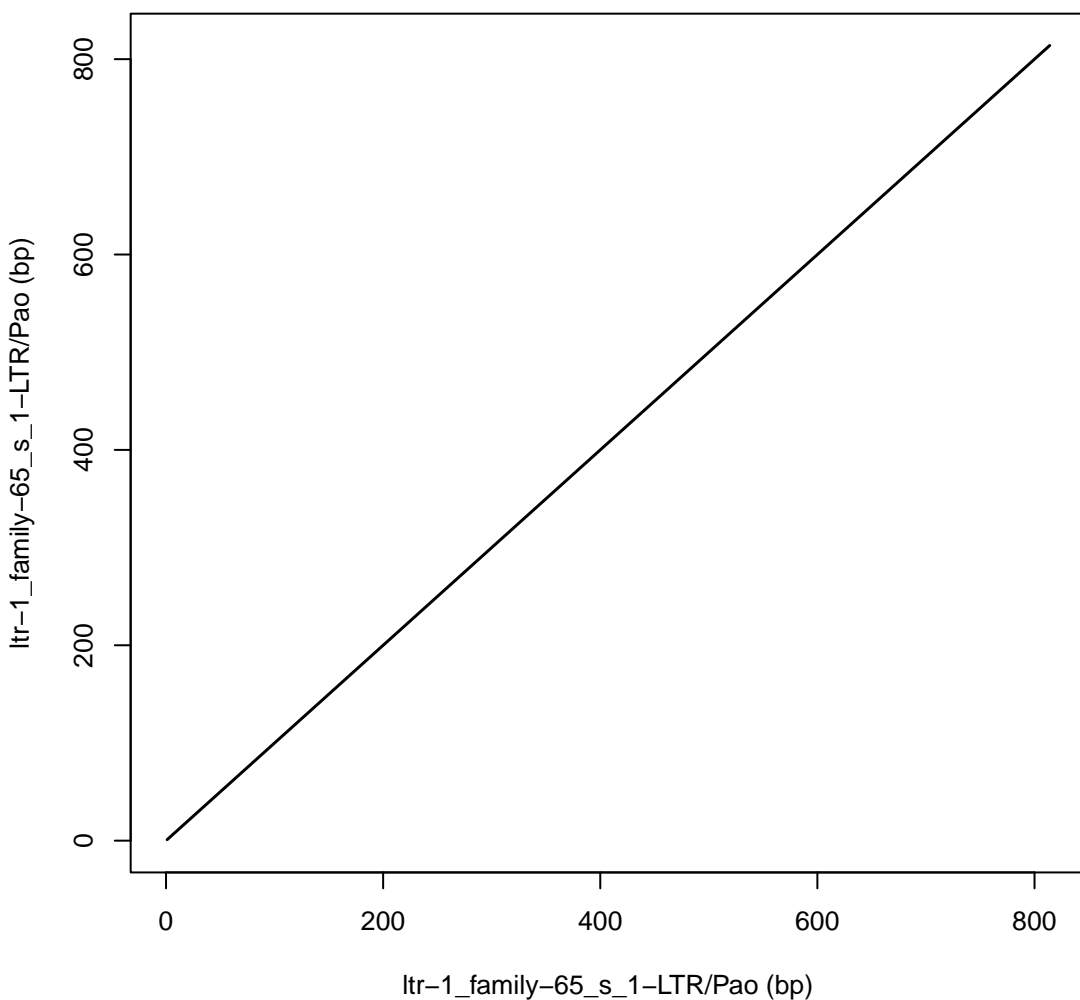
TE: ltr-1\_family-65\_s\_1-LTR/Pao  
consensus size: 814bp; fragments: 204; full length: 28 (>=732.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

