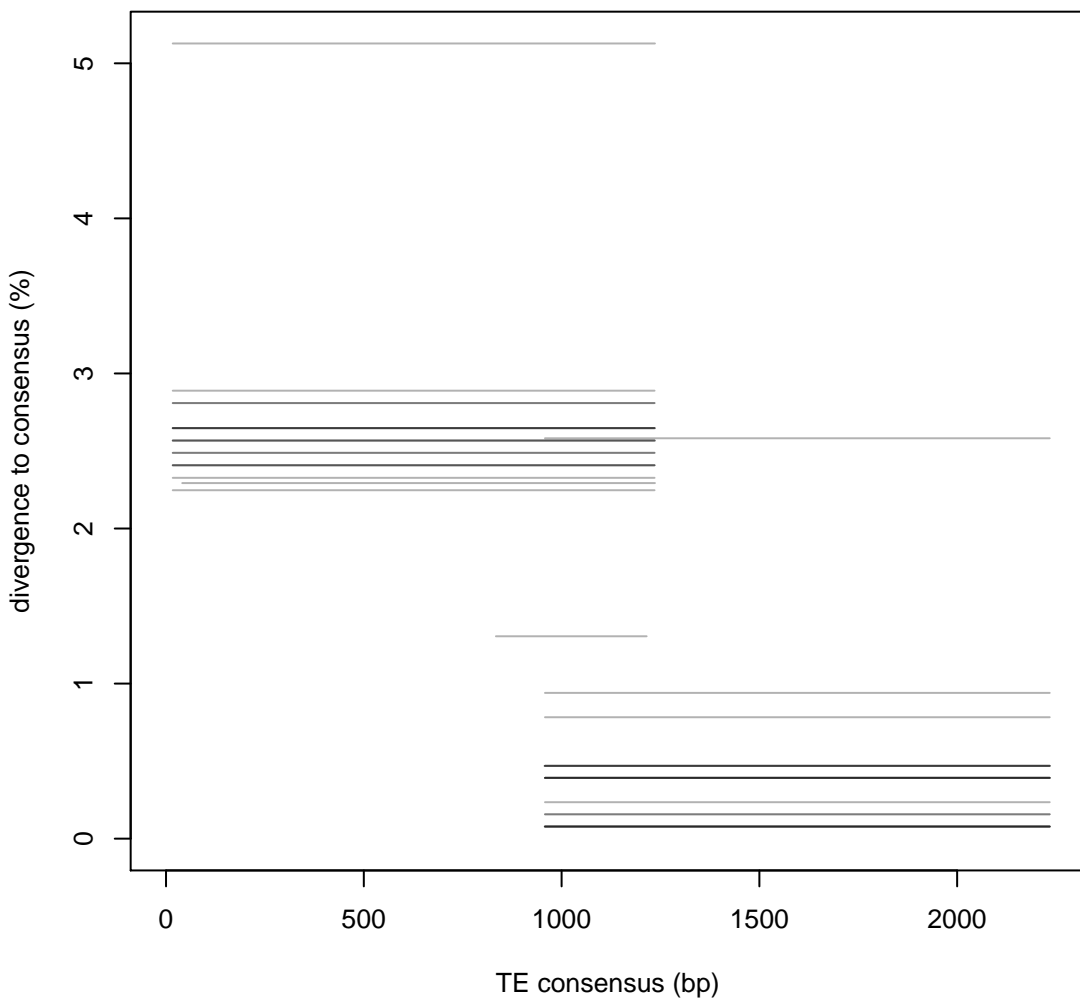
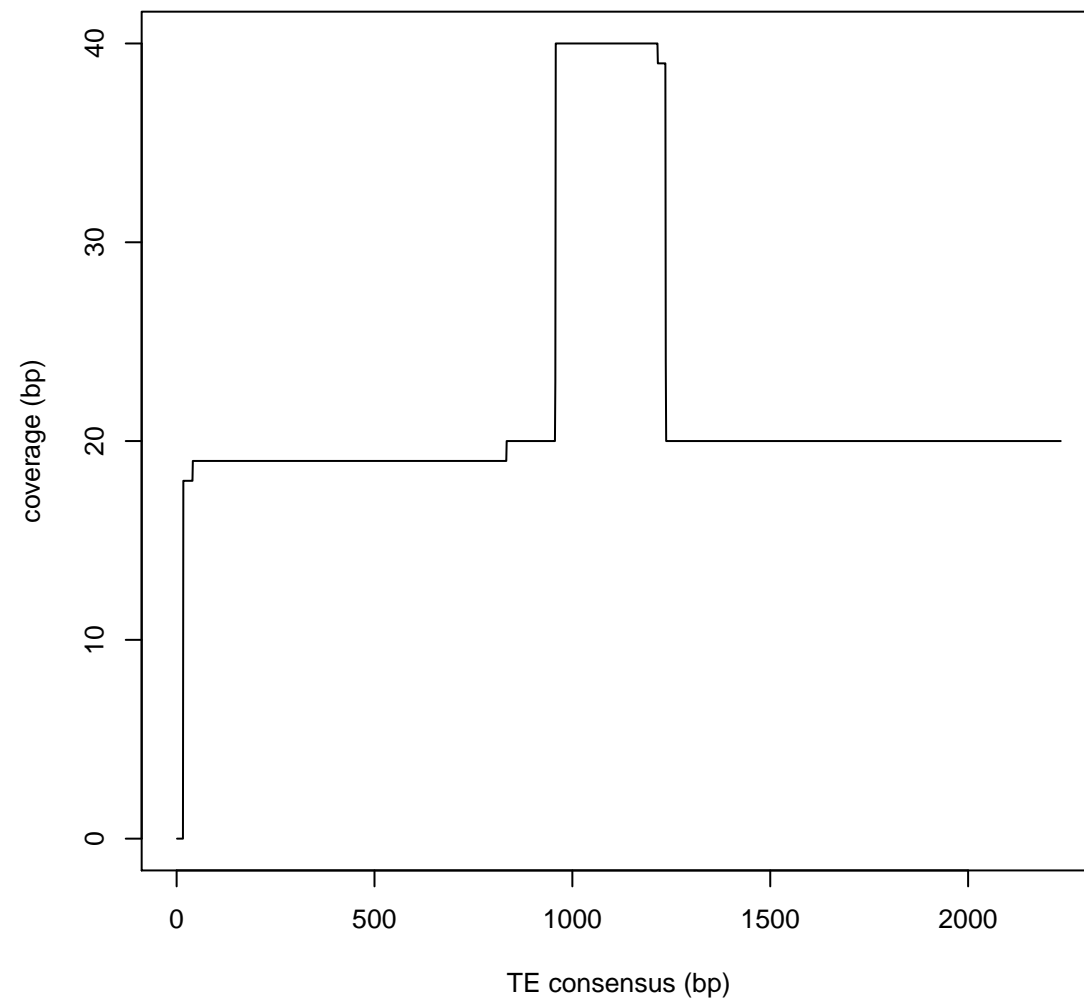


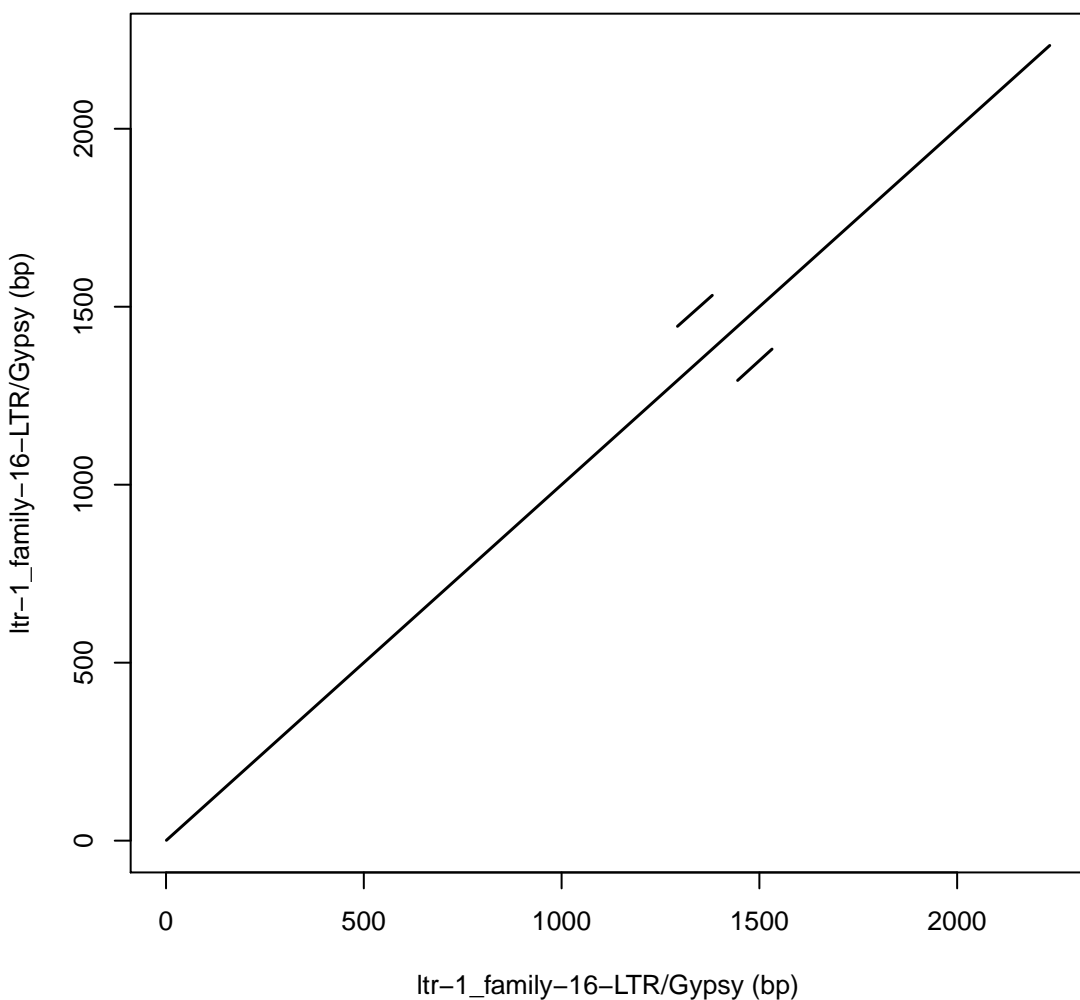
TE: ltr-1\_family-16-LTR/Gypsy  
consensus size: 2234bp; fragments: 40; full length: 0 (>=2010.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

