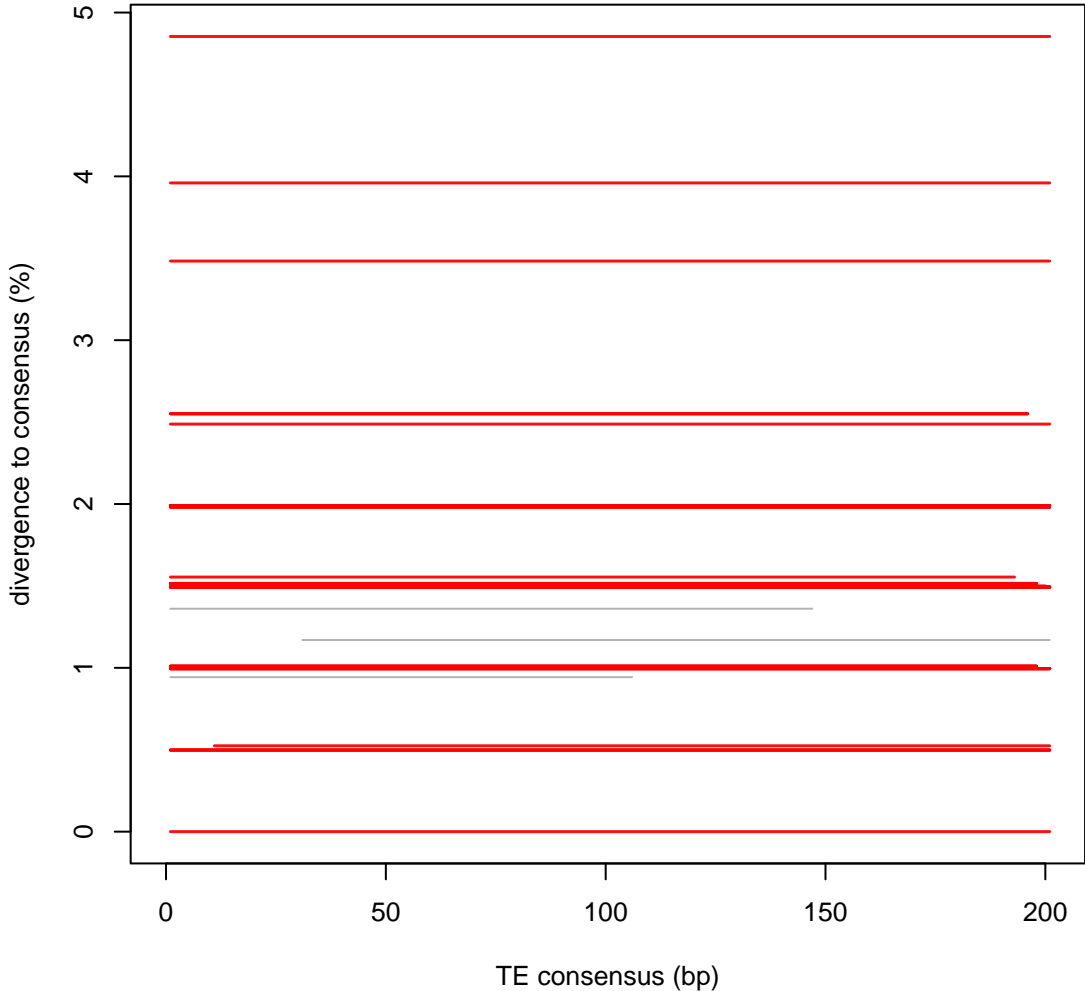
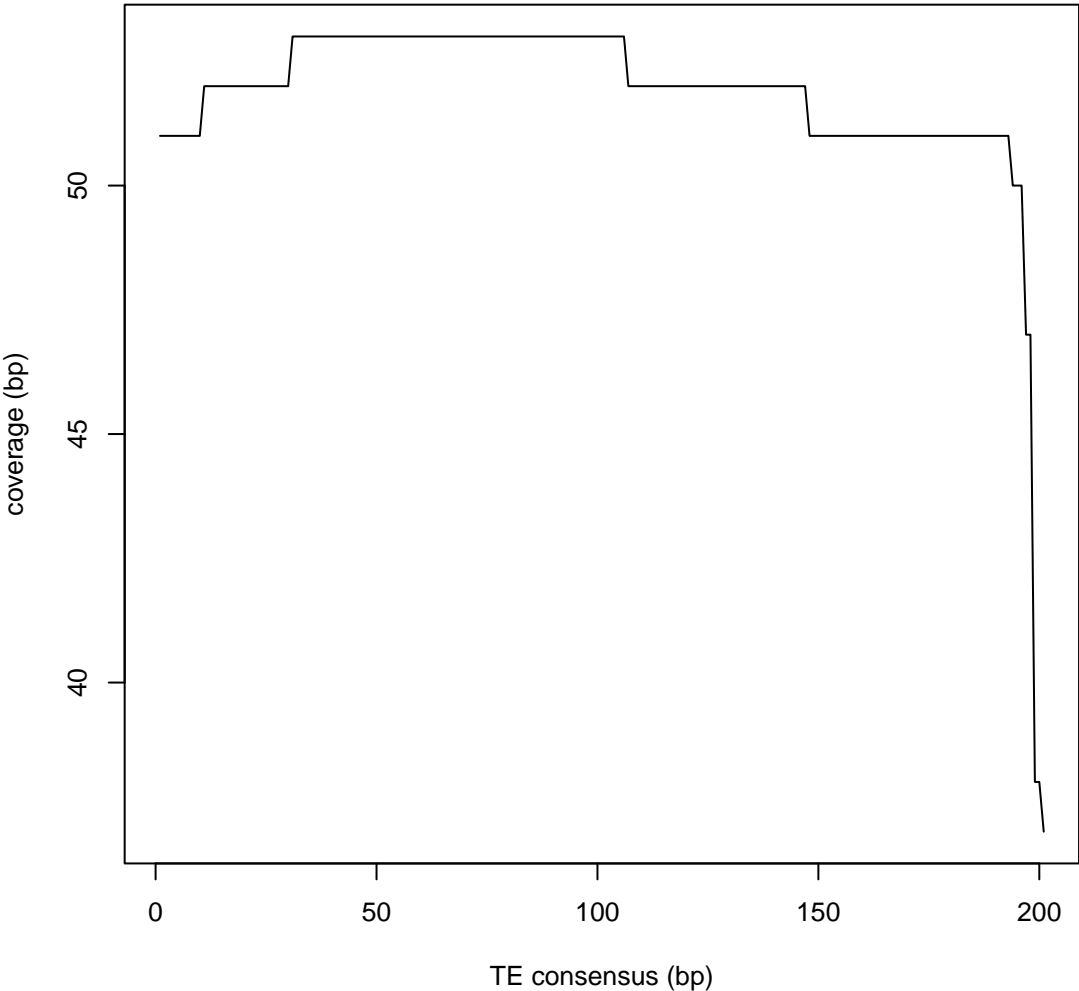


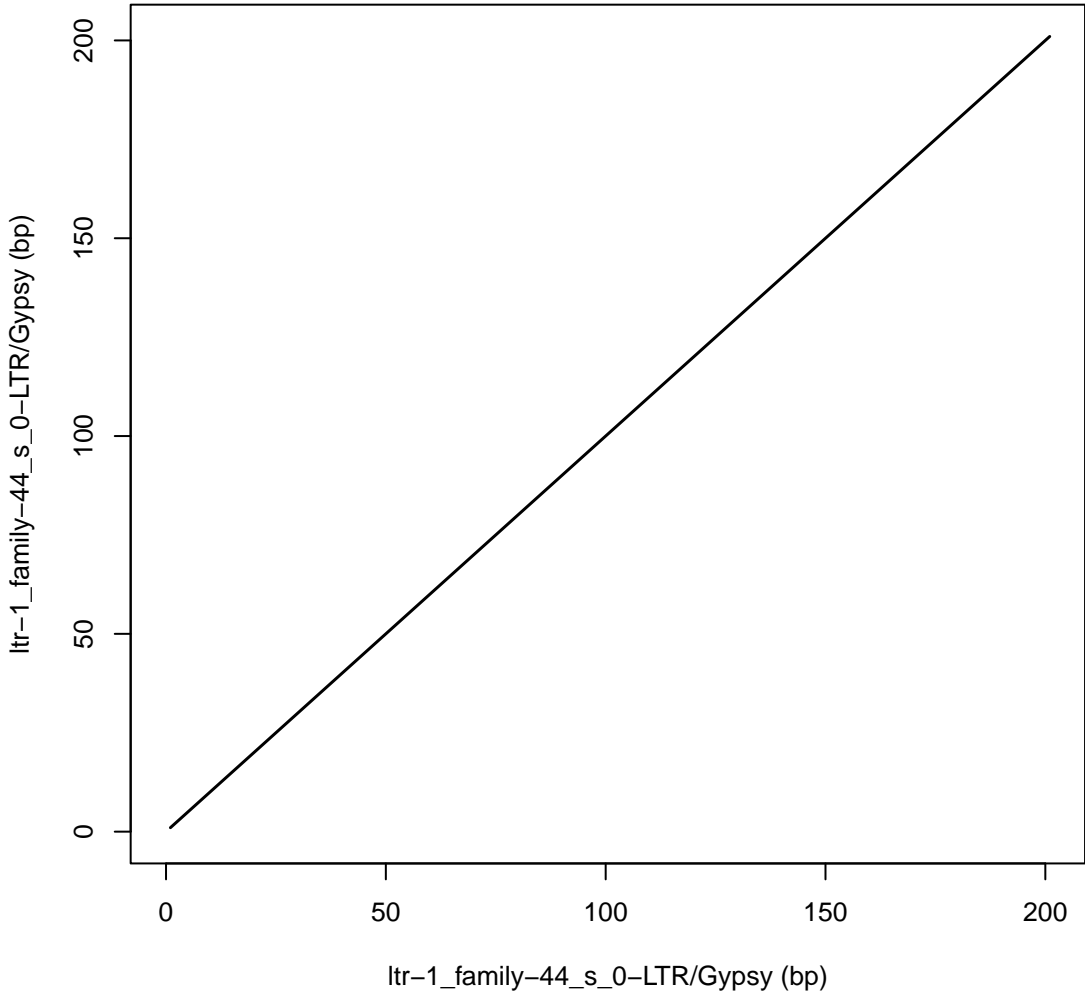
TE: ltr-1_family-44_s_0-LTR/Gypsy
consensus size: 201bp; fragments: 53; full length: 50 (>=180.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

