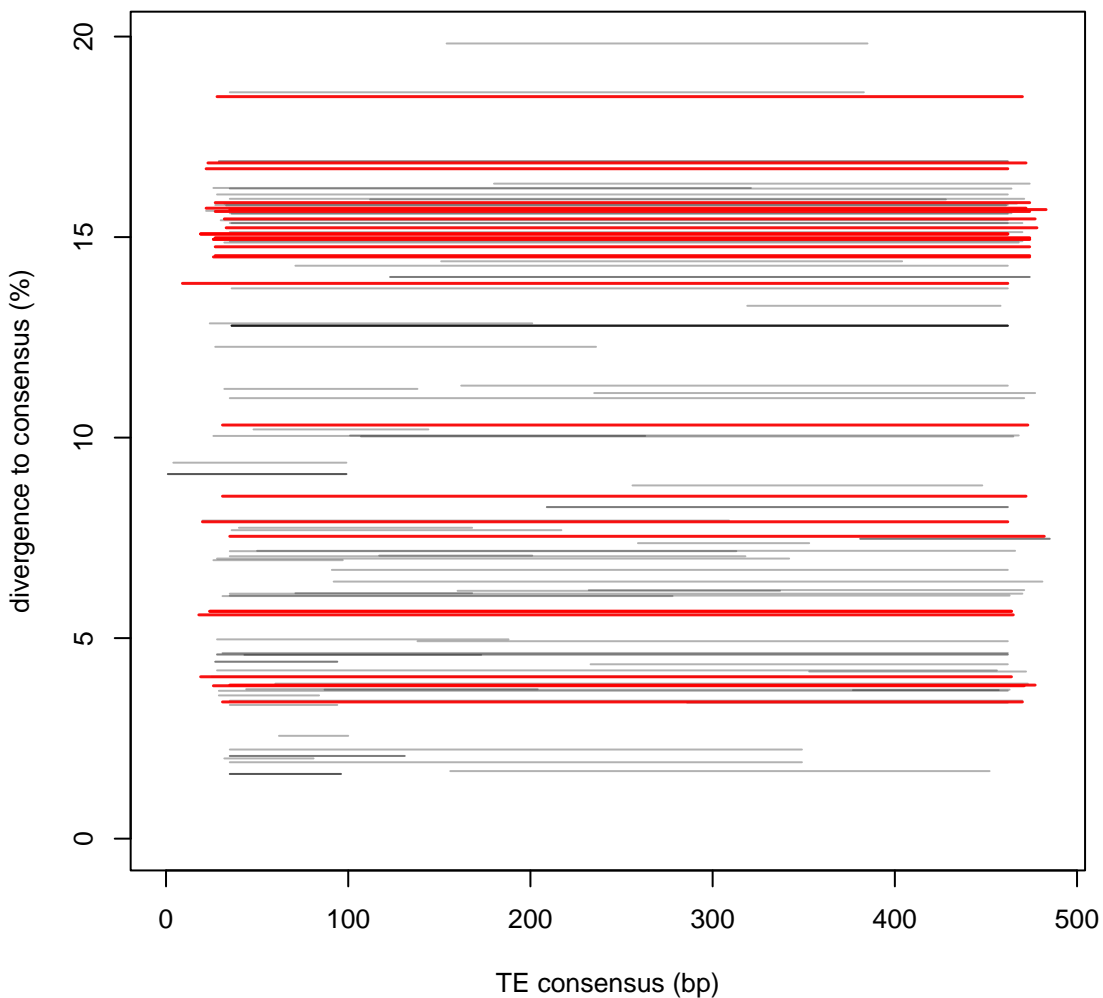
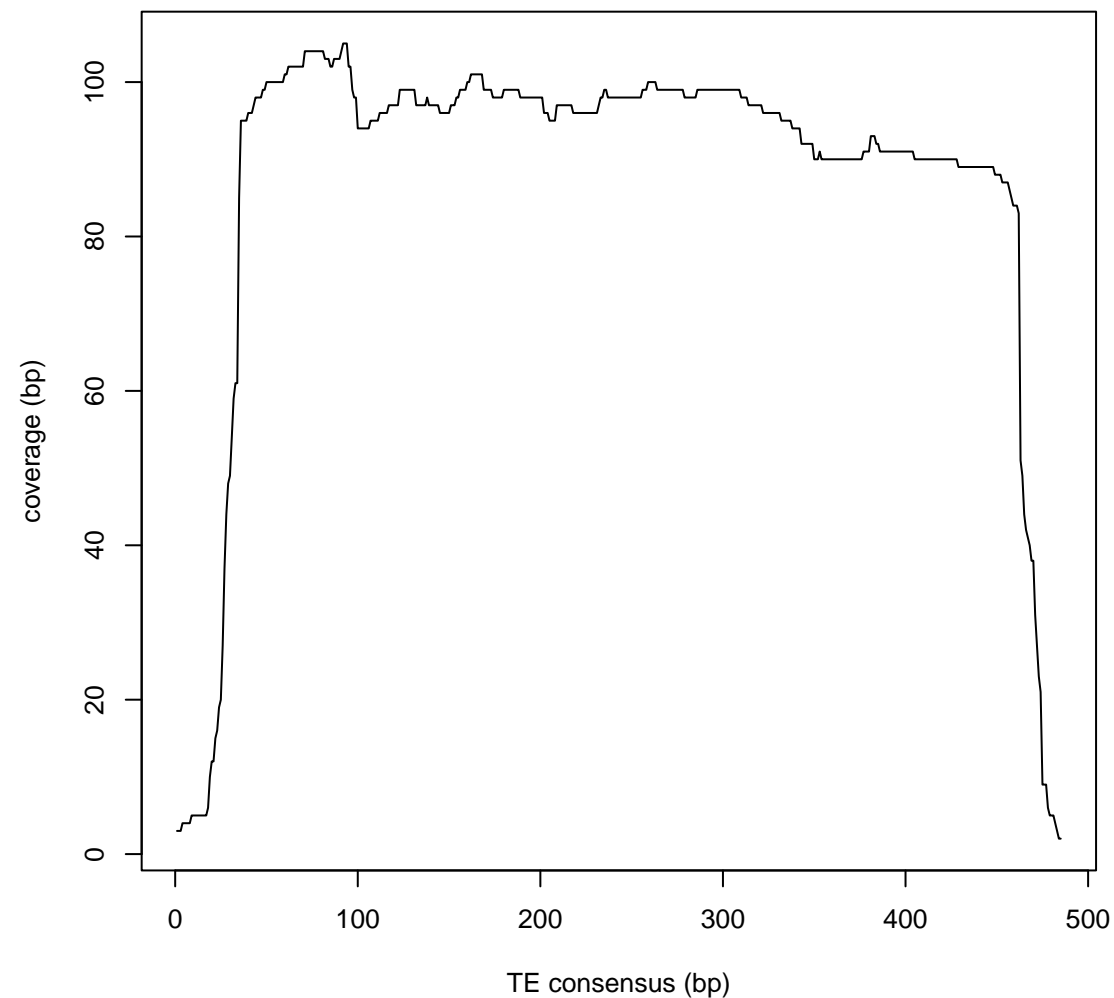


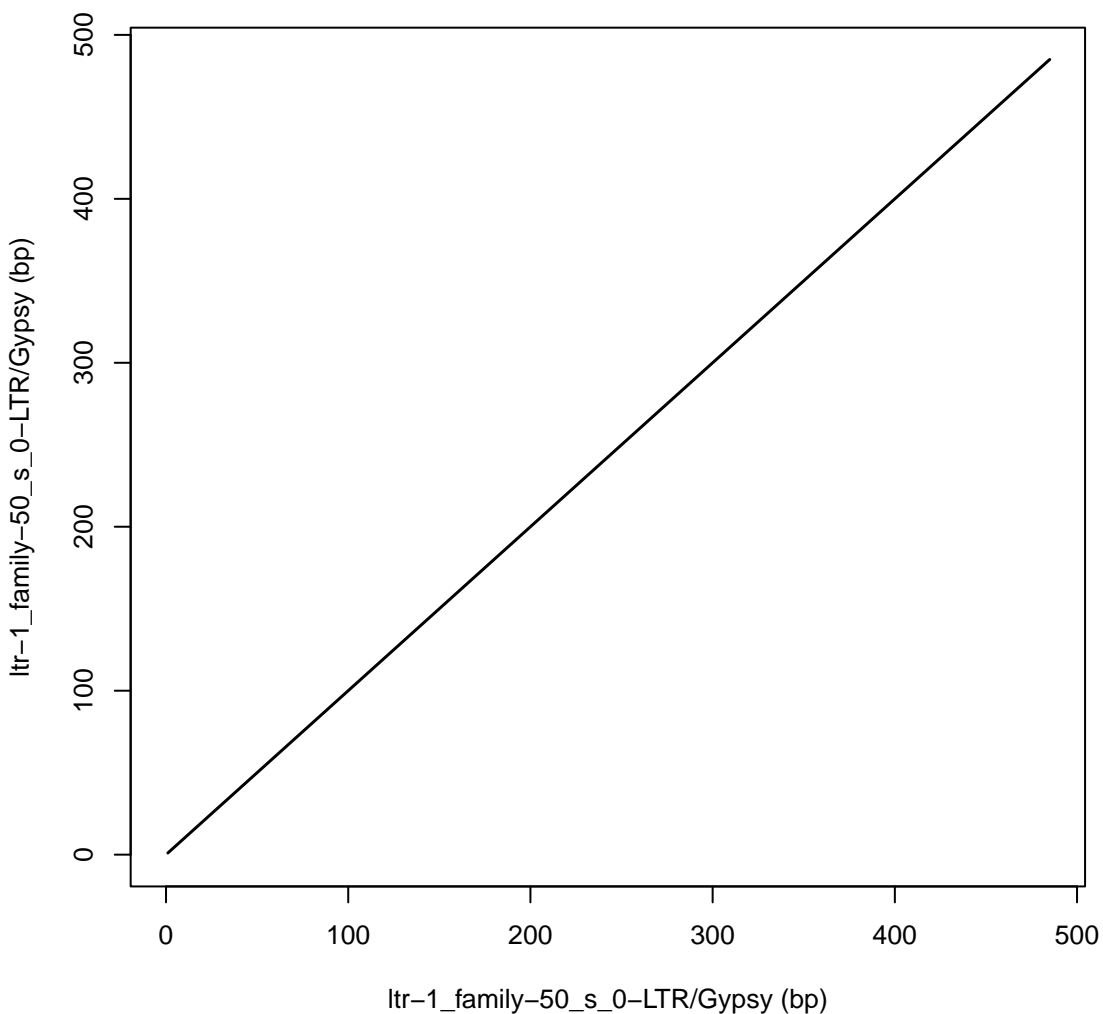
TE: ltr-1_family-50_s_0-LTR/Gypsy
consensus size: 485bp; fragments: 133; full length: 31 (>=436.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

