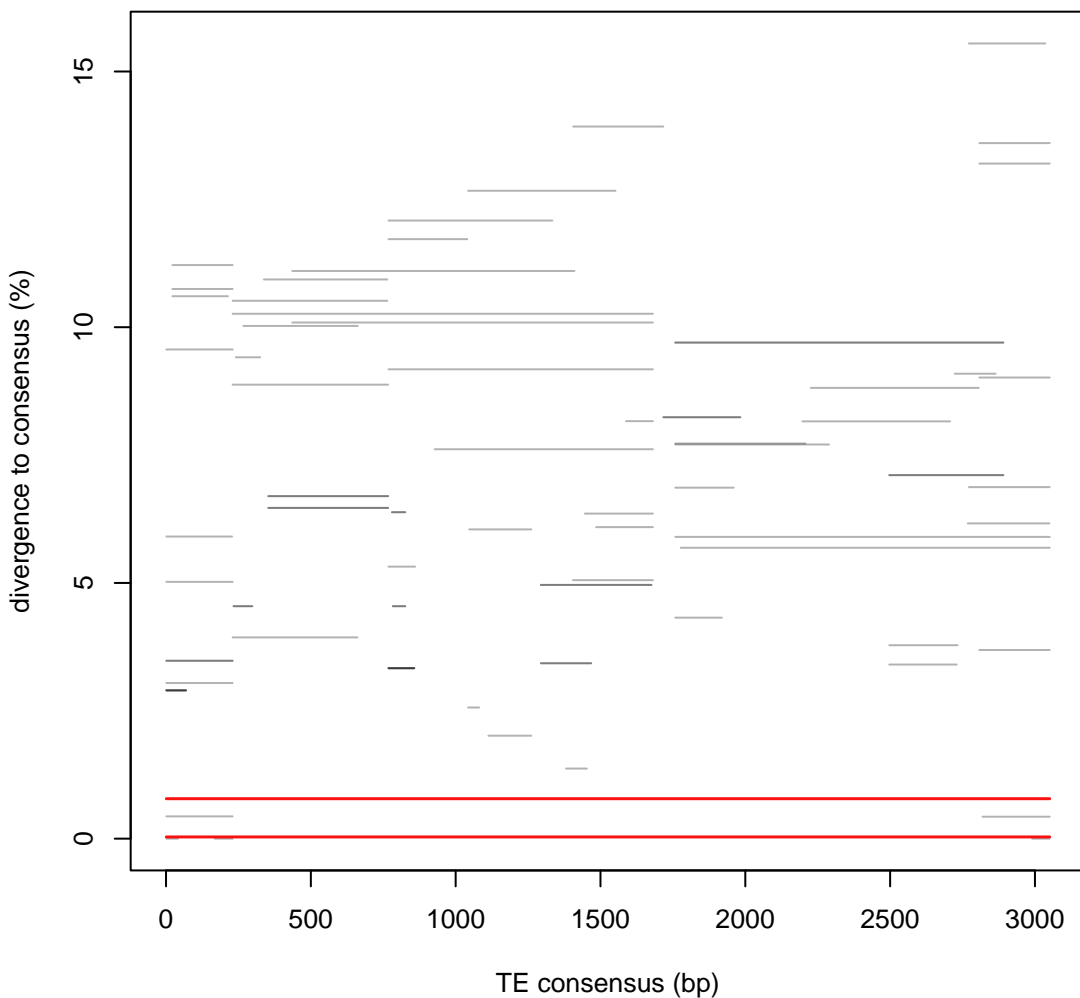
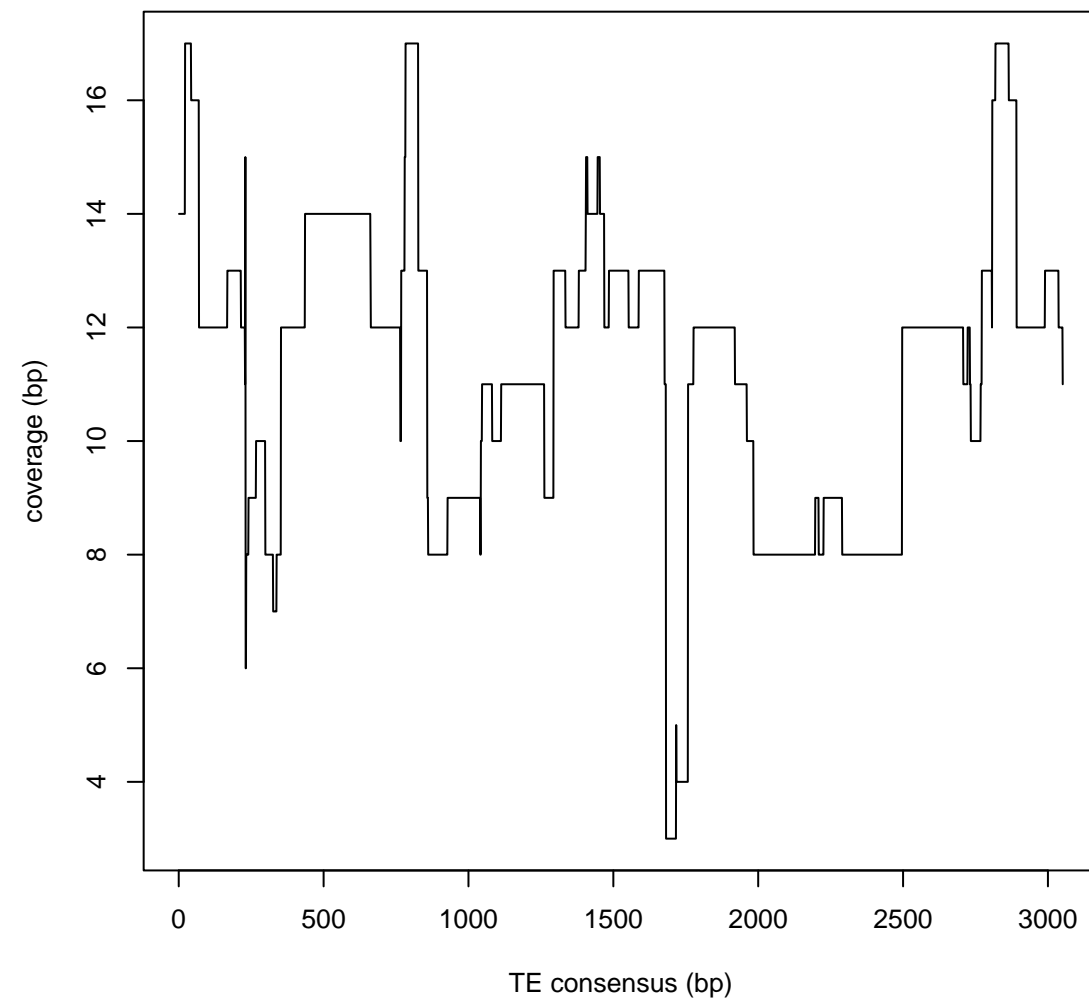


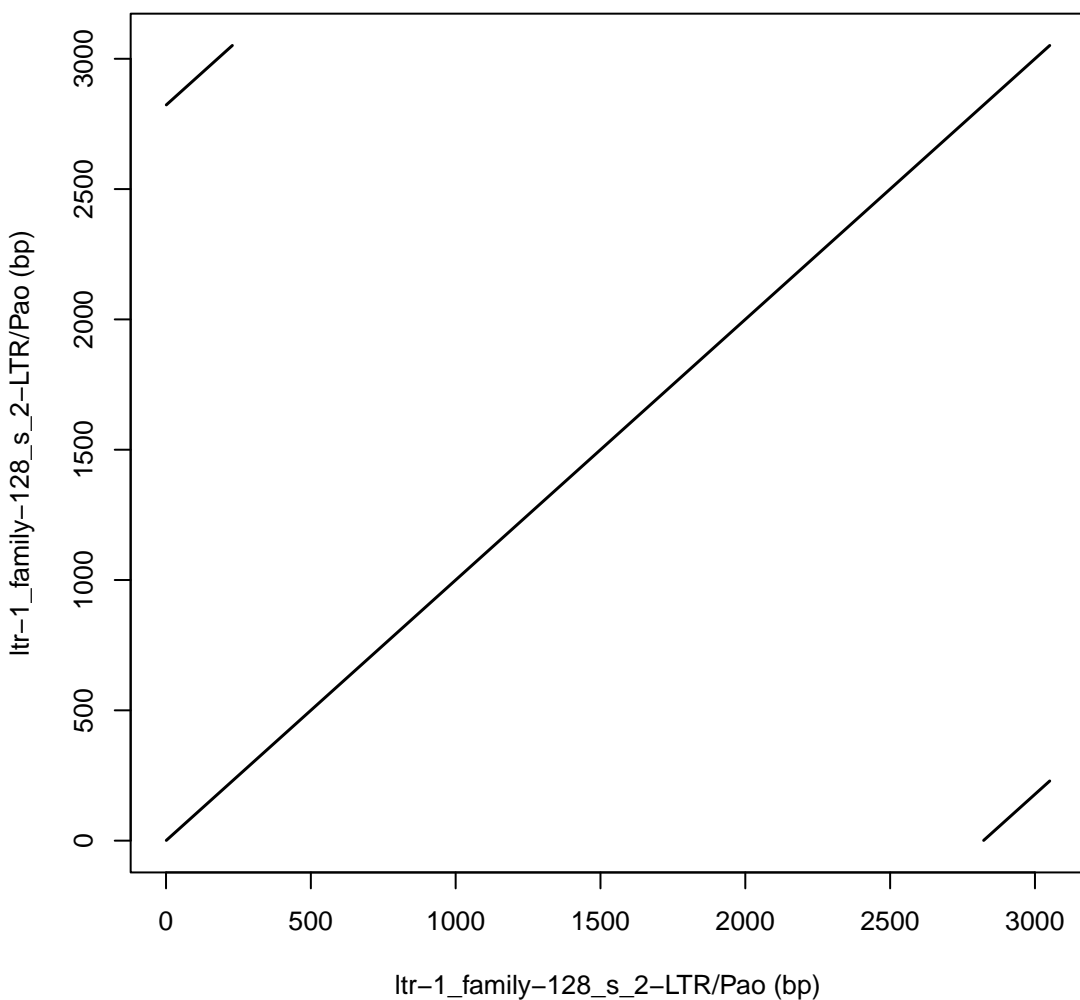
TE: ltr-1\_family-128\_s\_2-LTR/Pao  
consensus size: 3051bp; fragments: 86; full length: 2 (>=2745.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

