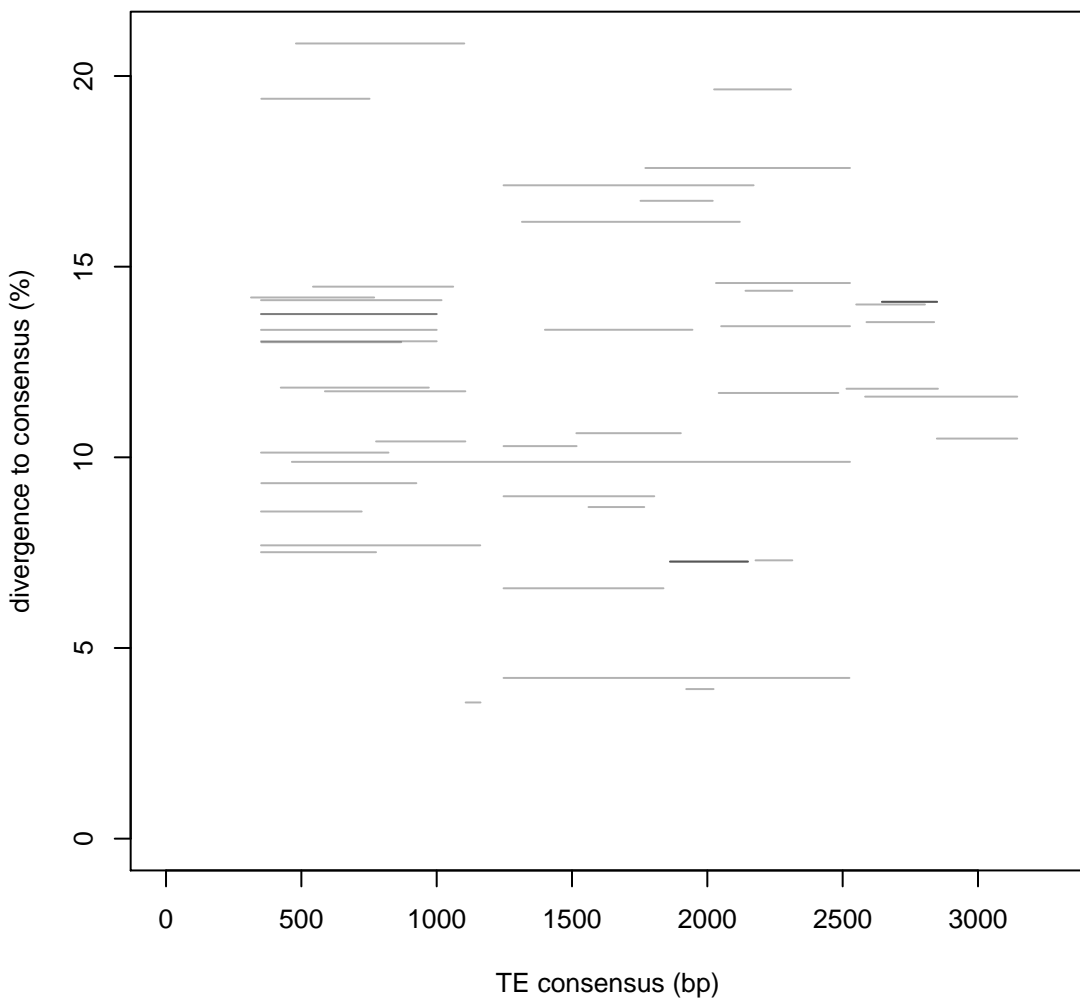
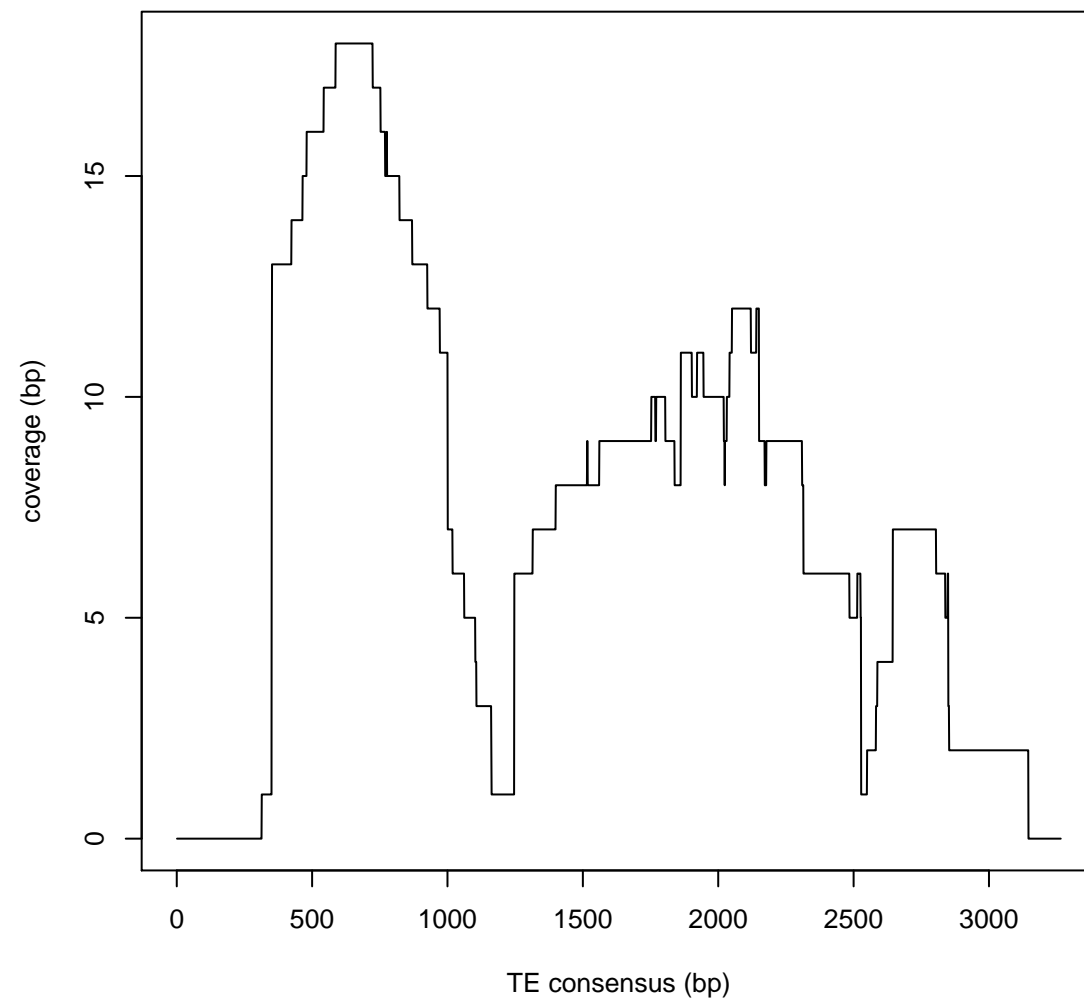


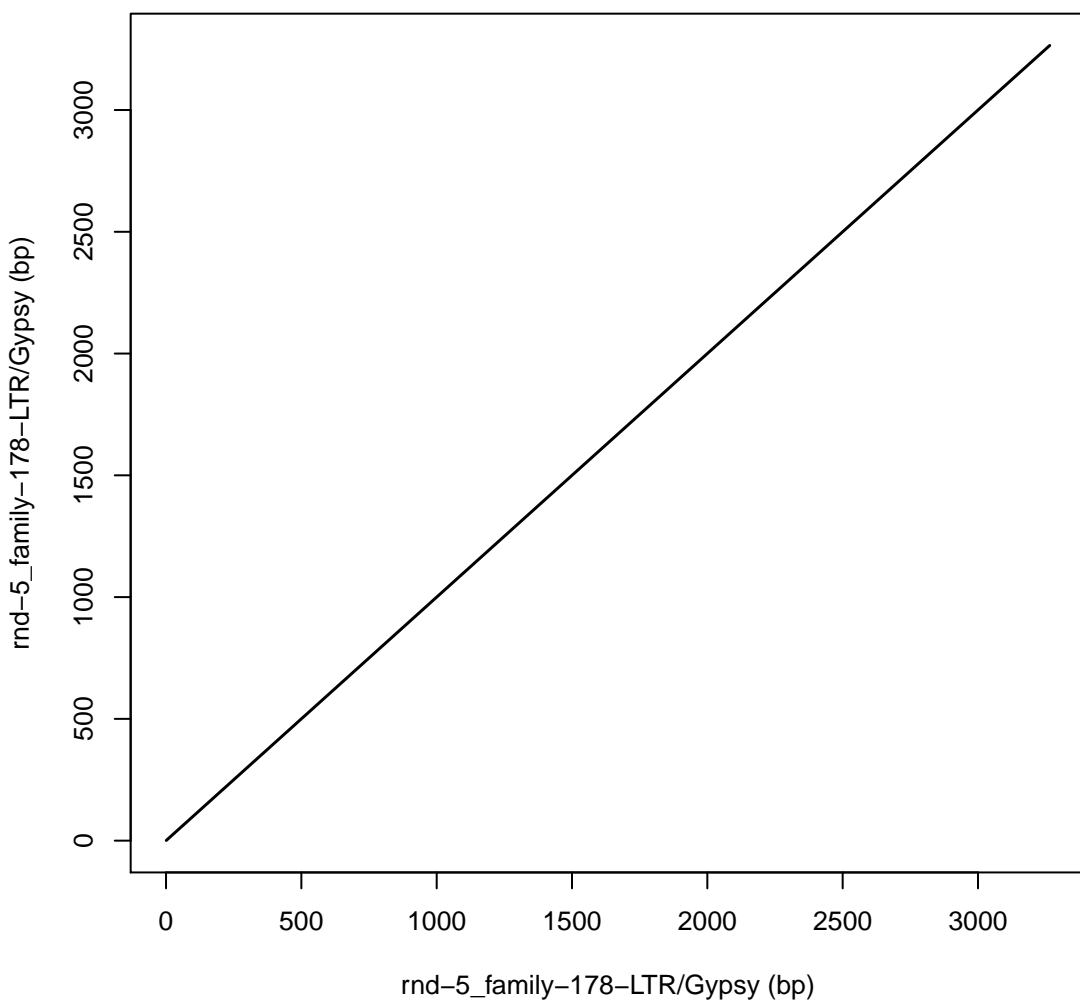
TE: rnd-5_family-178-LTR/Gypsy
consensus size: 3265bp; fragments: 49; full length: 0 (>=2938.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

