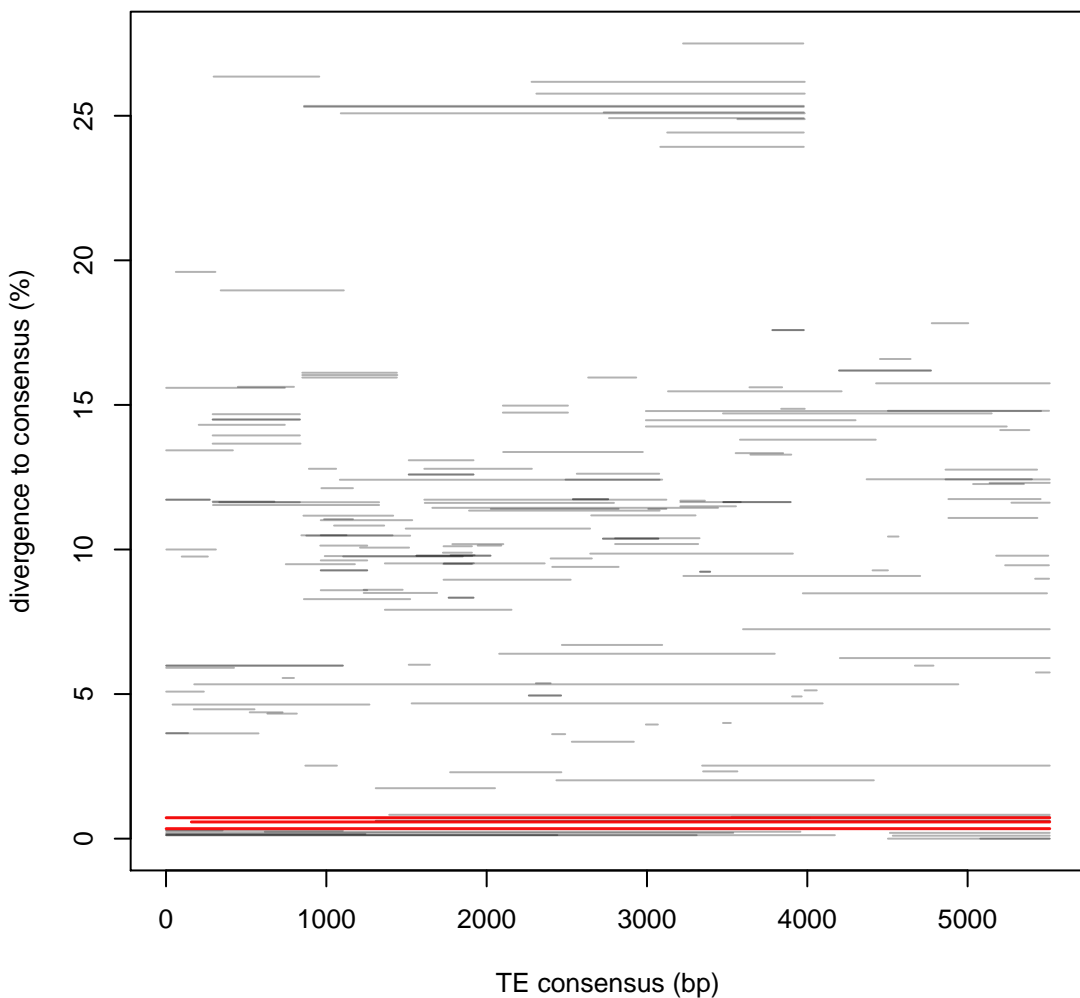
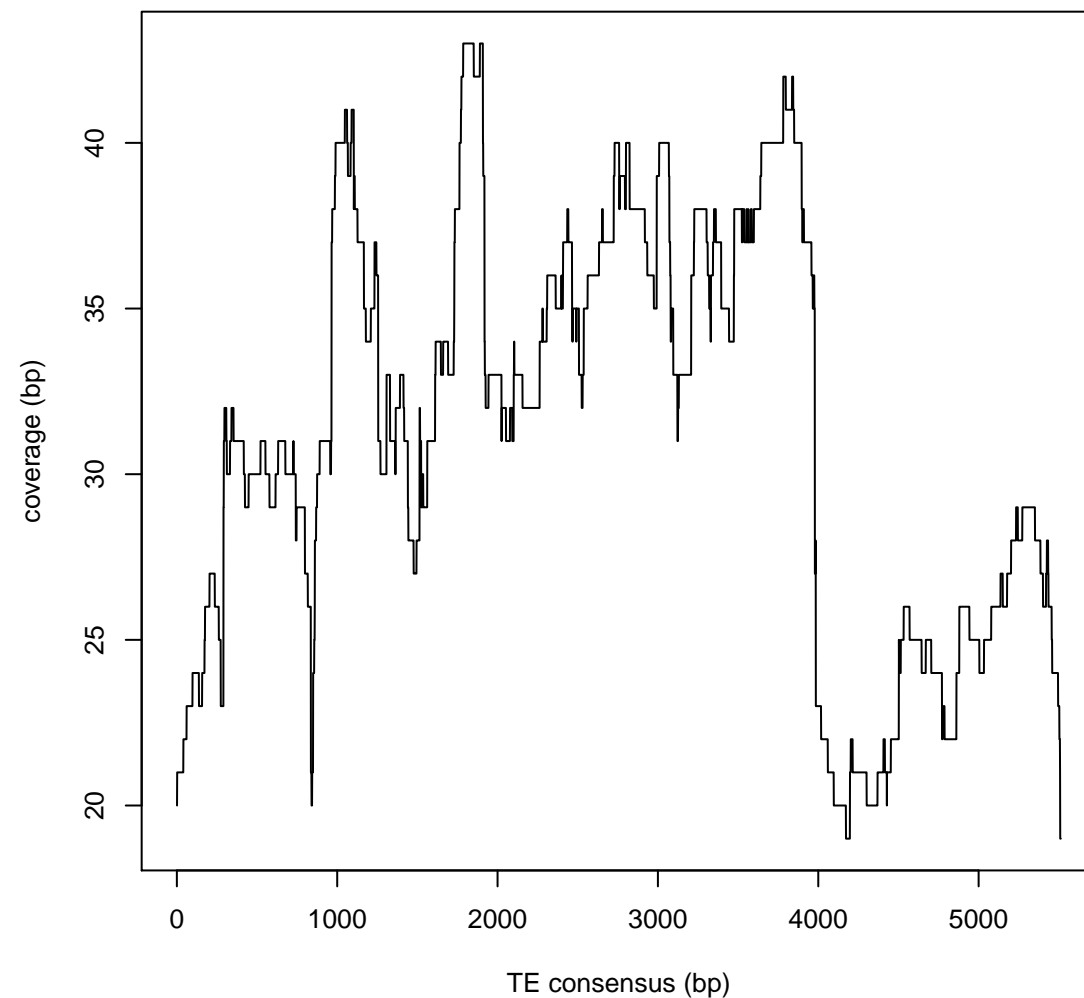


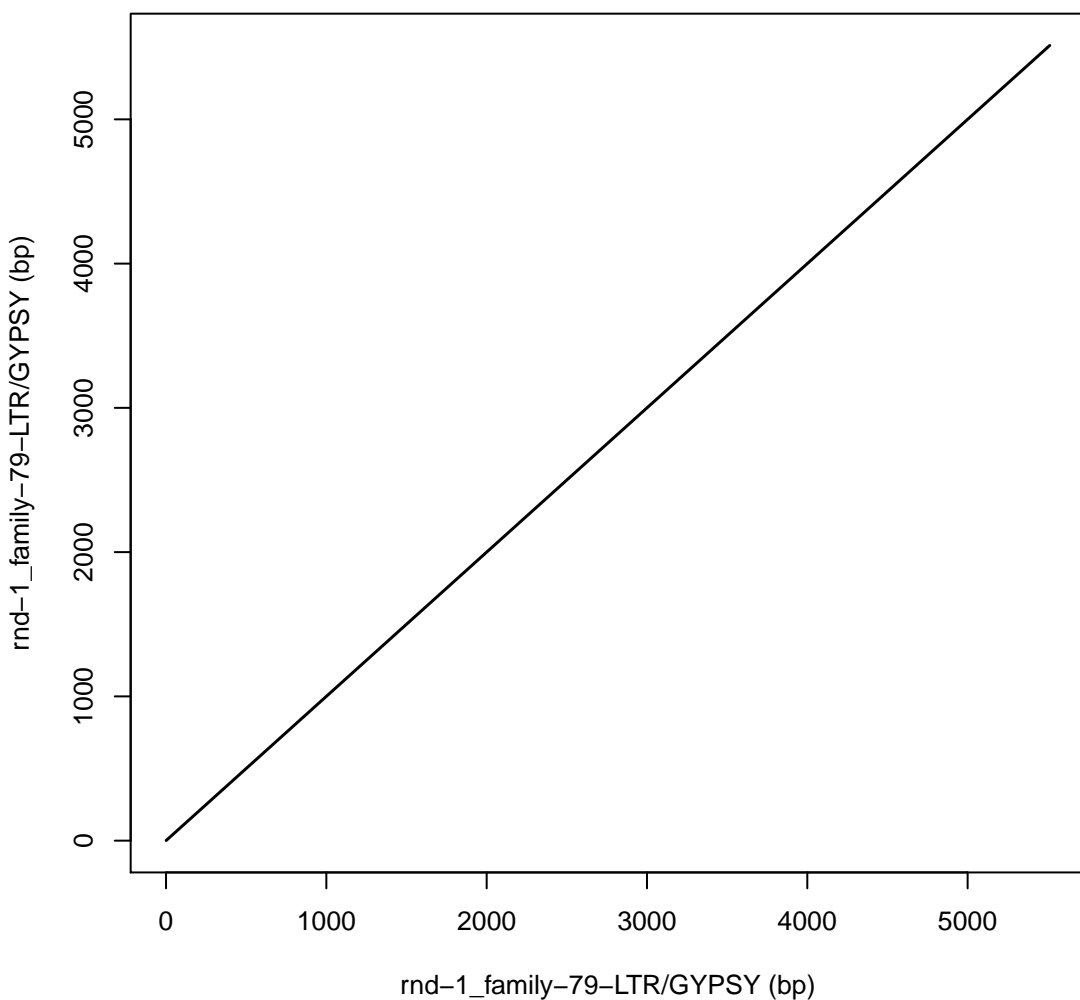
TE: rnd-1_family-79-LTR/GYPSY
consensus size: 5512bp; fragments: 193; full length: 3 (>=4960.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

