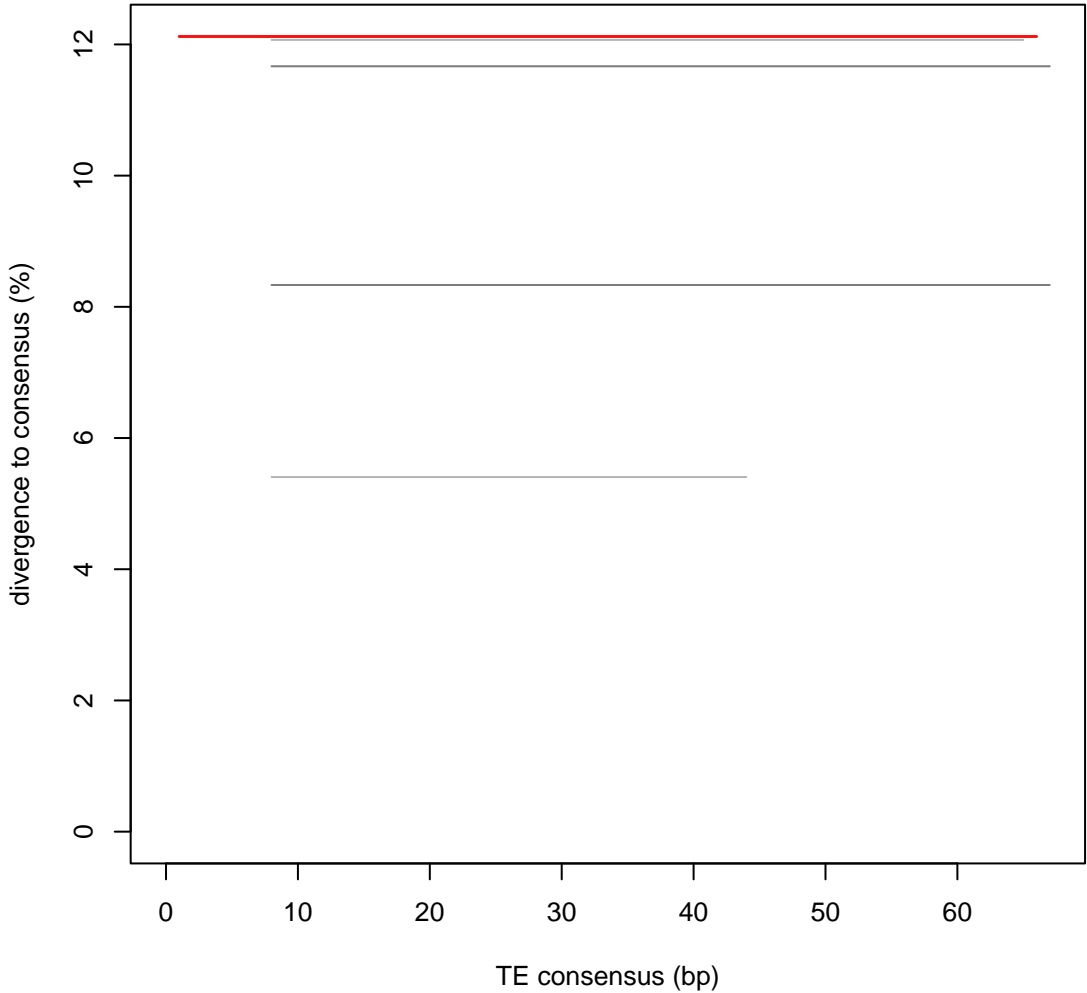
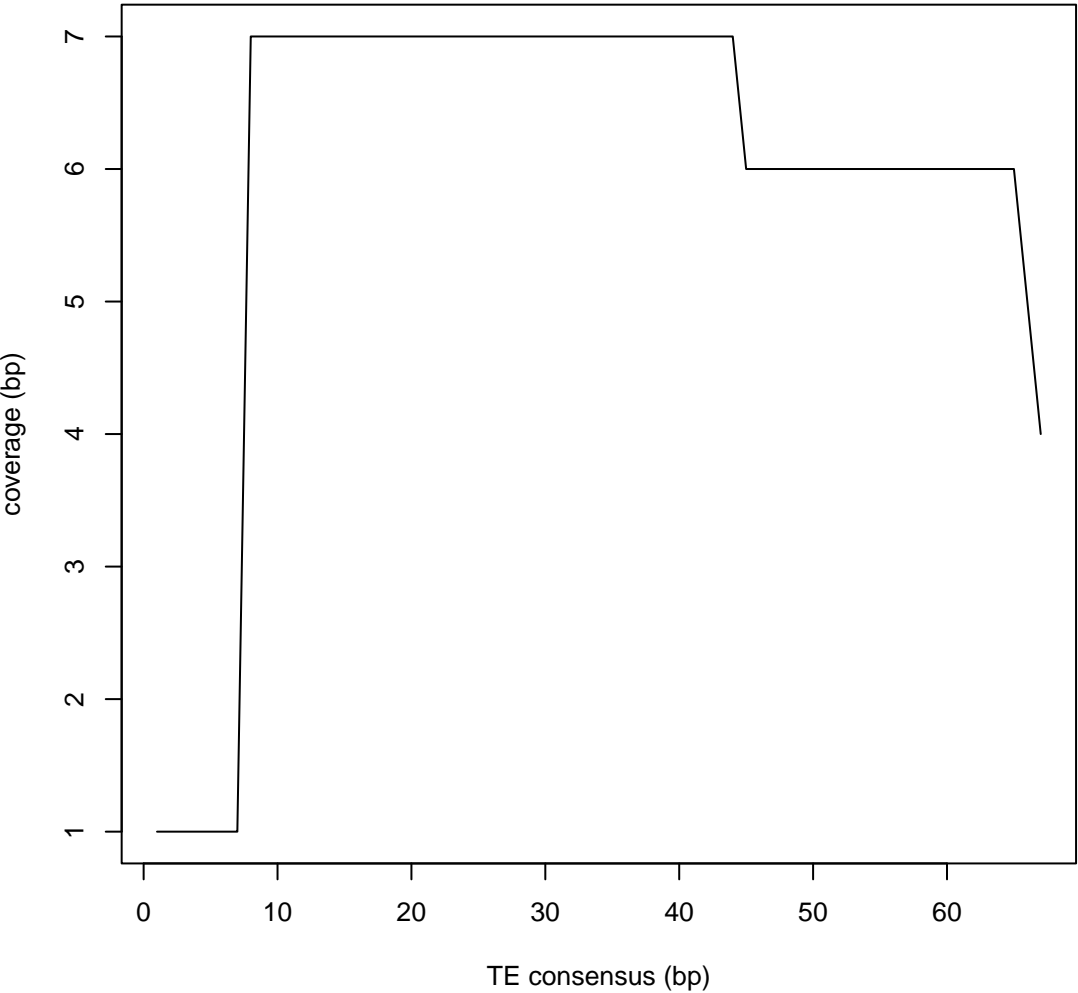


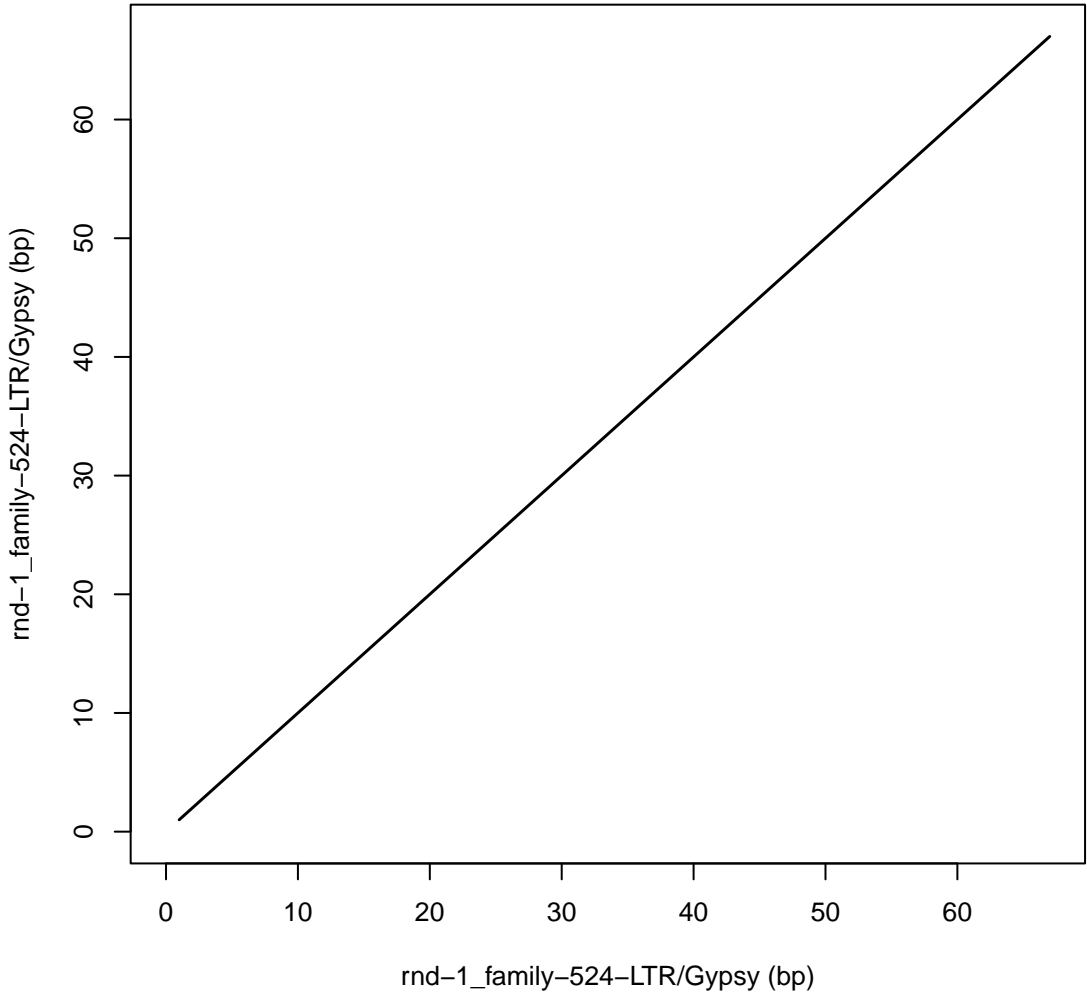
TE: rnd-1_family-524-LTR/Gypsy
consensus size: 67bp; fragments: 7; full length: 1 (>=60.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

