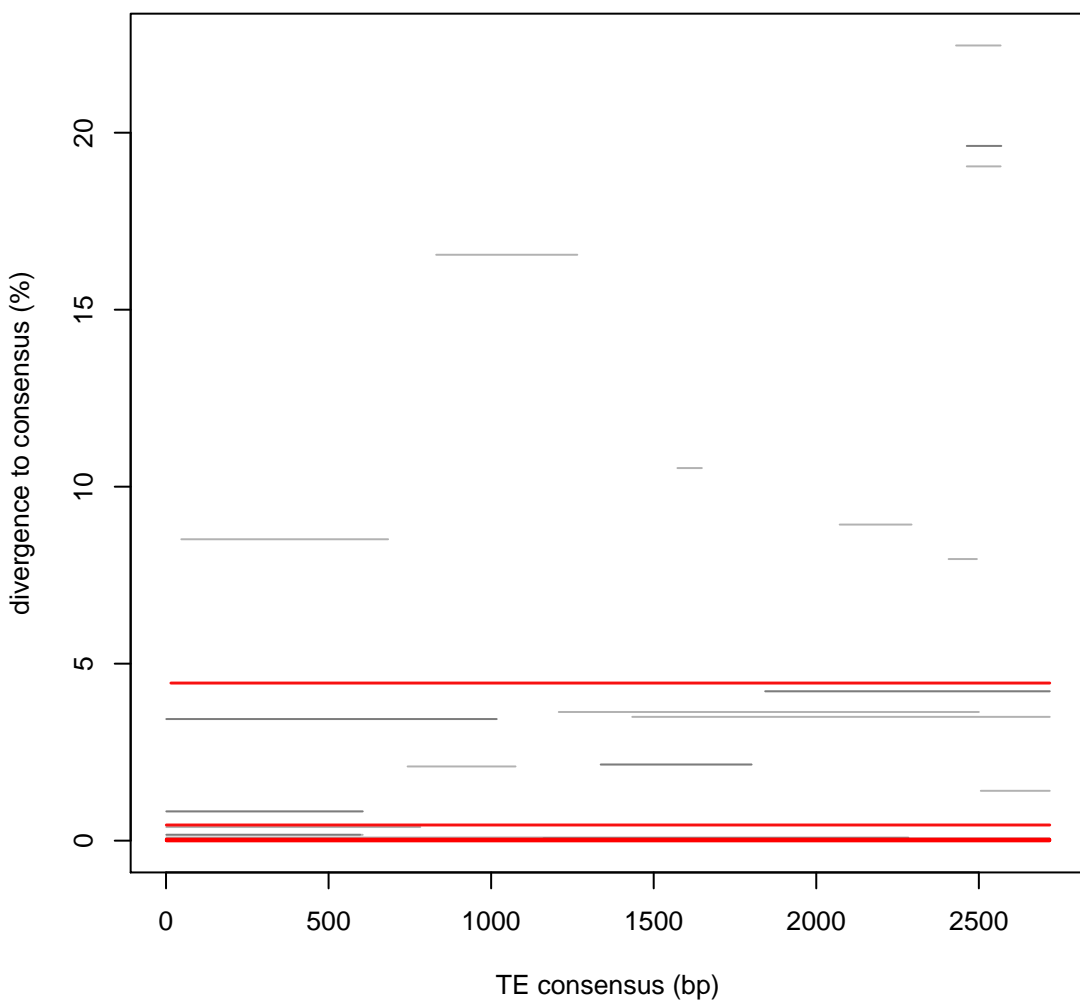
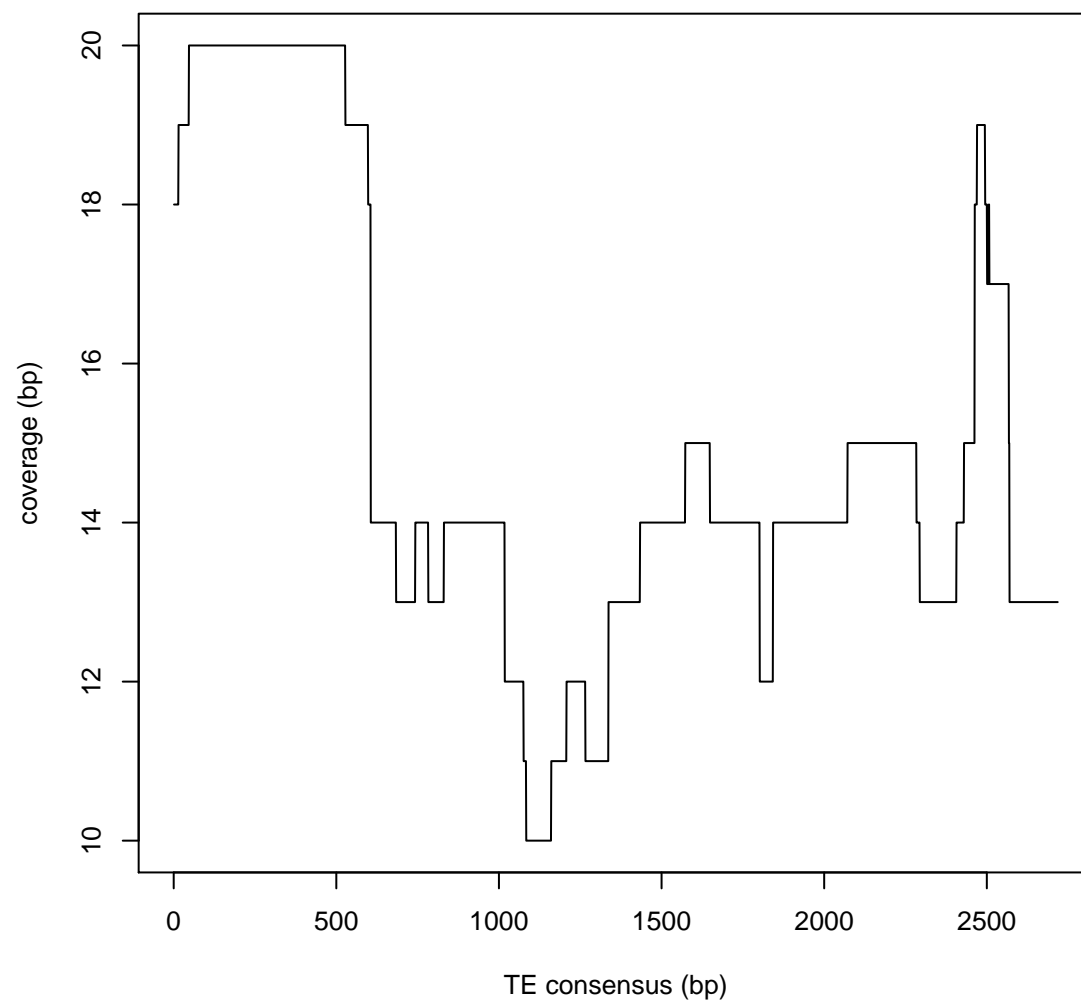


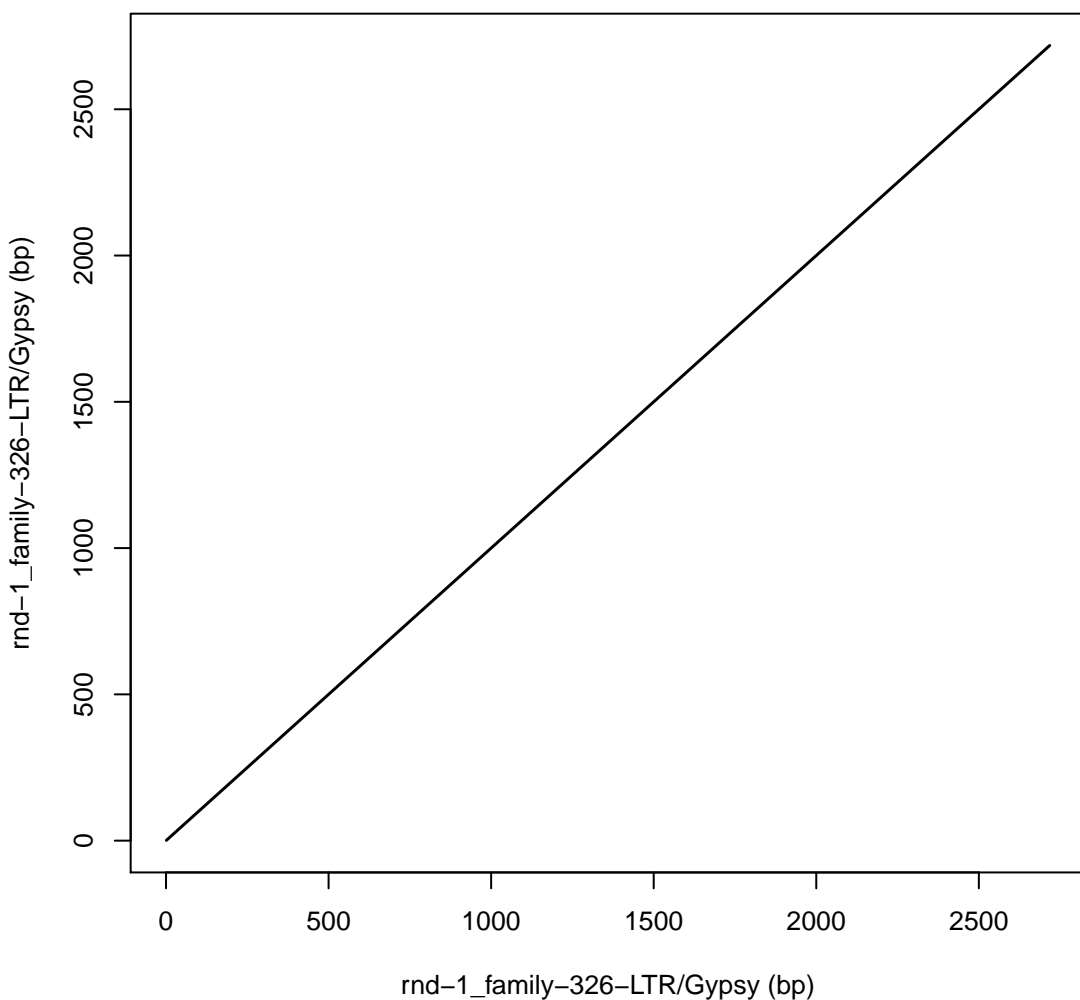
TE: rnd-1_family-326-LTR/Gypsy
consensus size: 2718bp; fragments: 38; full length: 8 (>=2446.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

