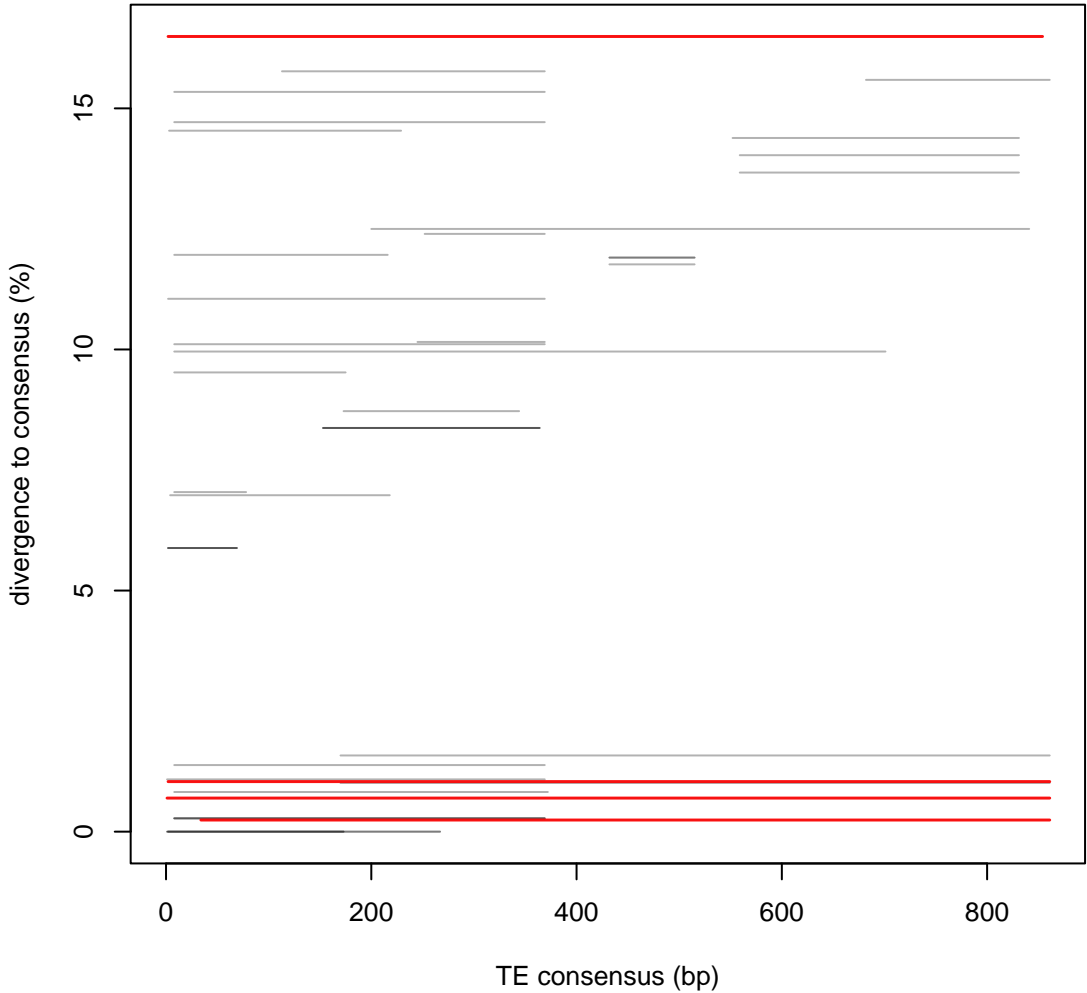
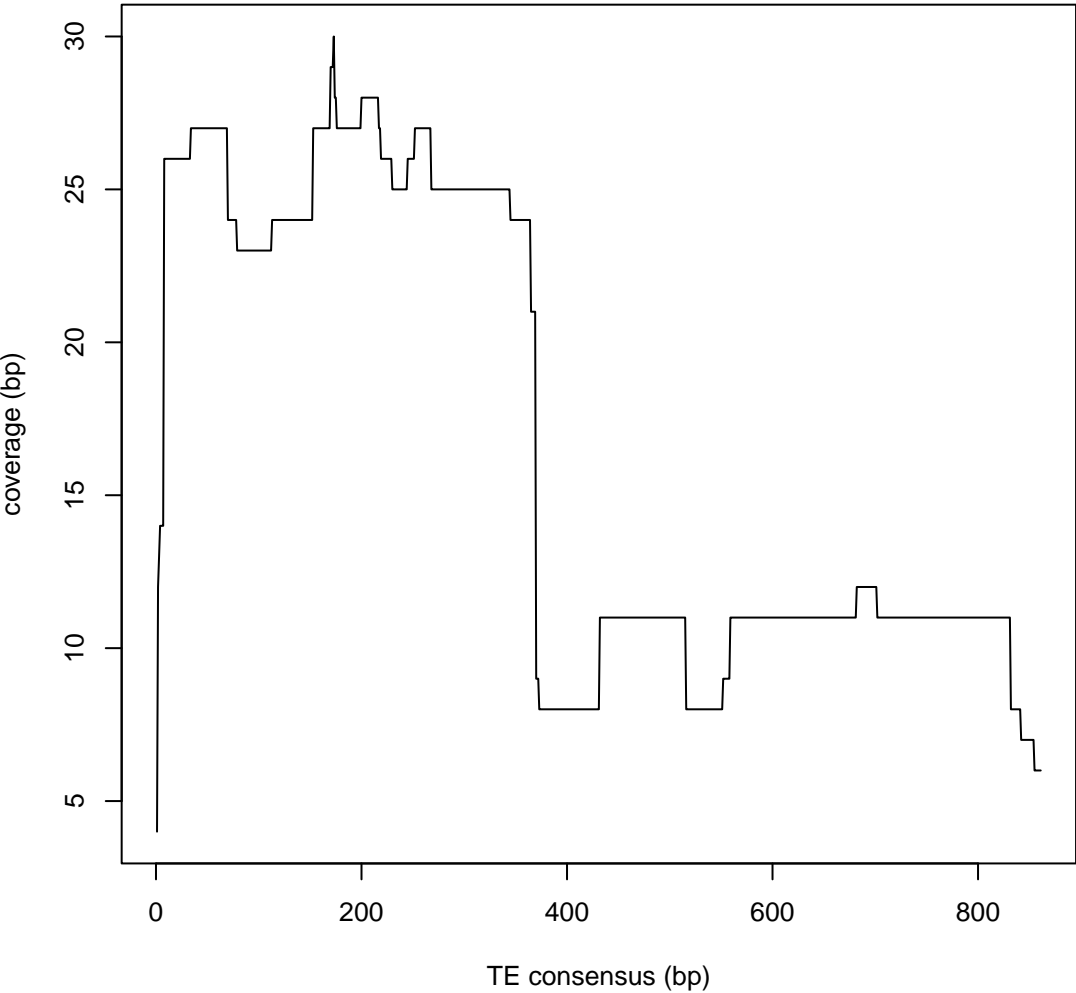


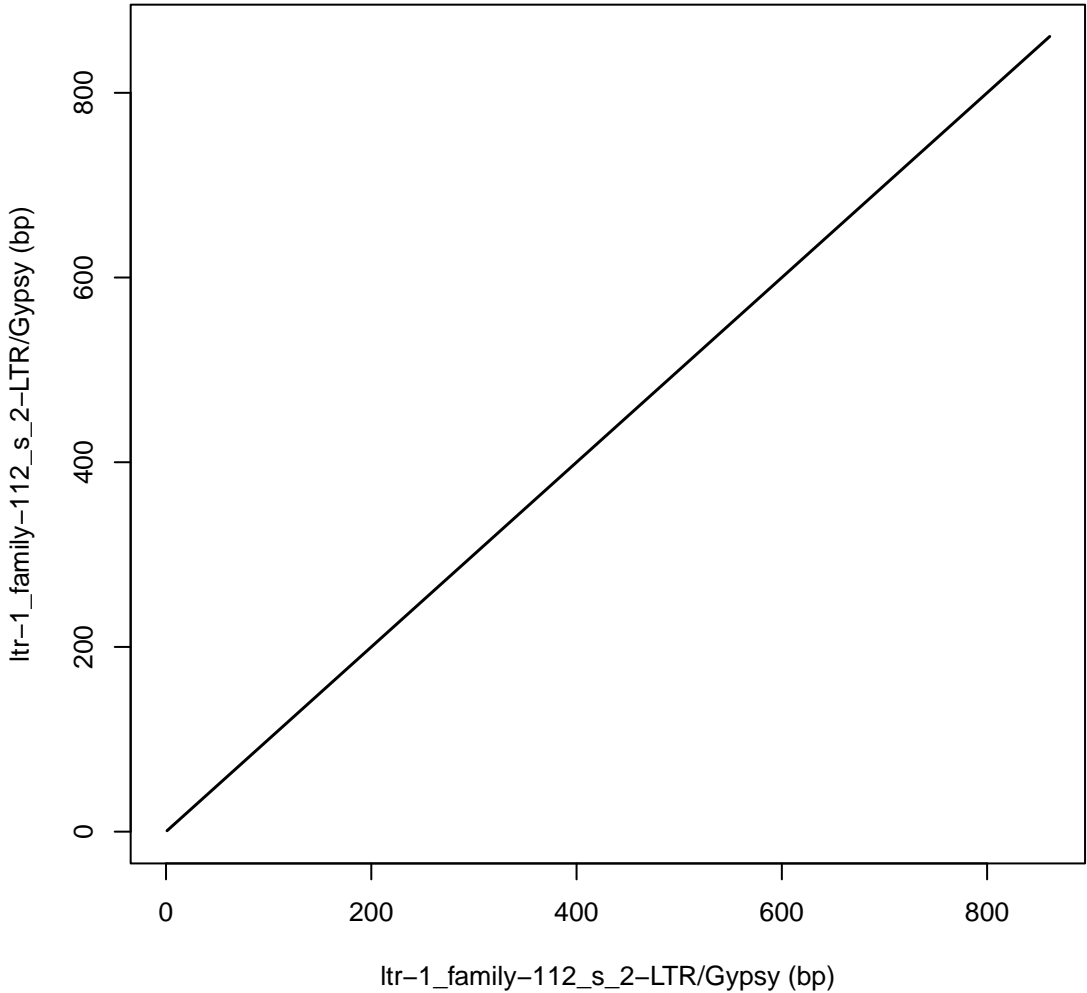
TE: ltr-1_family-112_s_2-LTR/Gypsy
consensus size: 861bp; fragments: 44; full length: 4 (>=774.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

