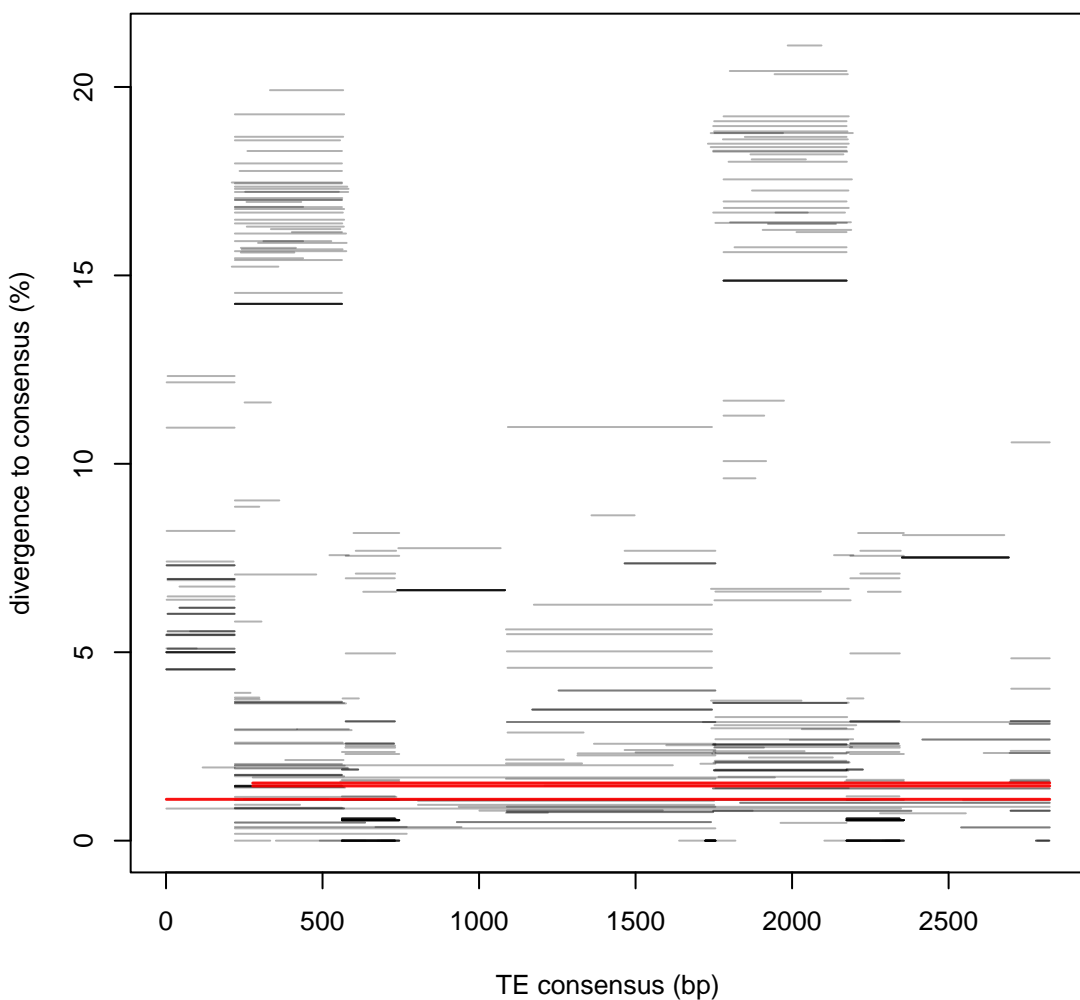
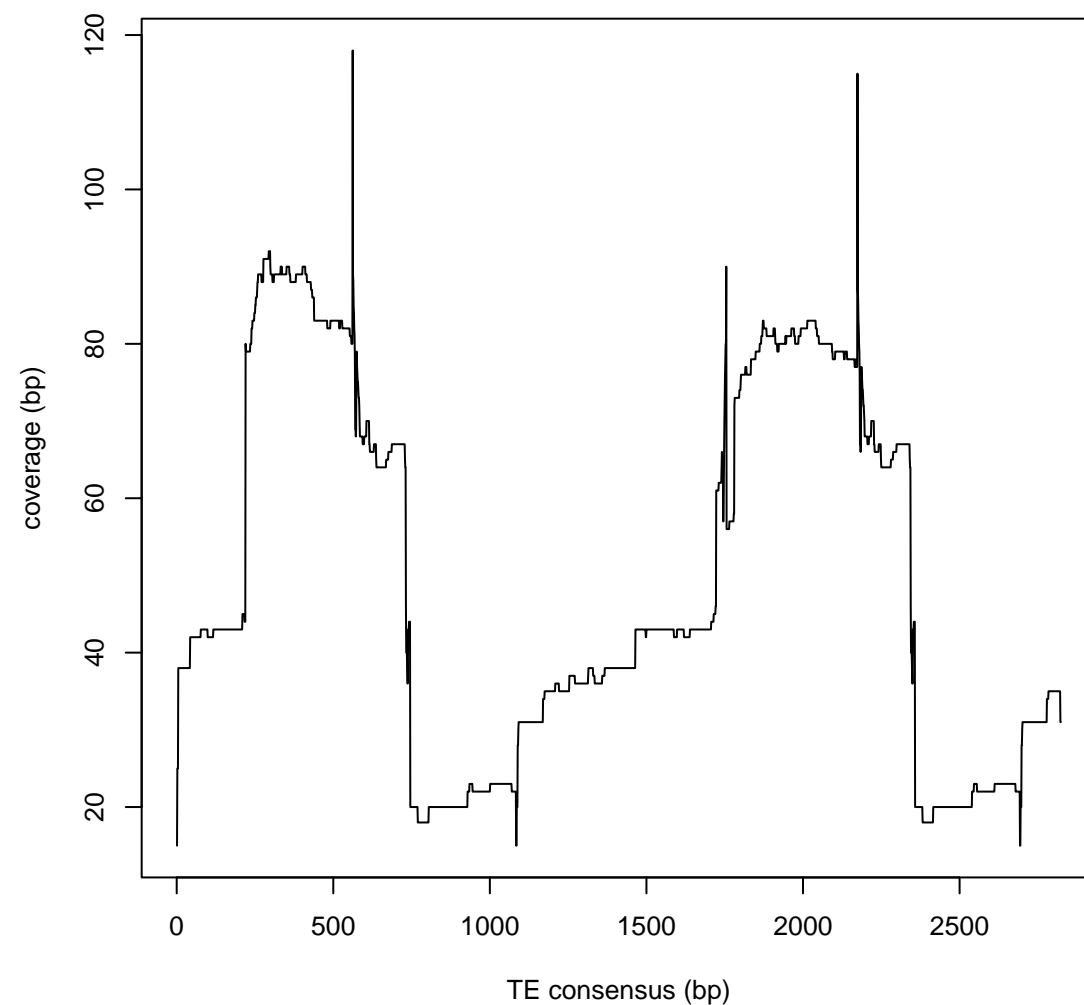


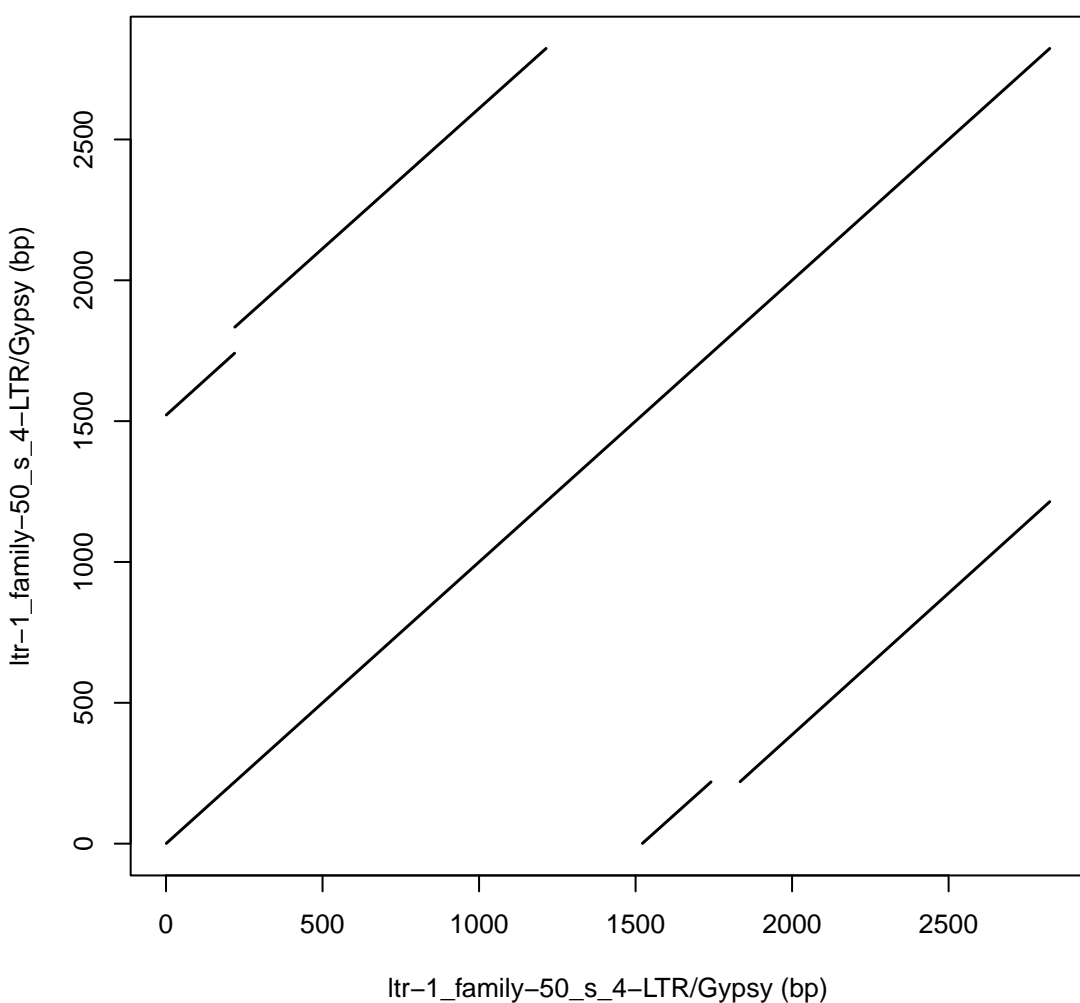
TE: ltr-1_family-50_s_4-LTR/Gypsy
consensus size: 2823bp; fragments: 450; full length: 3 (>=2540.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

