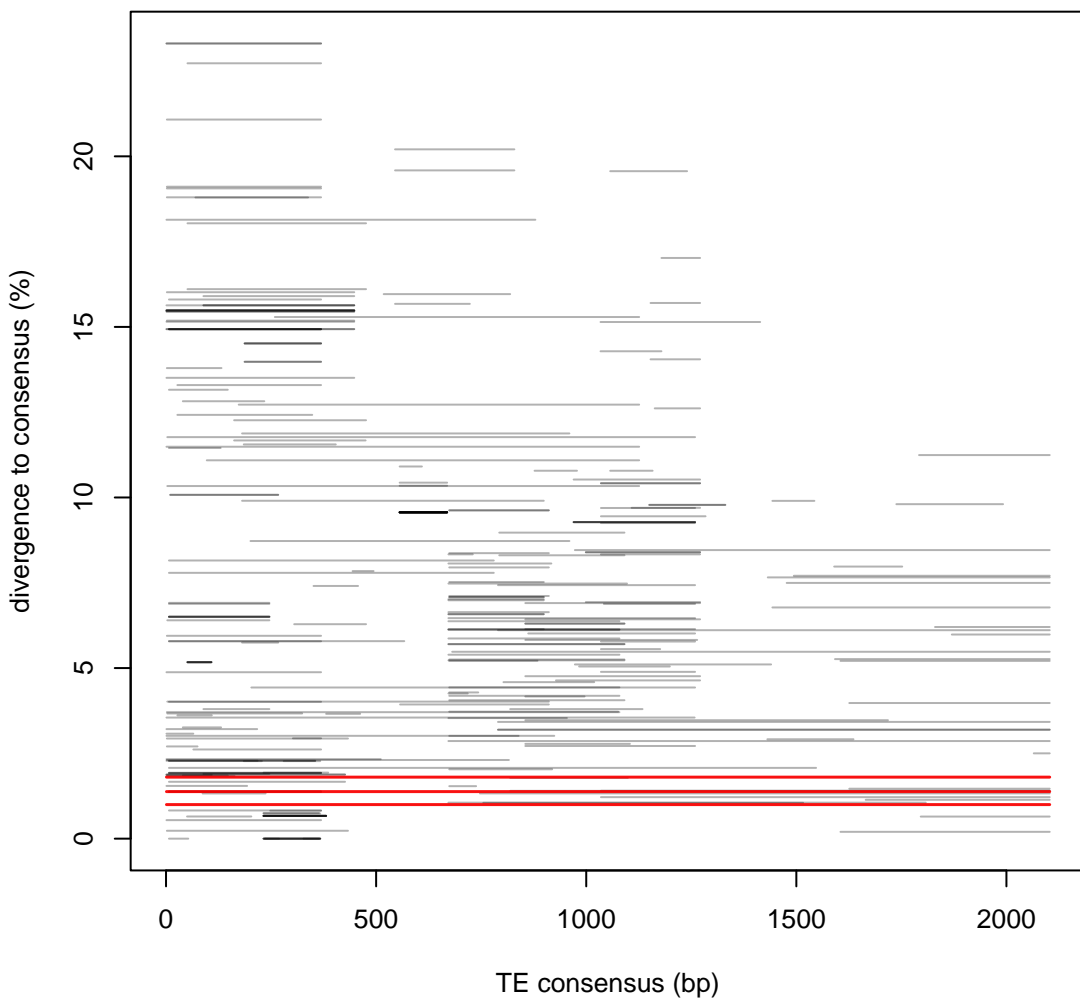
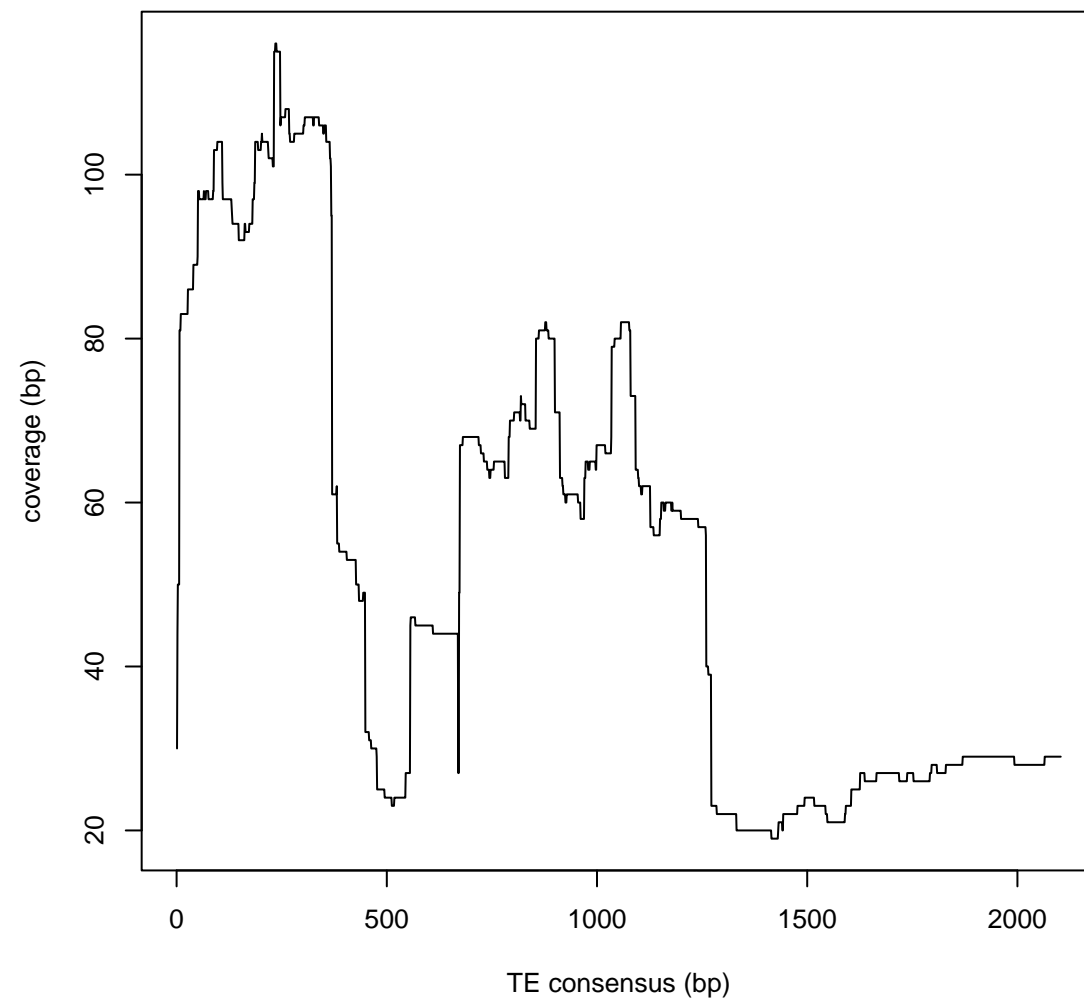


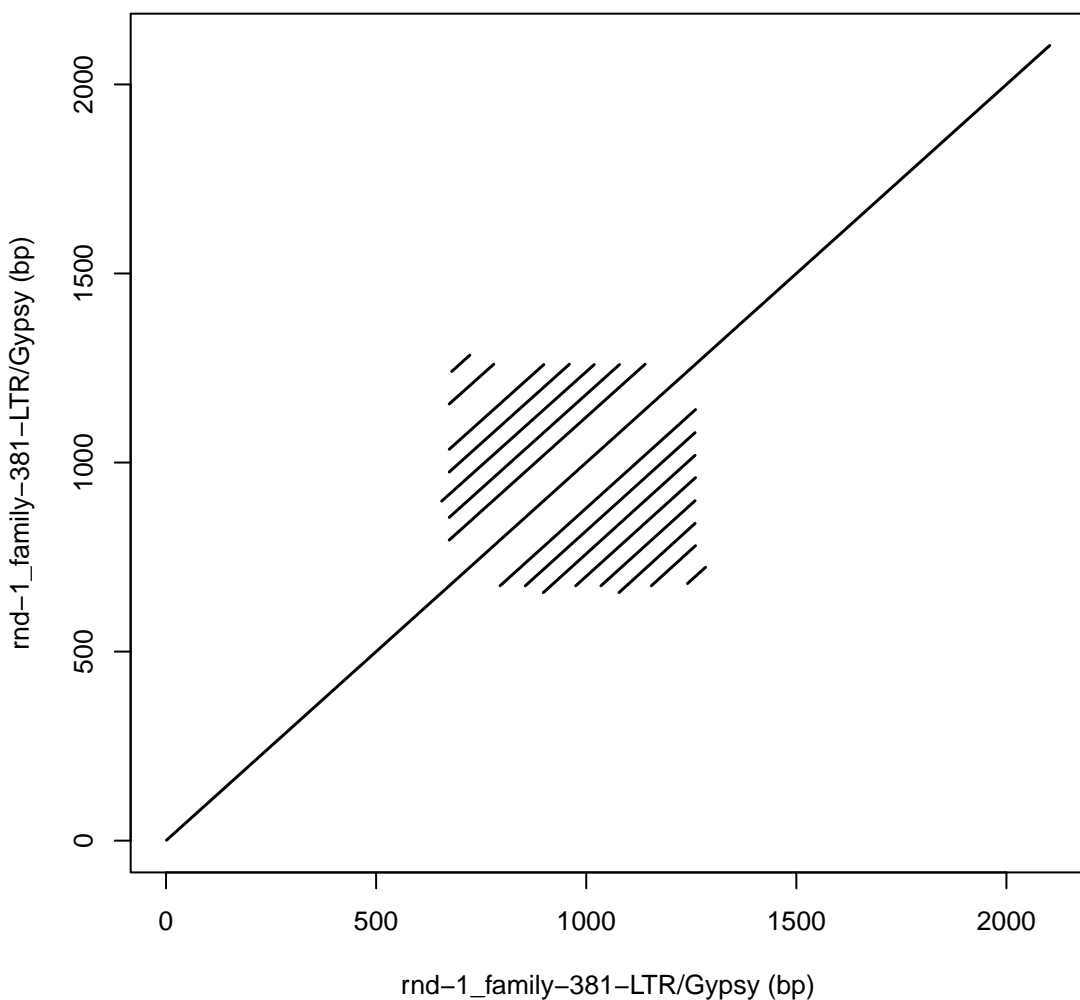
TE: rnd-1_family-381-LTR/Gypsy
consensus size: 2103bp; fragments: 291; full length: 3 (>=1892.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

