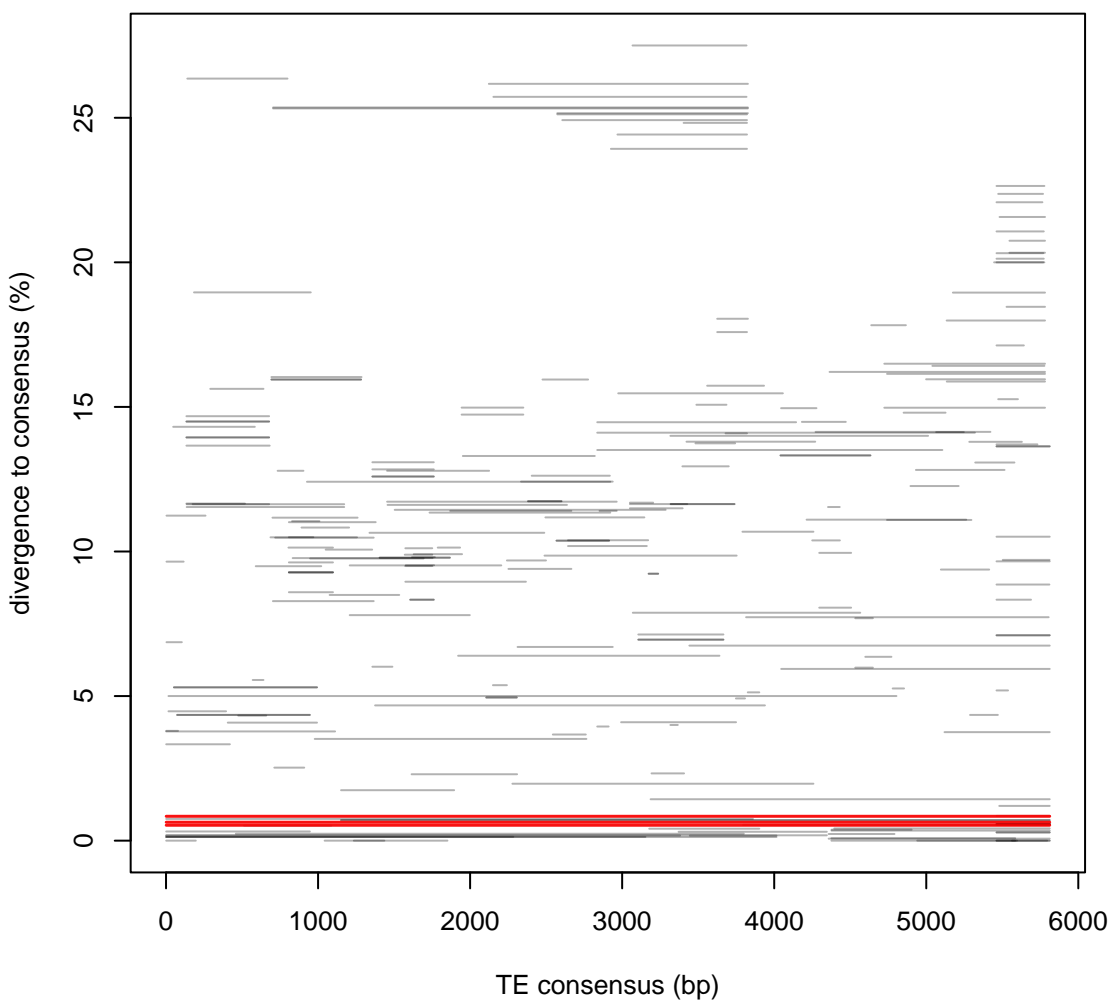
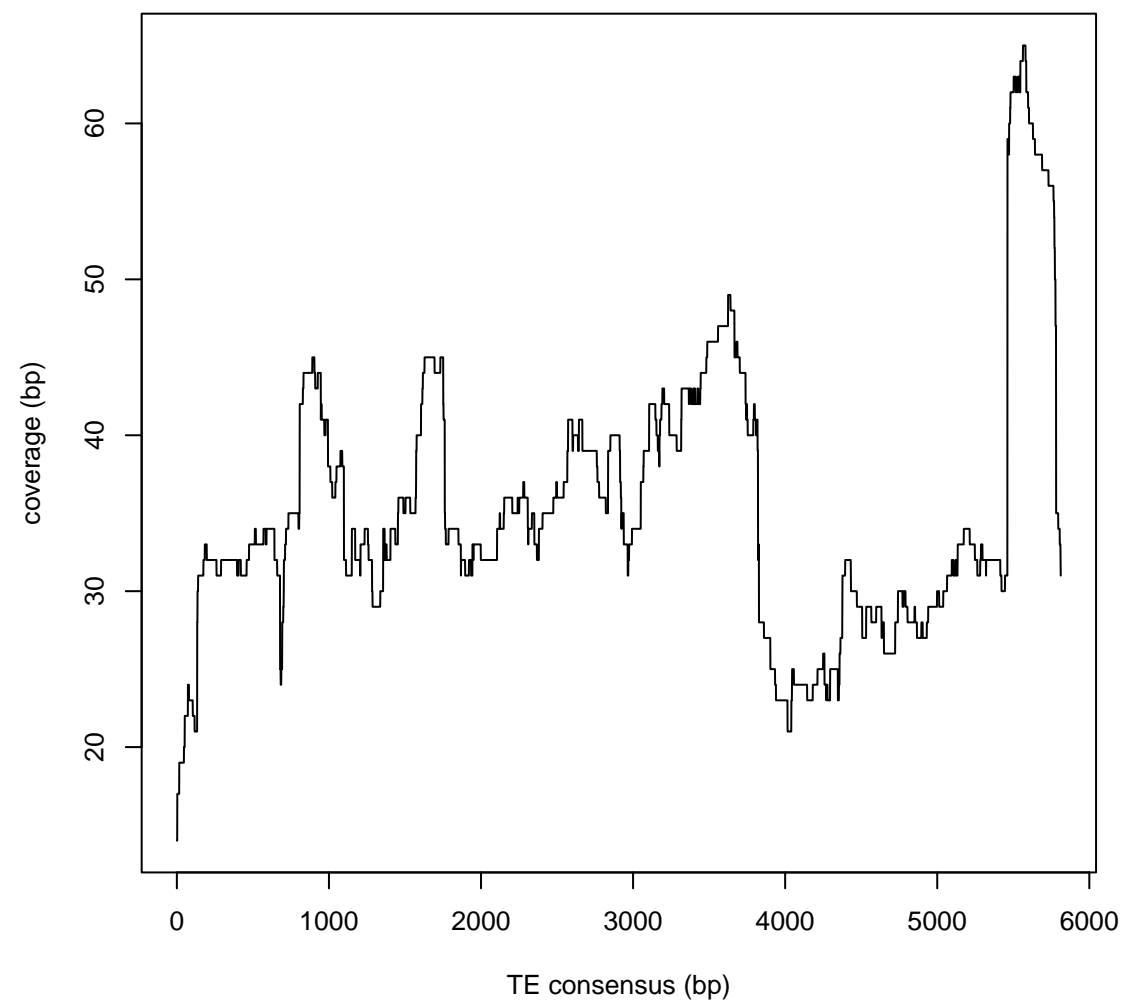


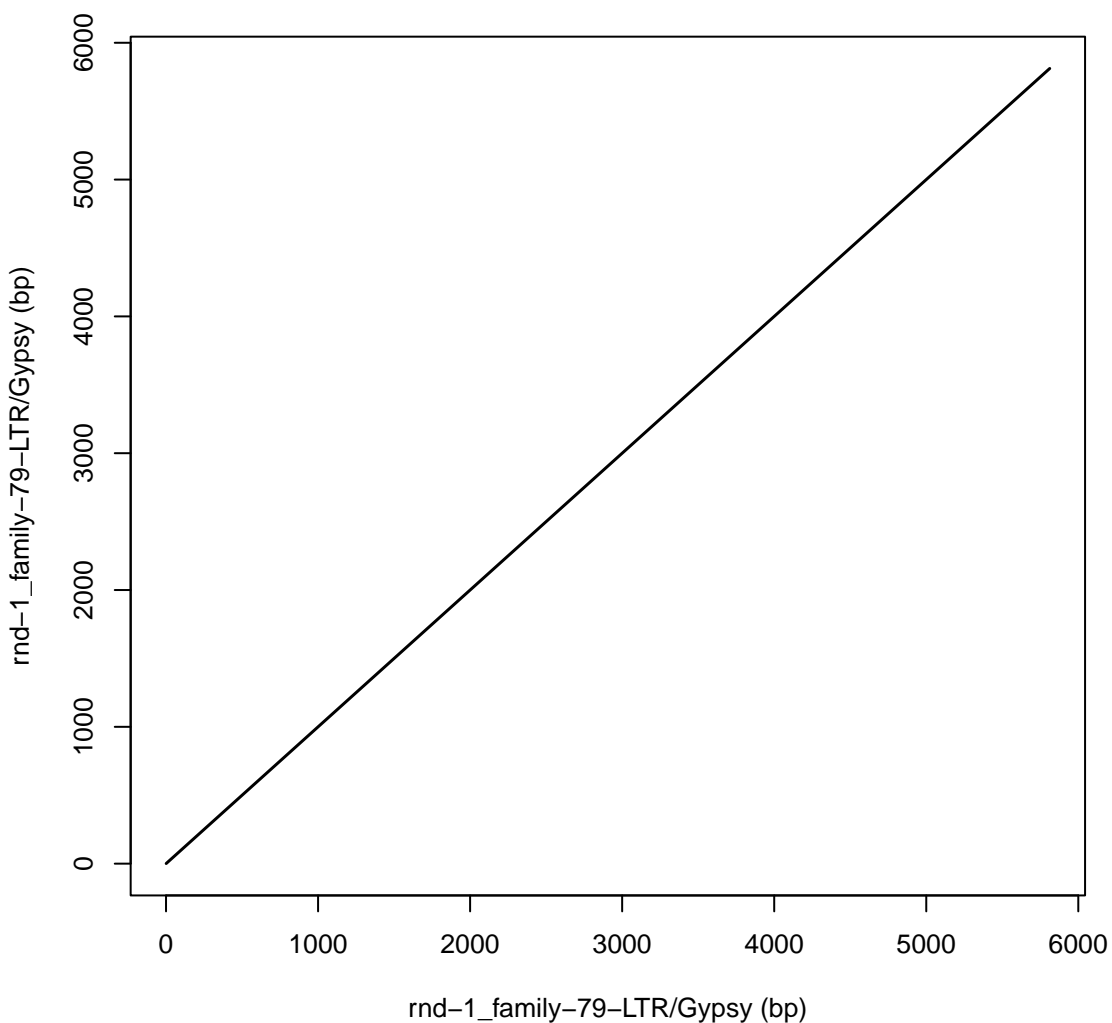
TE: rnd-1_family-79-LTR/Gypsy
consensus size: 5812bp; fragments: 255; full length: 3 (>=5230.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

