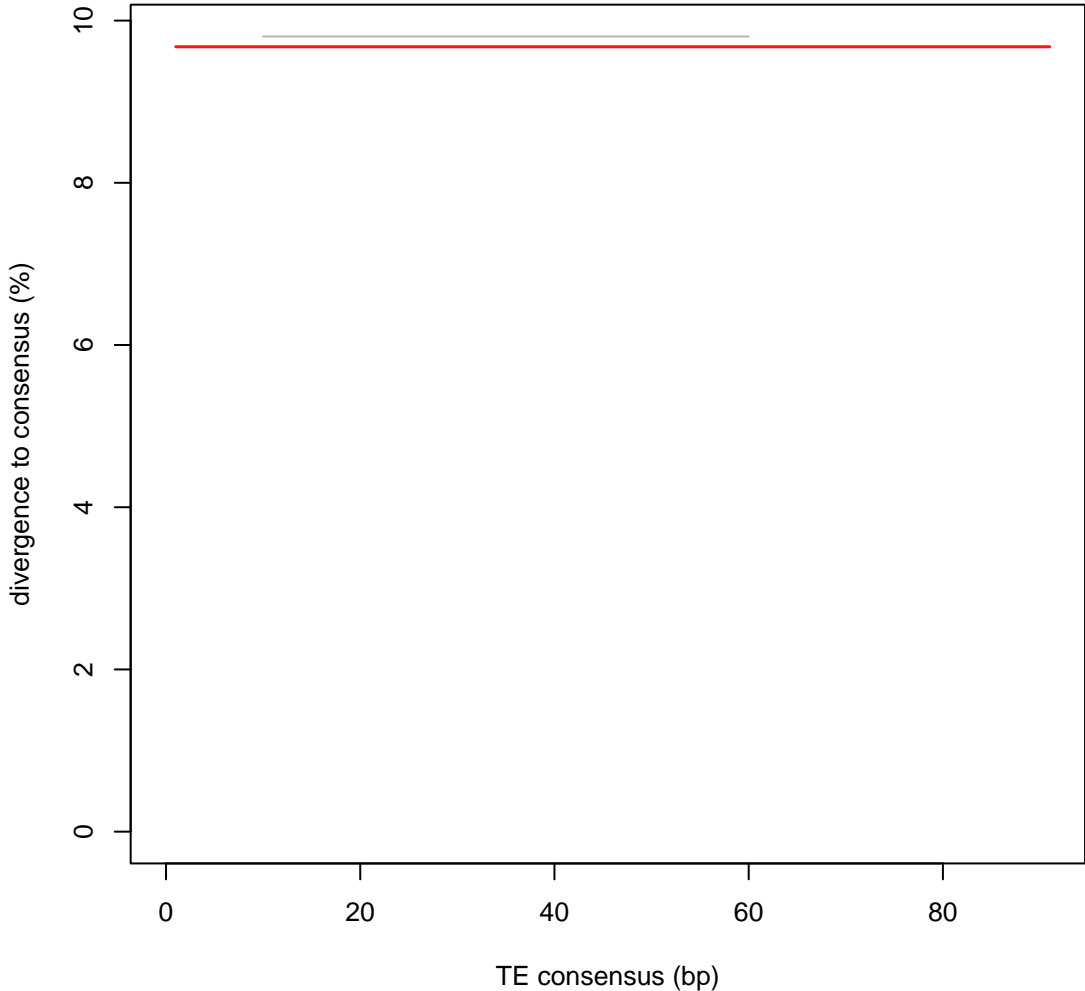
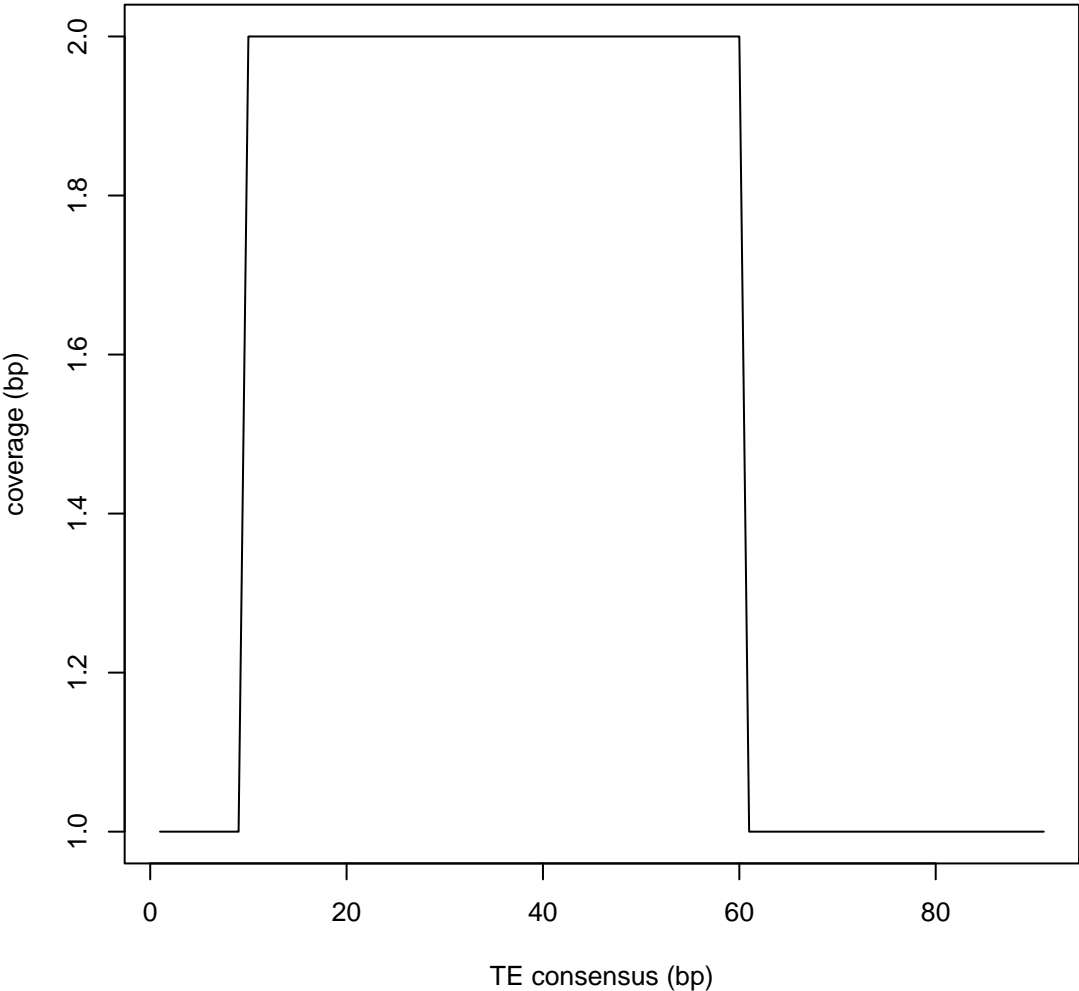


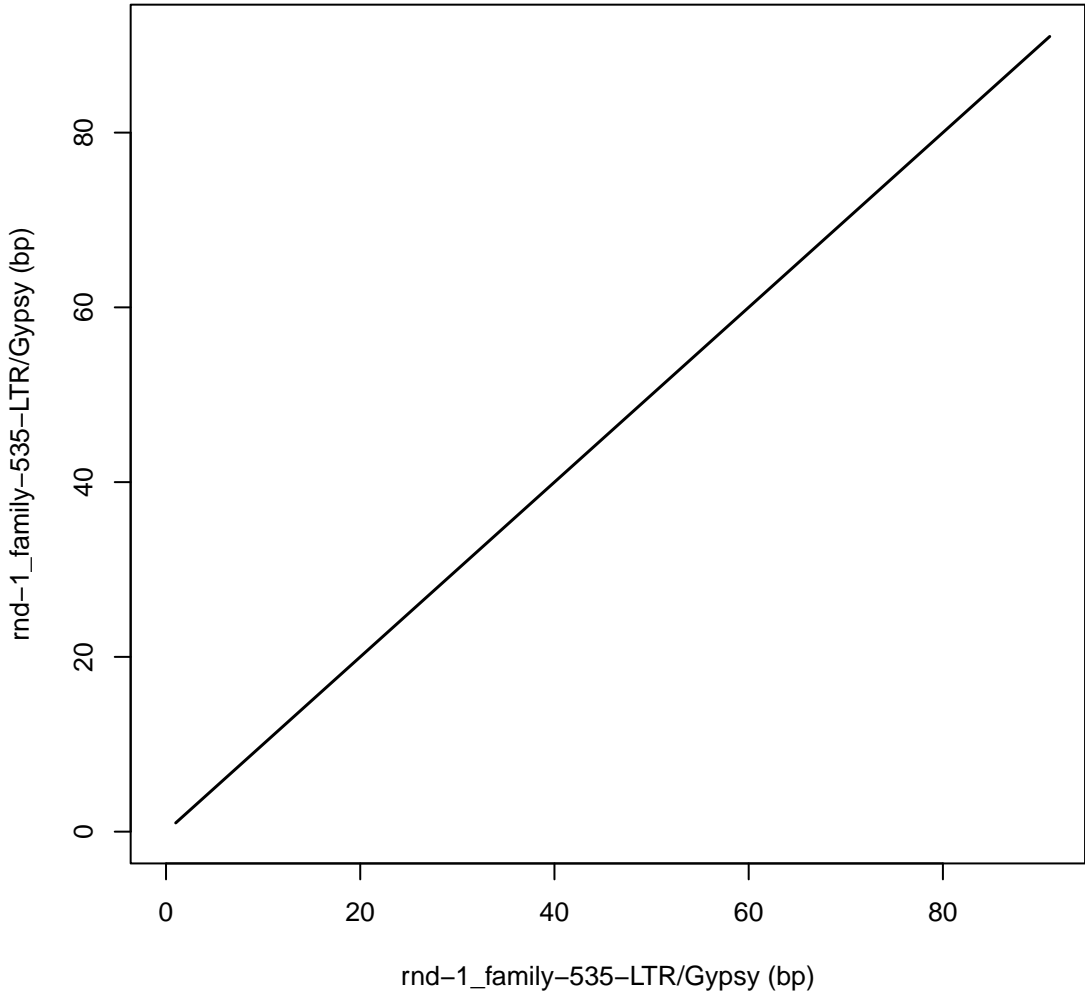
TE: rnd-1\_family-535-LTR/Gypsy  
consensus size: 91bp; fragments: 2; full length: 1 (>=81.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

