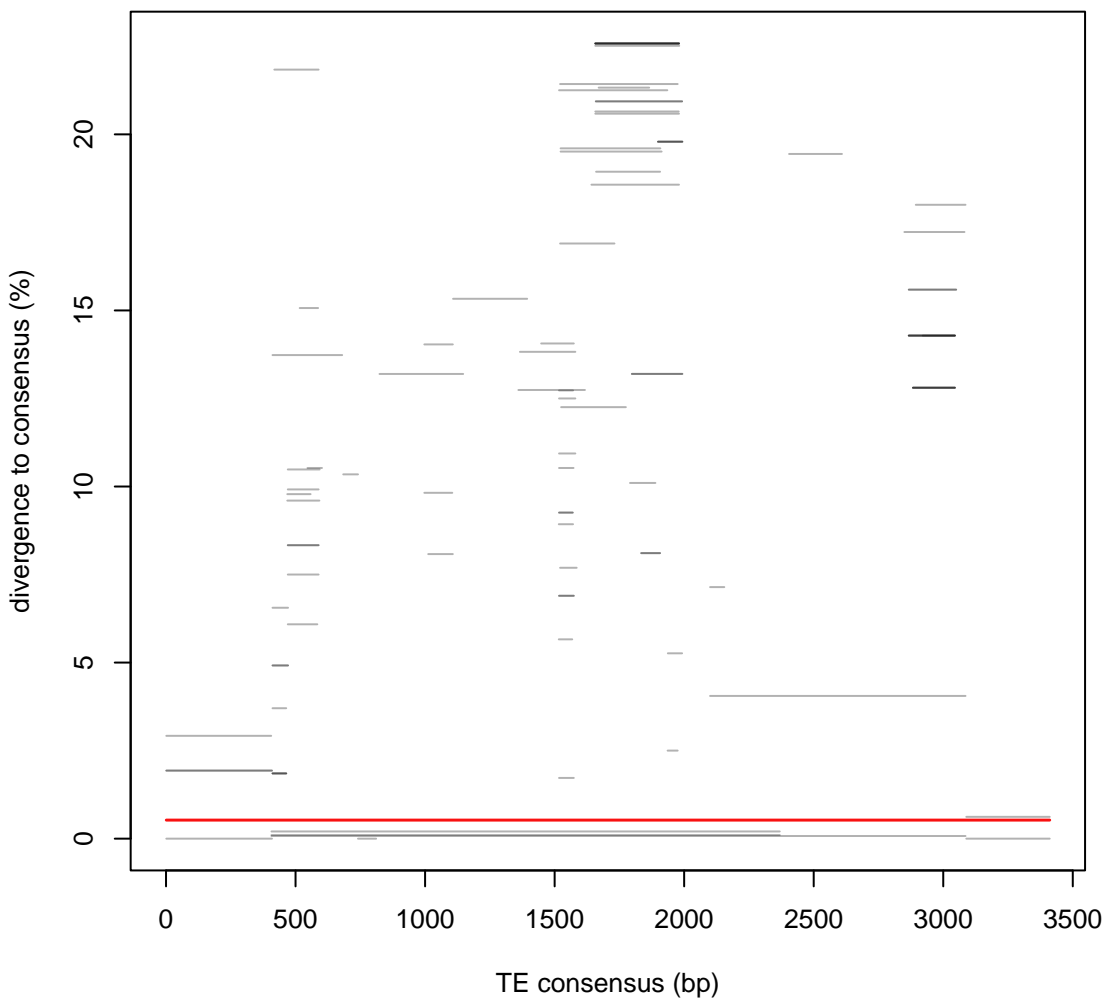
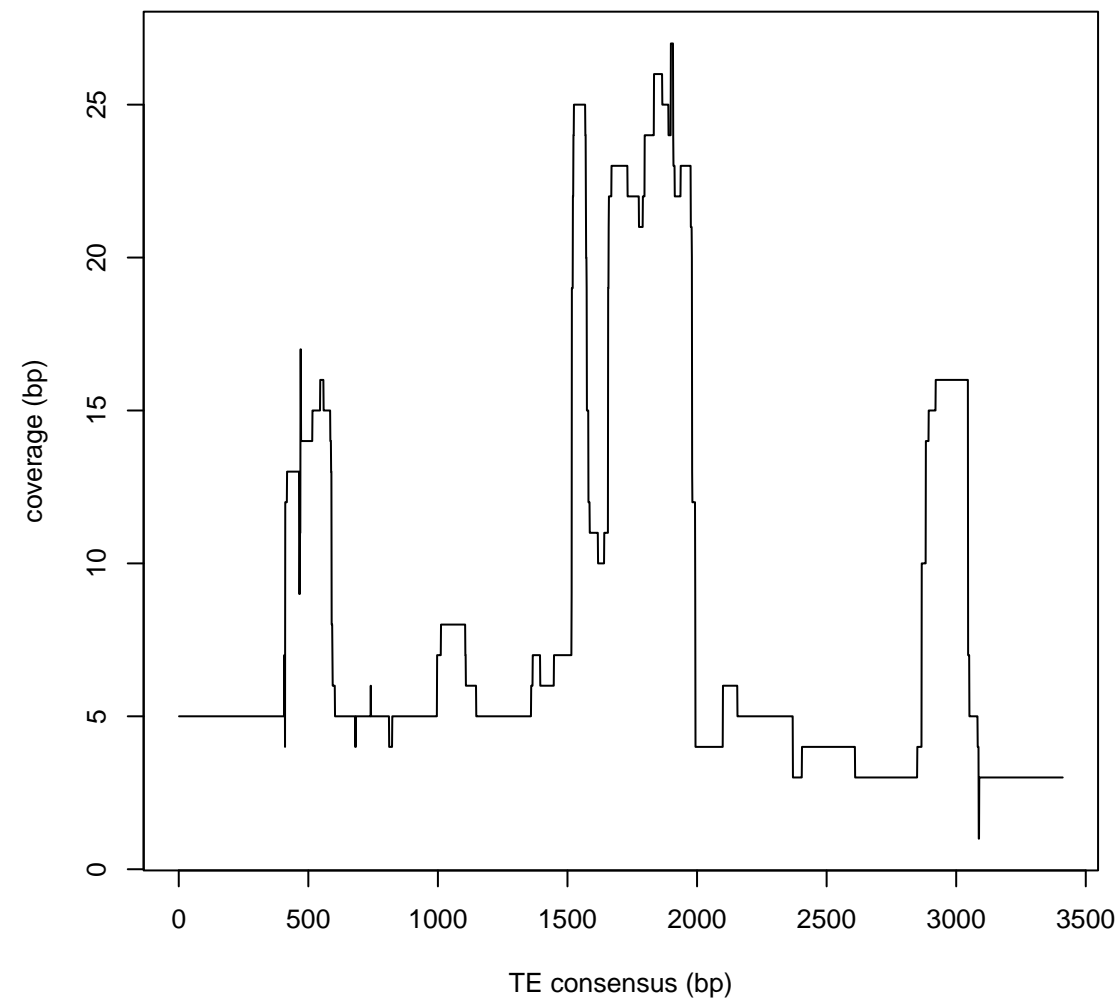


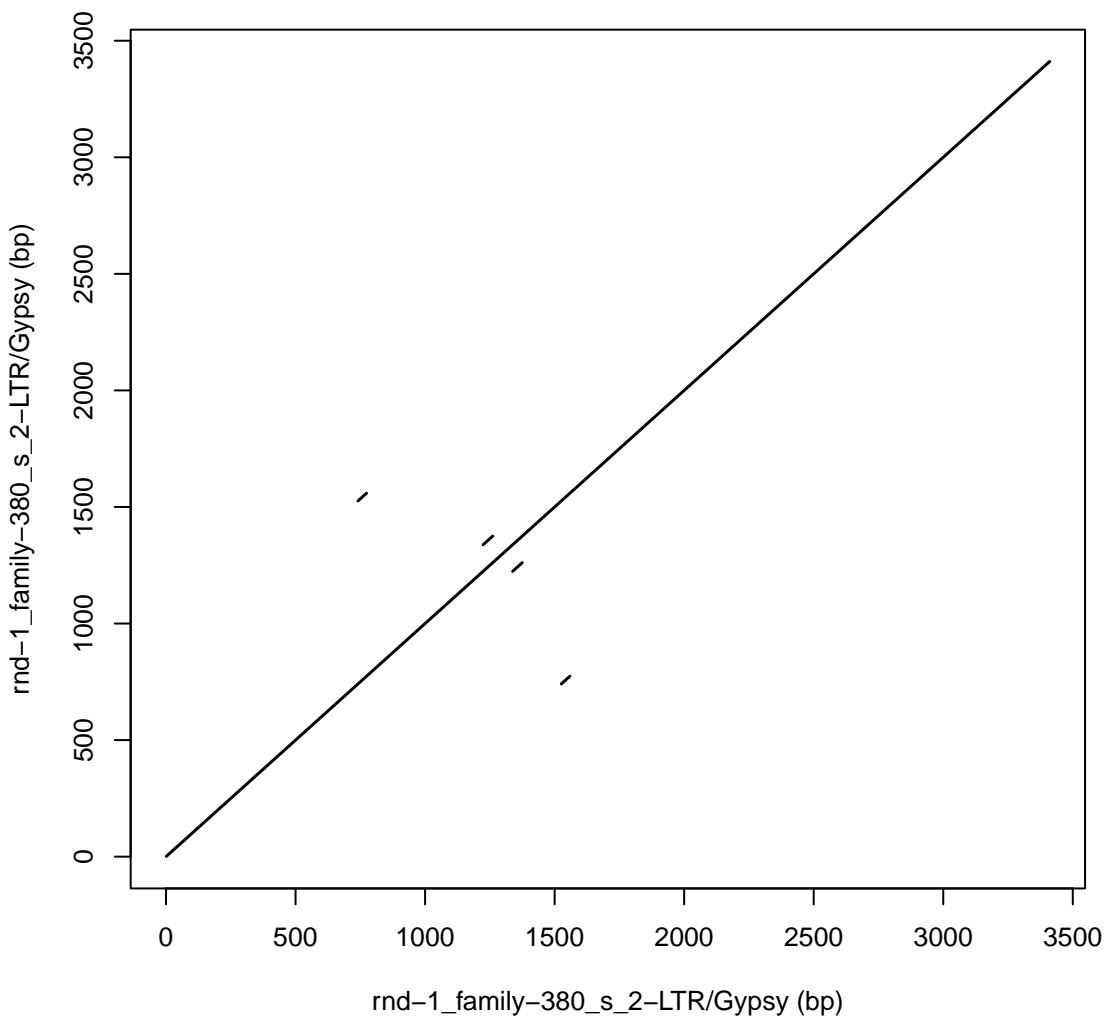
TE: rnd-1_family-380_s_2-LTR/Gypsy
consensus size: 3411bp; fragments: 96; full length: 1 (>=3069.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

