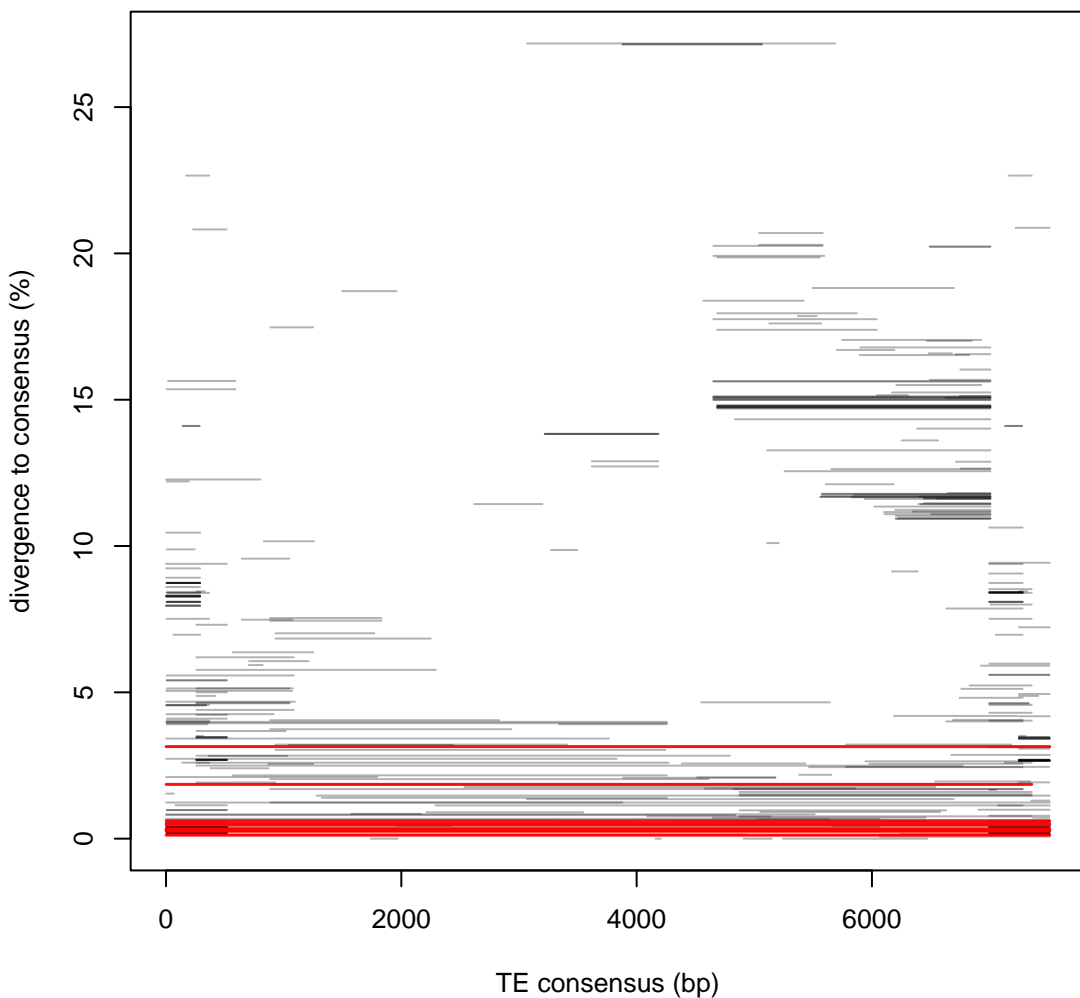
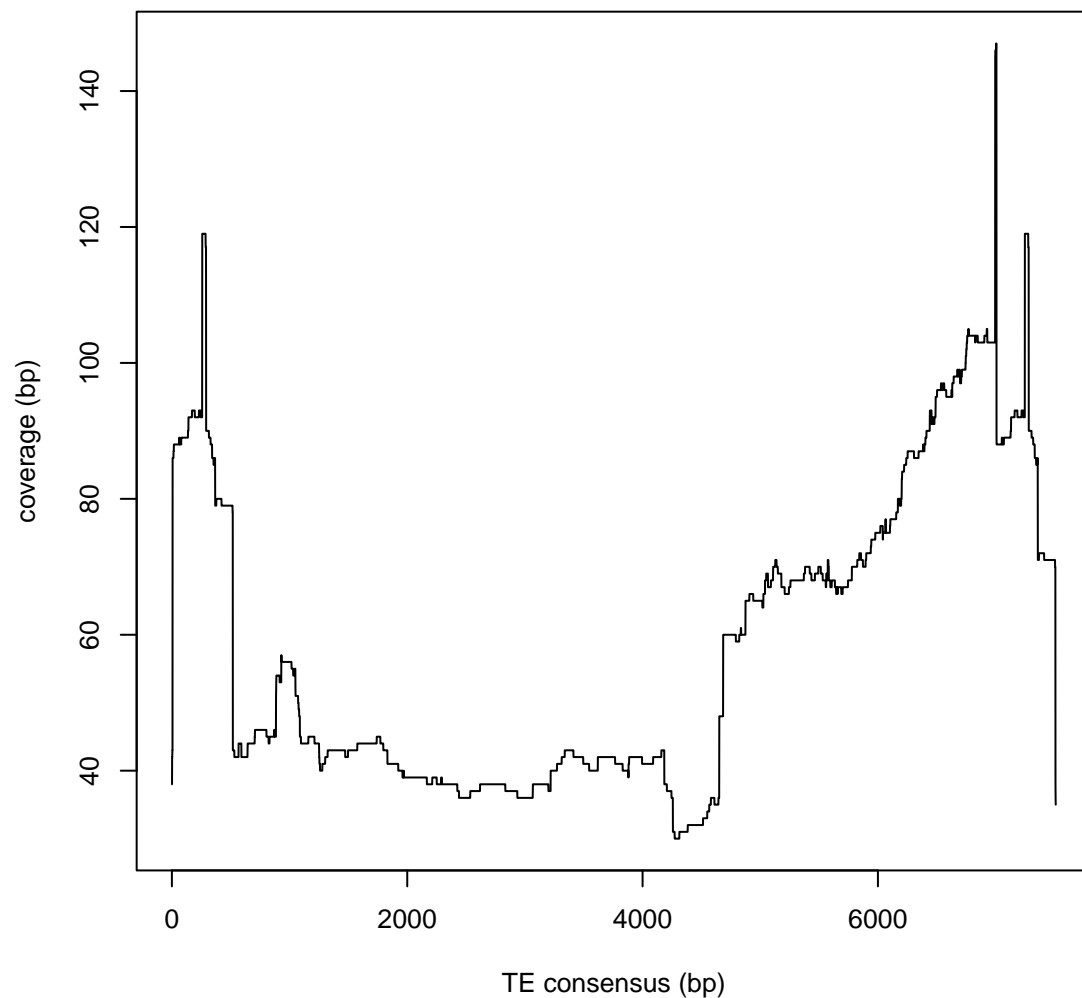


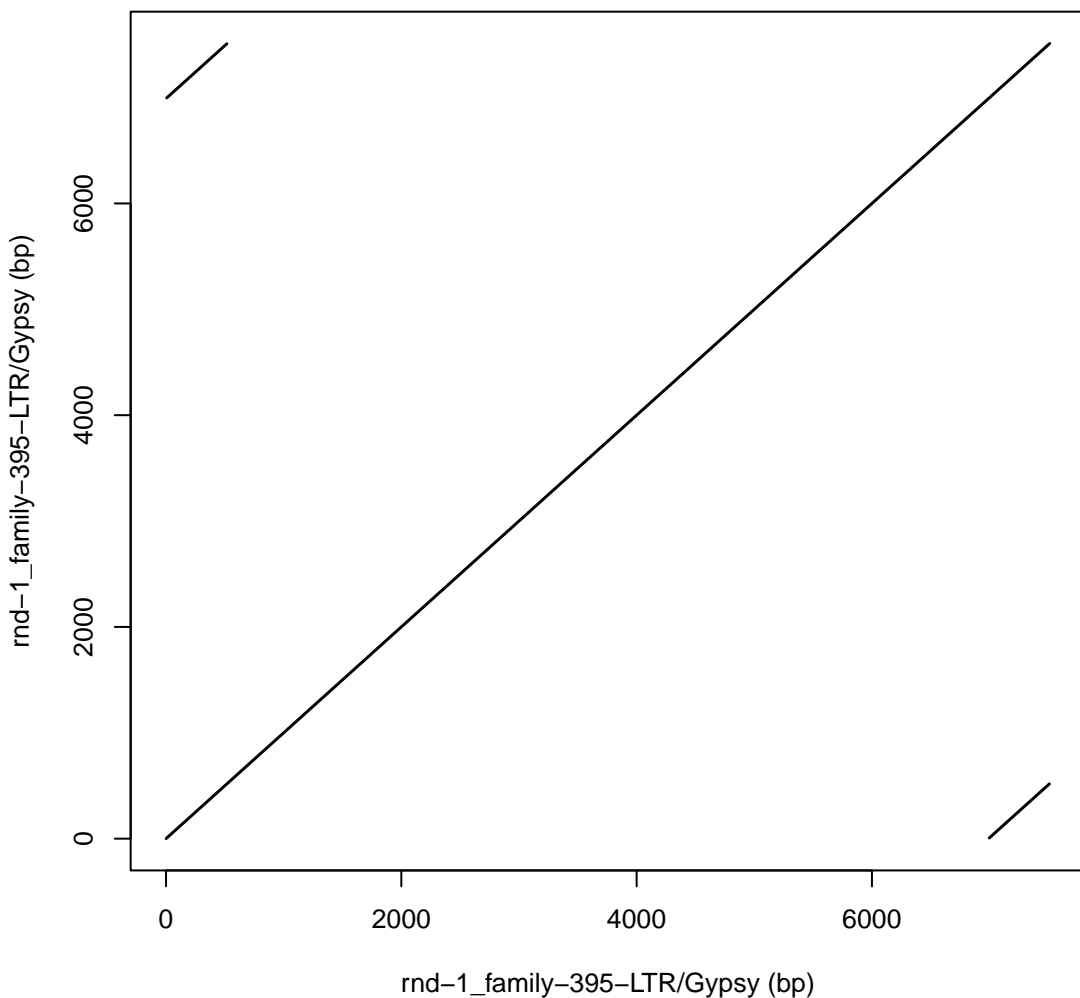
TE: rnd-1_family-395-LTR/Gypsy
consensus size: 7511bp; fragments: 378; full length: 10 (>=6759.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

