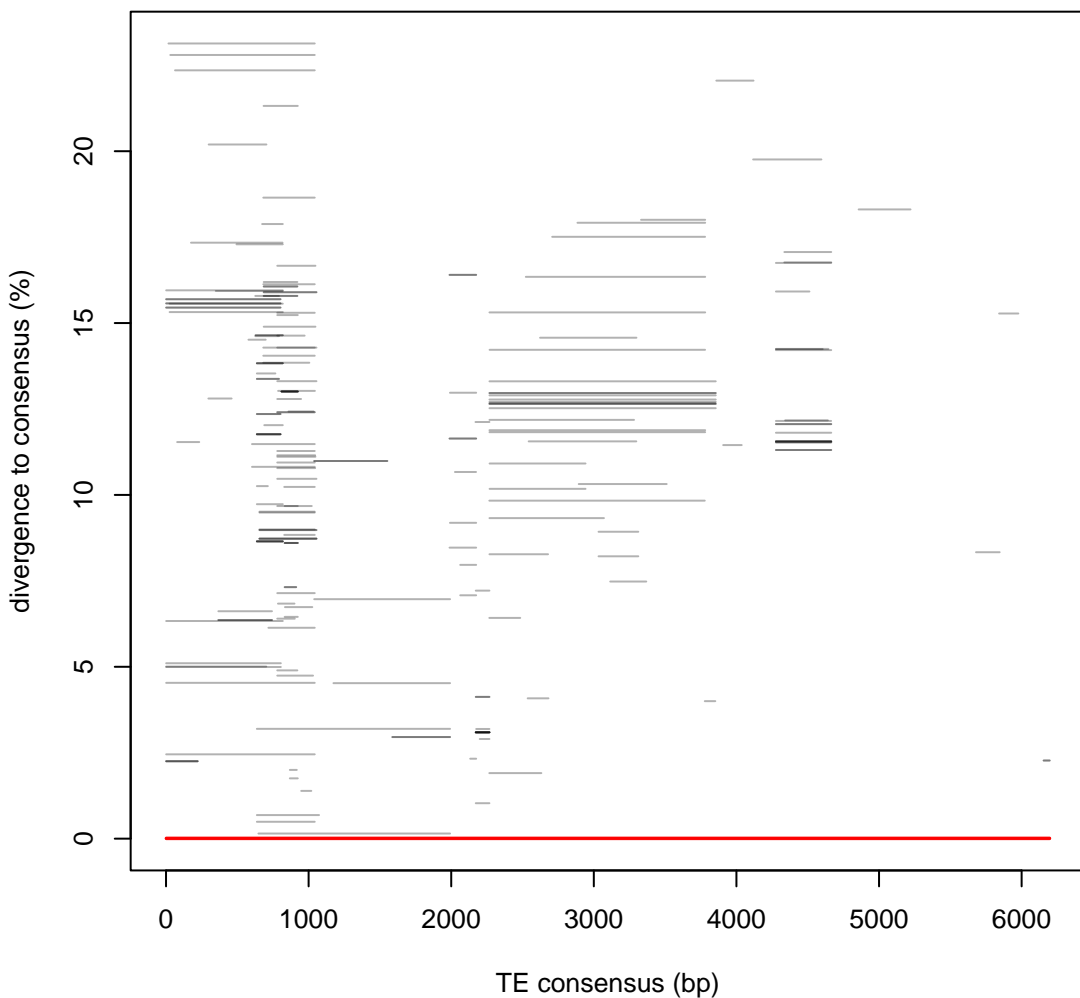
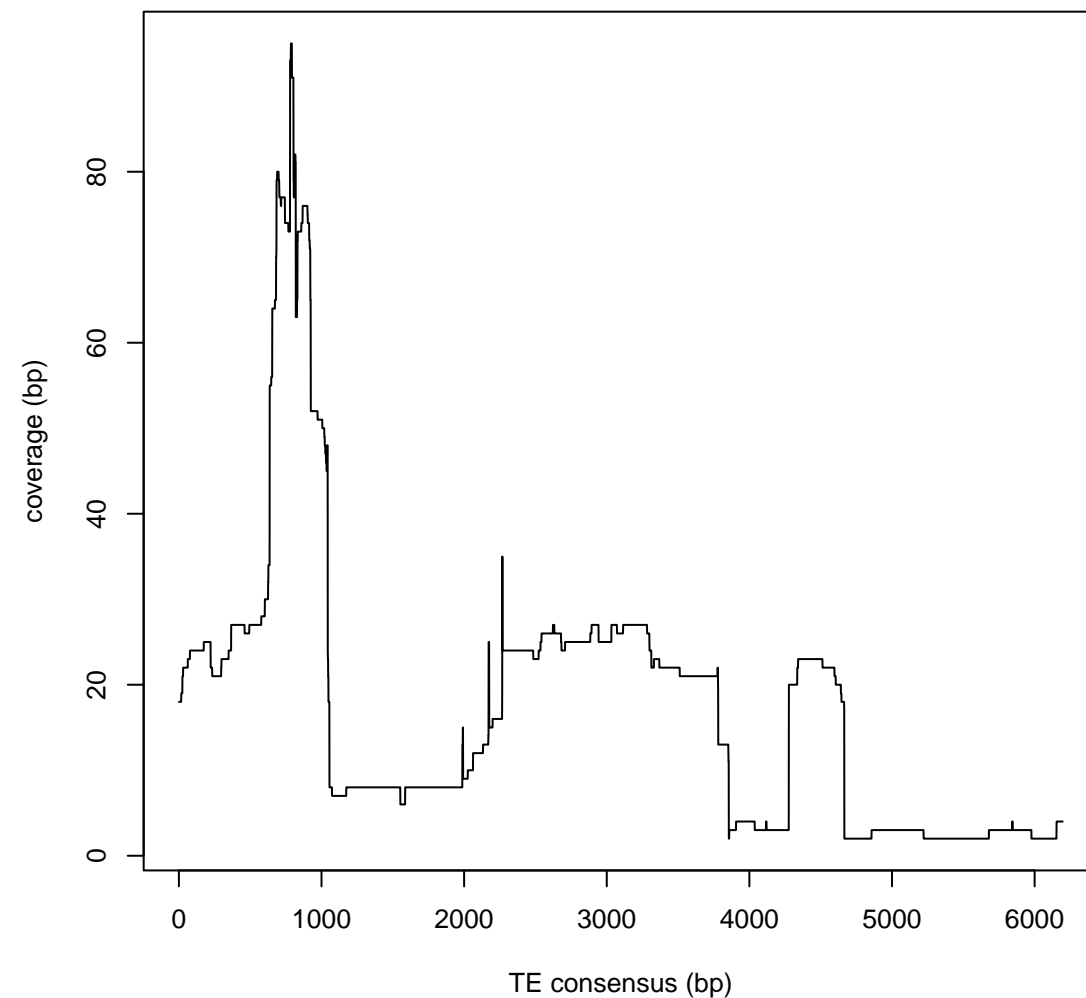


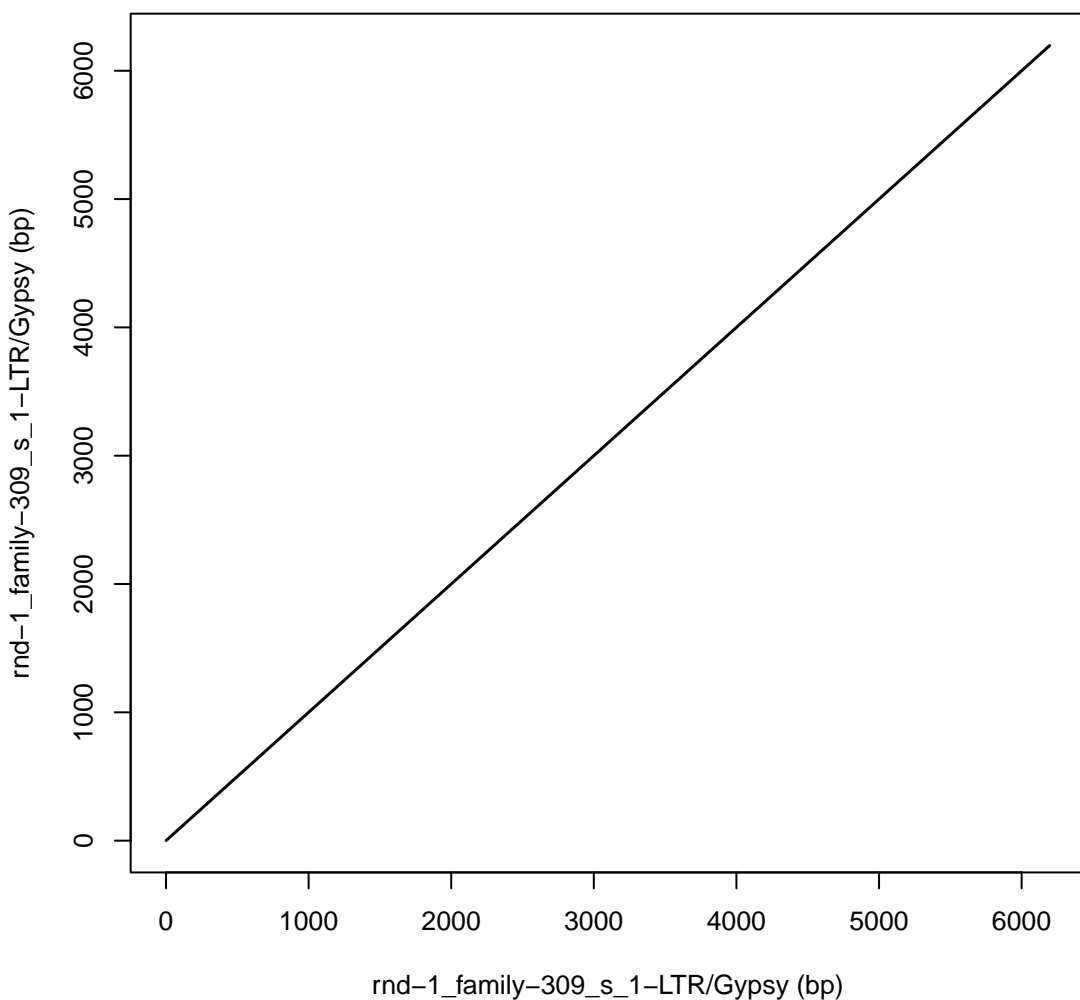
TE: rnd-1_family-309_s_1-LTR/Gypsy
consensus size: 6197bp; fragments: 220; full length: 2 (>=5577.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

