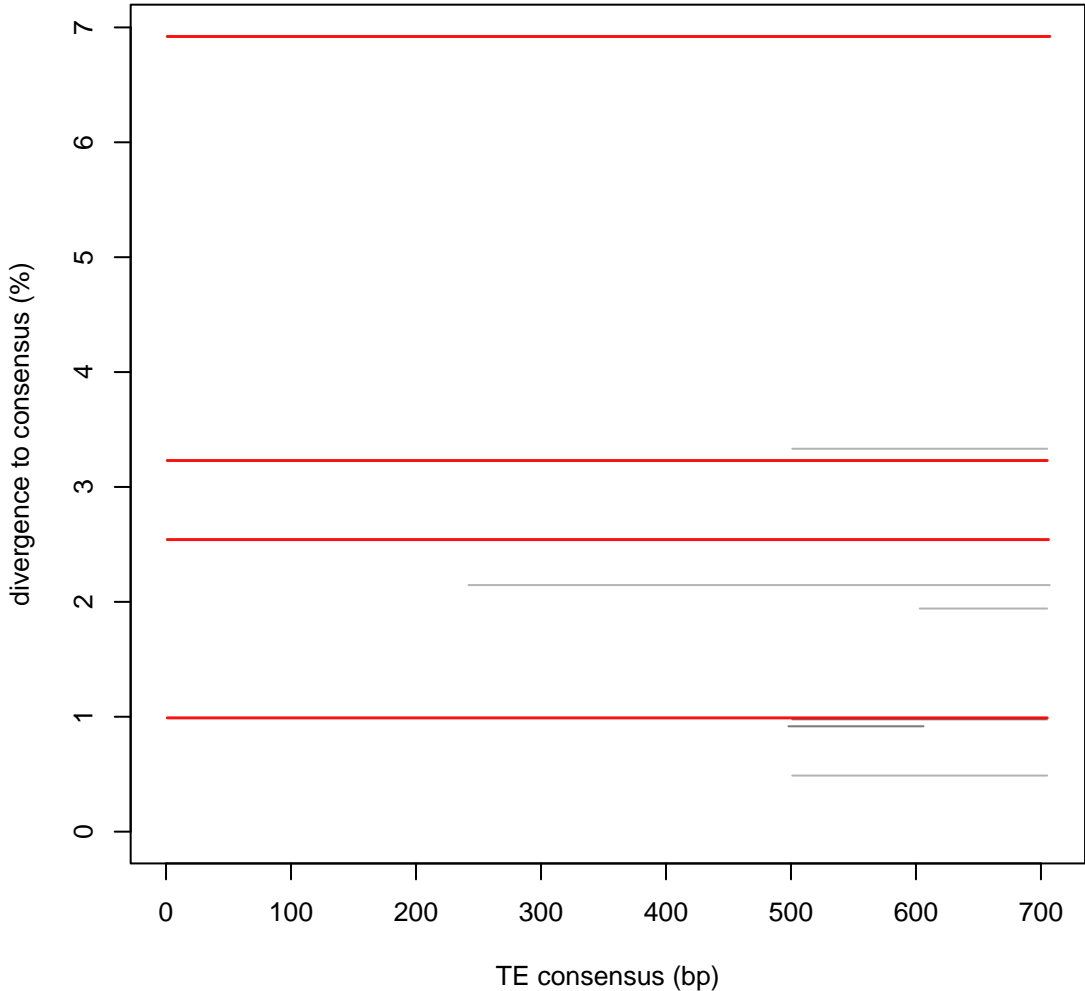
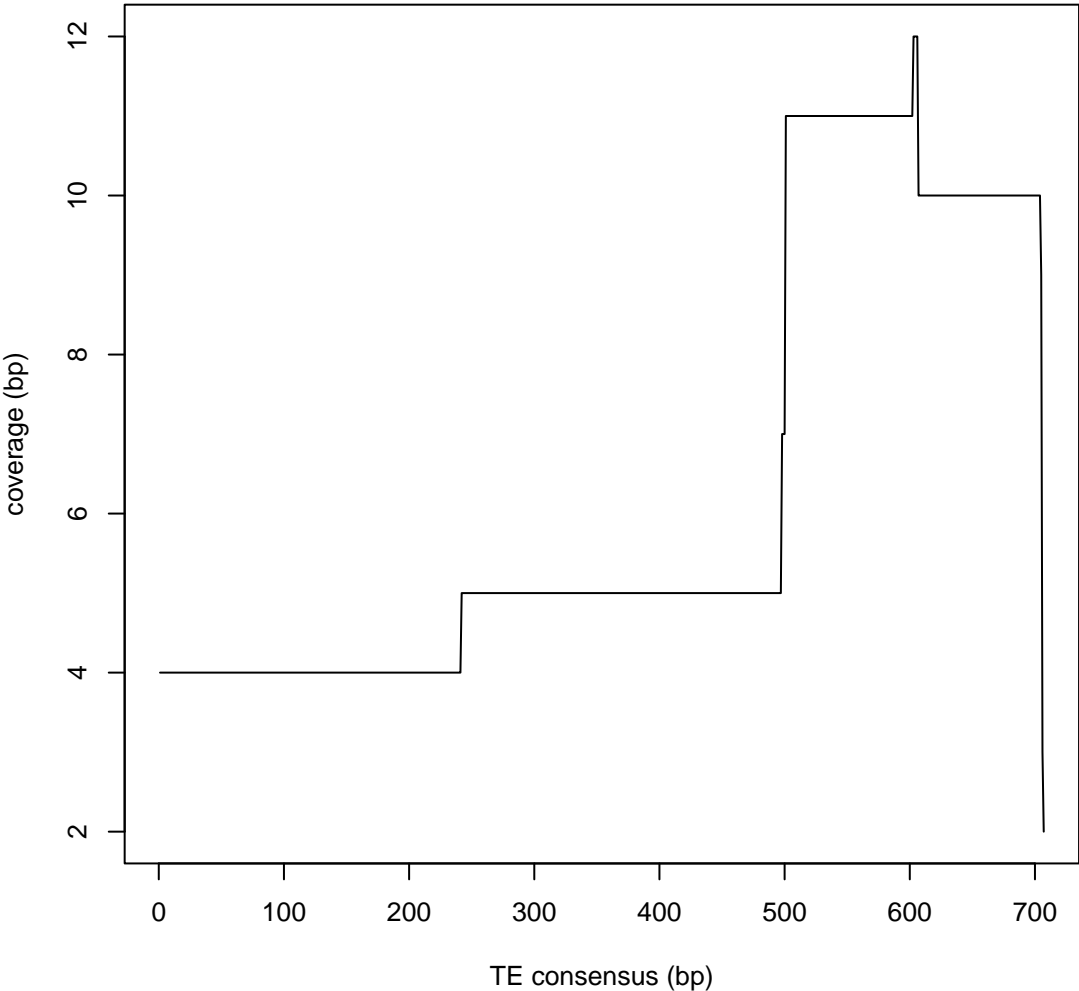


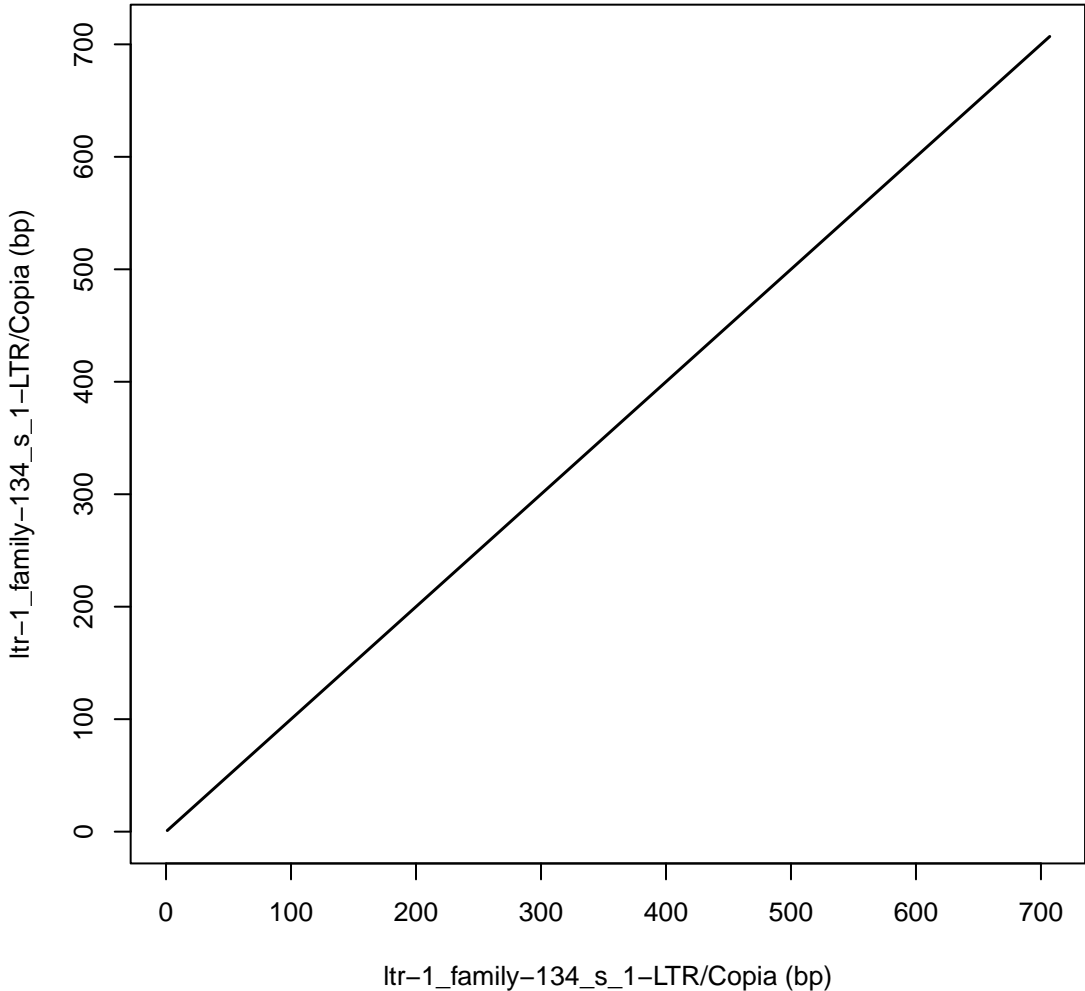
TE: ltr-1\_family-134\_s\_1-LTR/Copia  
consensus size: 707bp; fragments: 12; full length: 4 (>=636.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

