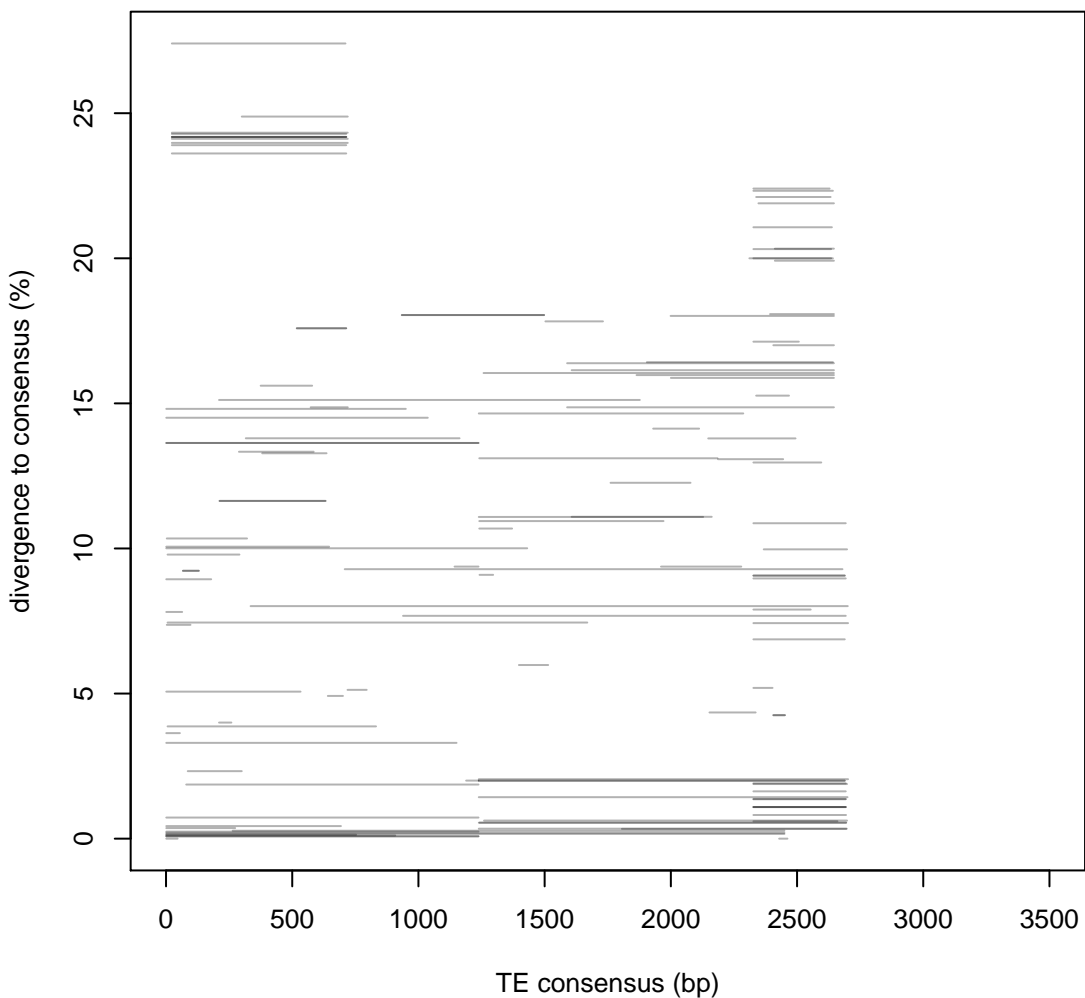
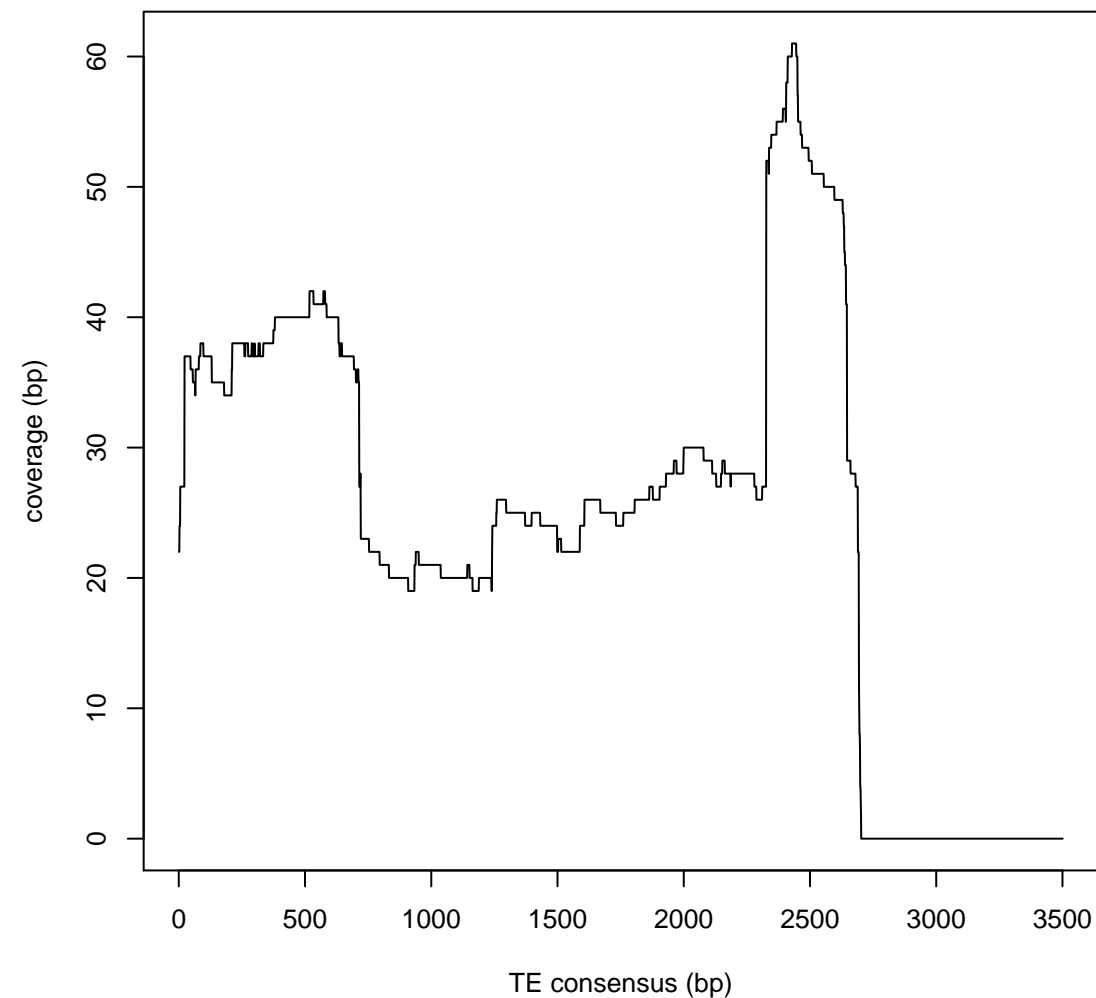


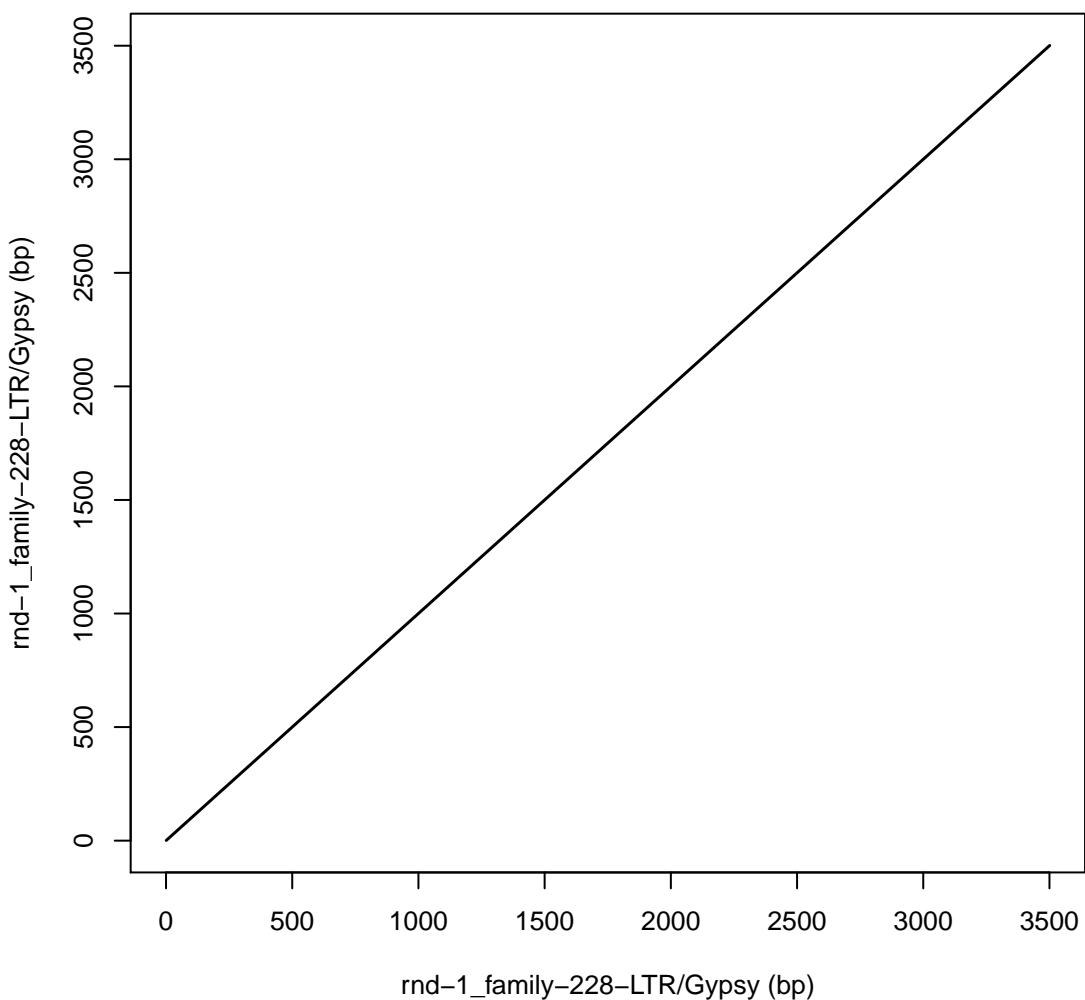
TE: rnd-1\_family-228-LTR/Gypsy  
consensus size: 3501bp; fragments: 132; full length: 0 (>=3150.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

