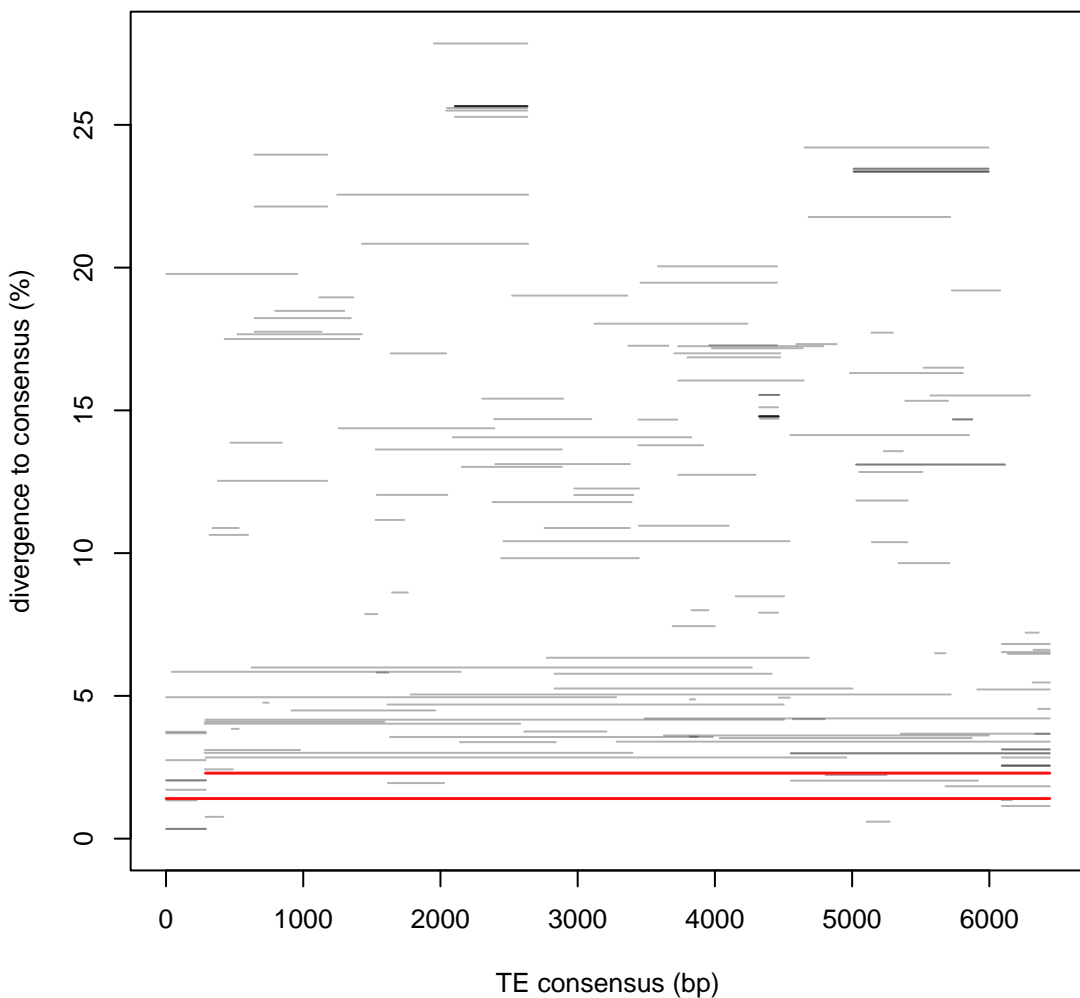
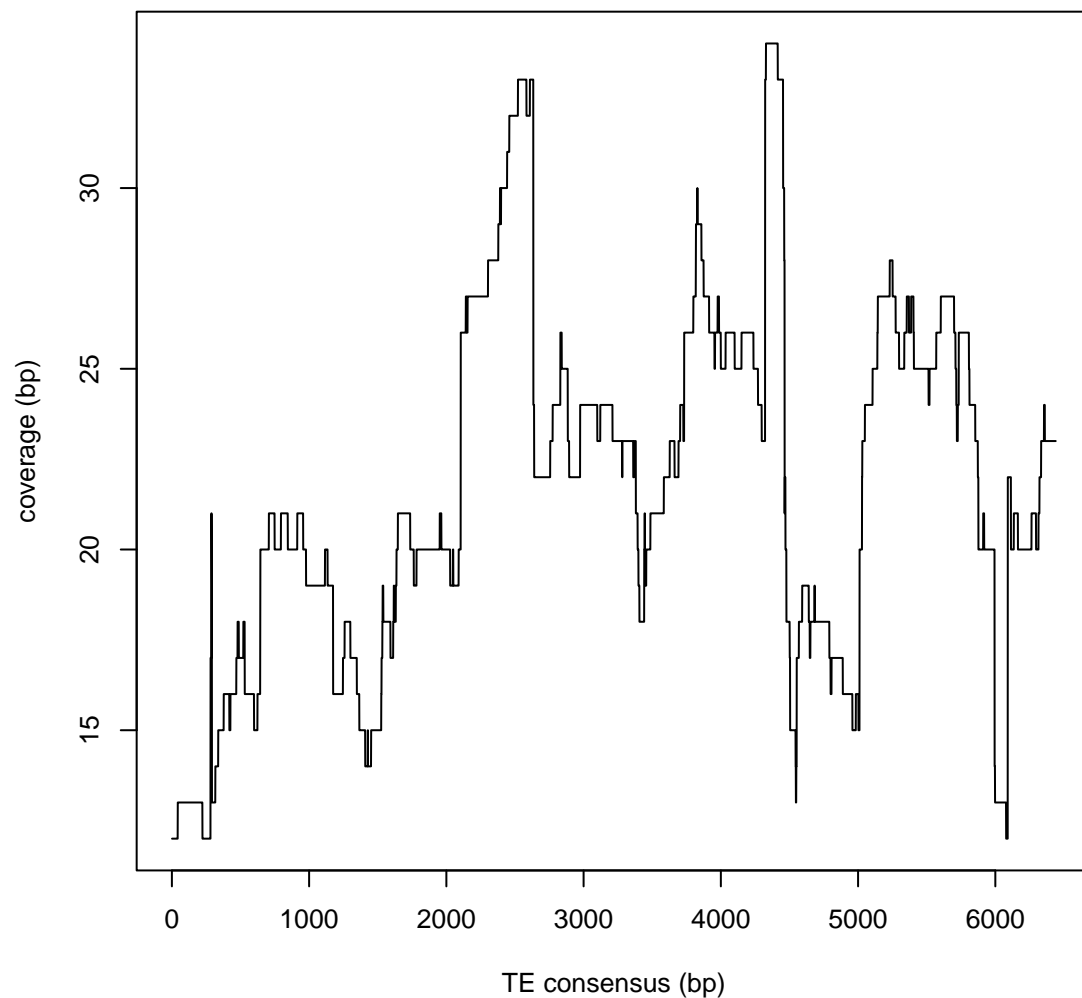


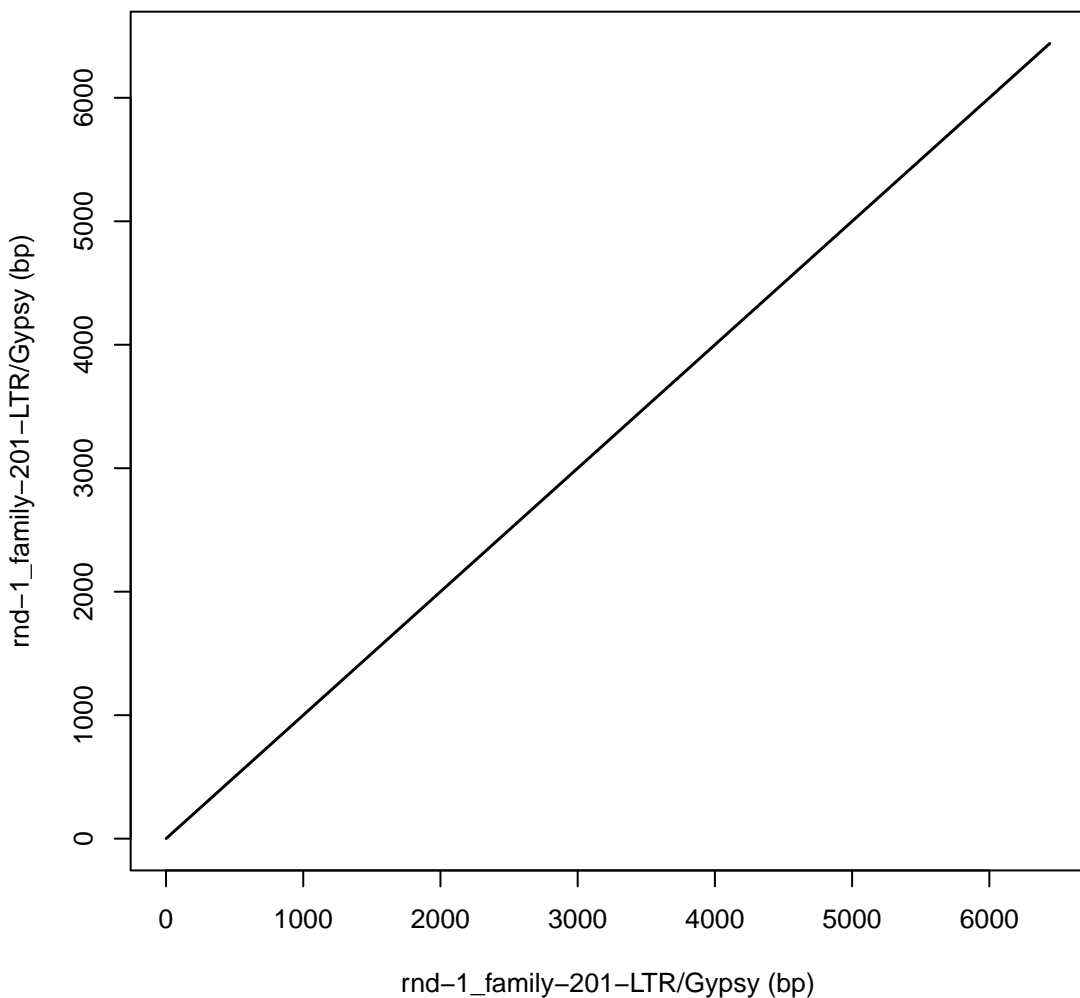
TE: rnd-1_family-201-LTR/Gypsy
consensus size: 6440bp; fragments: 163; full length: 2 (>=5796bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

