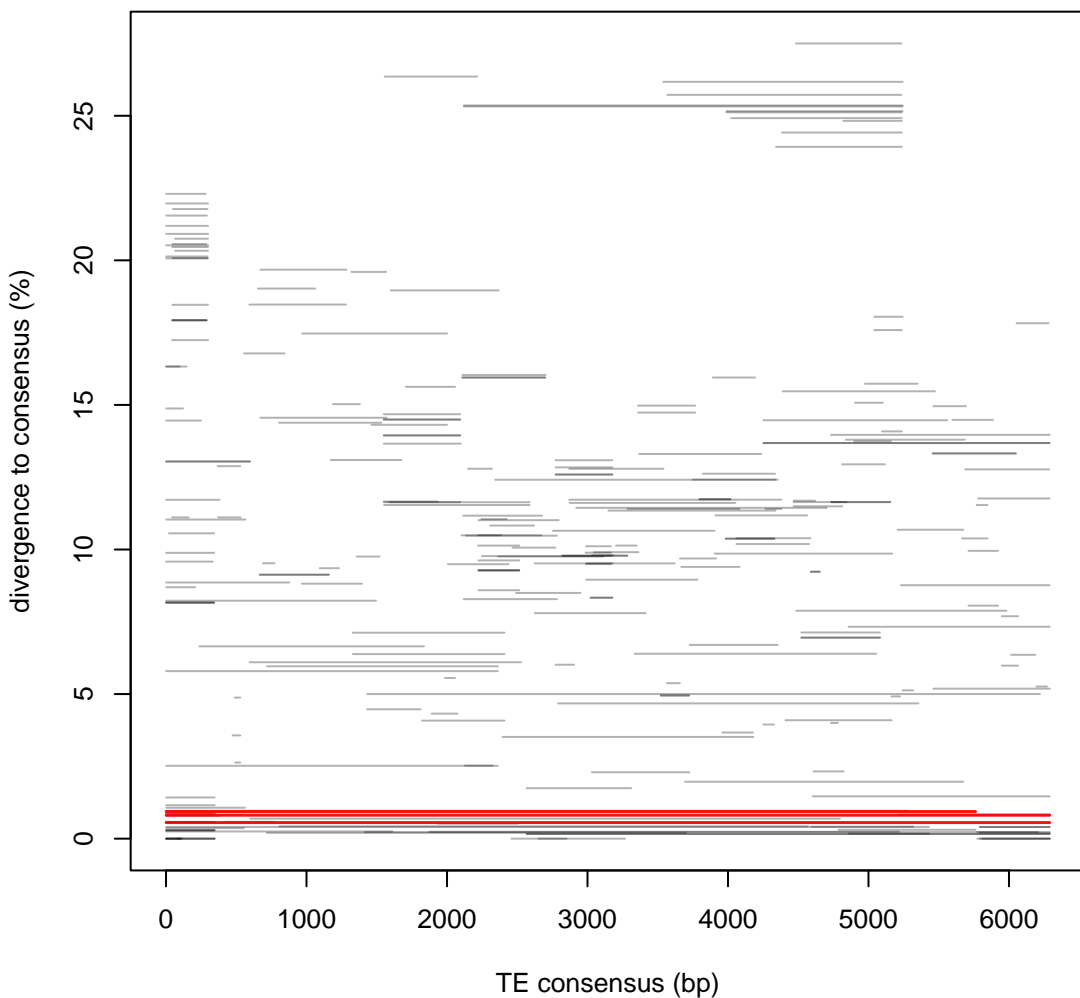
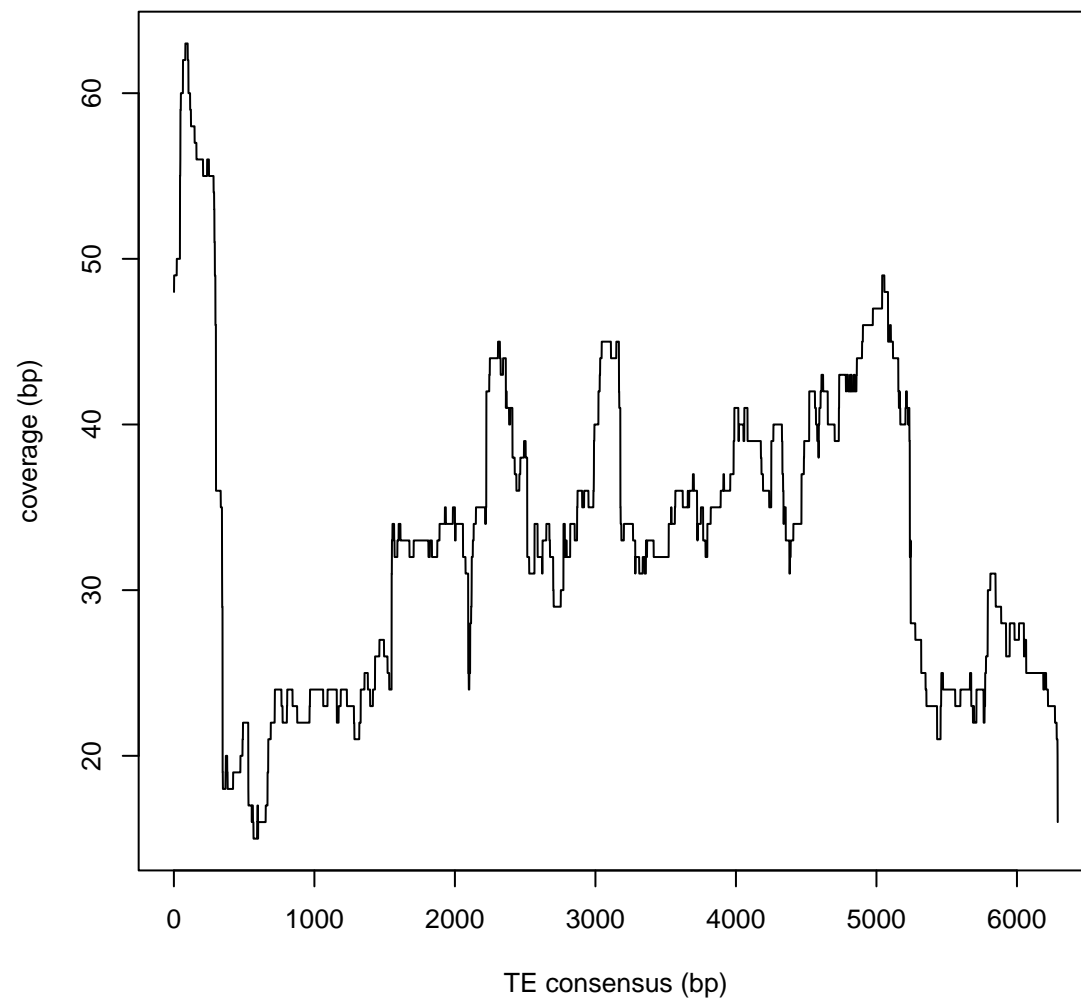


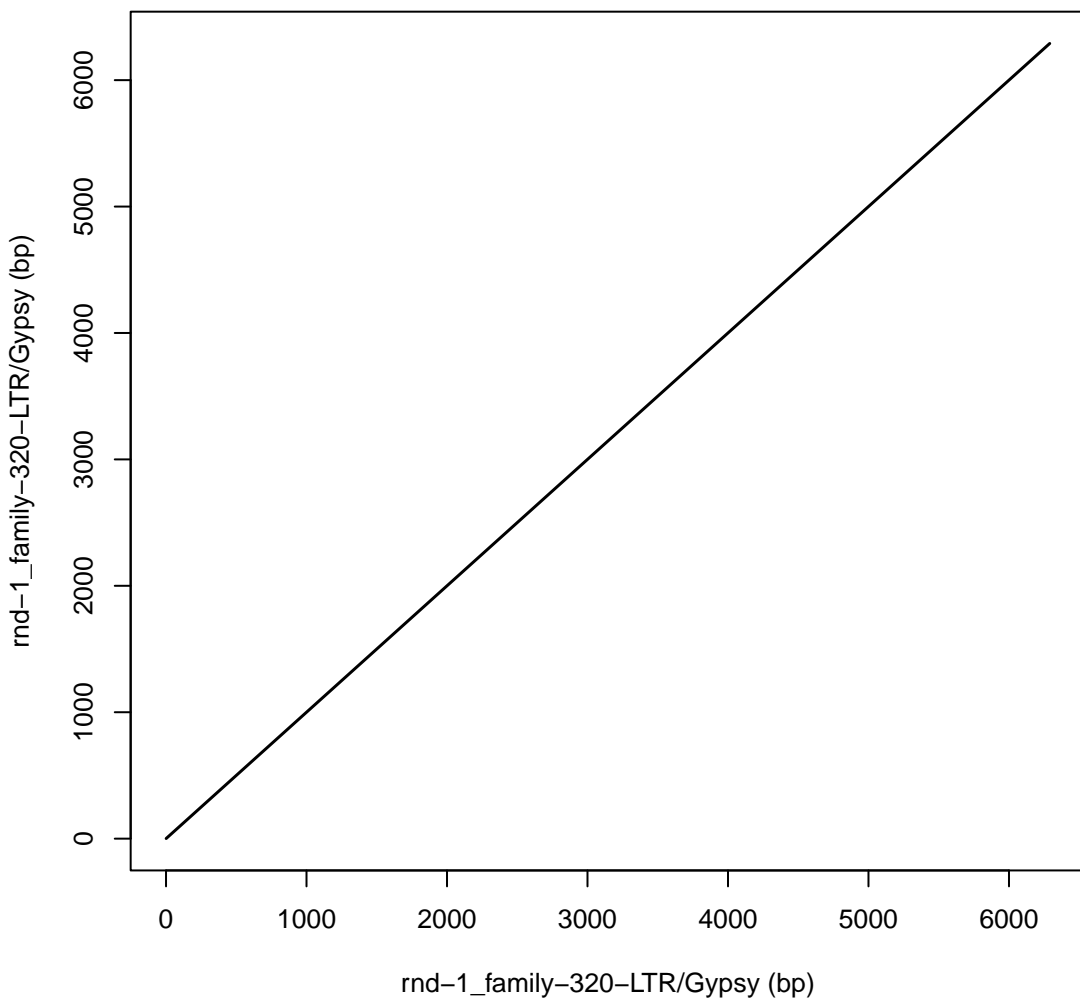
TE: rnd-1_family-320-LTR/Gypsy
consensus size: 6290bp; fragments: 270; full length: 3 (>=5661bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

