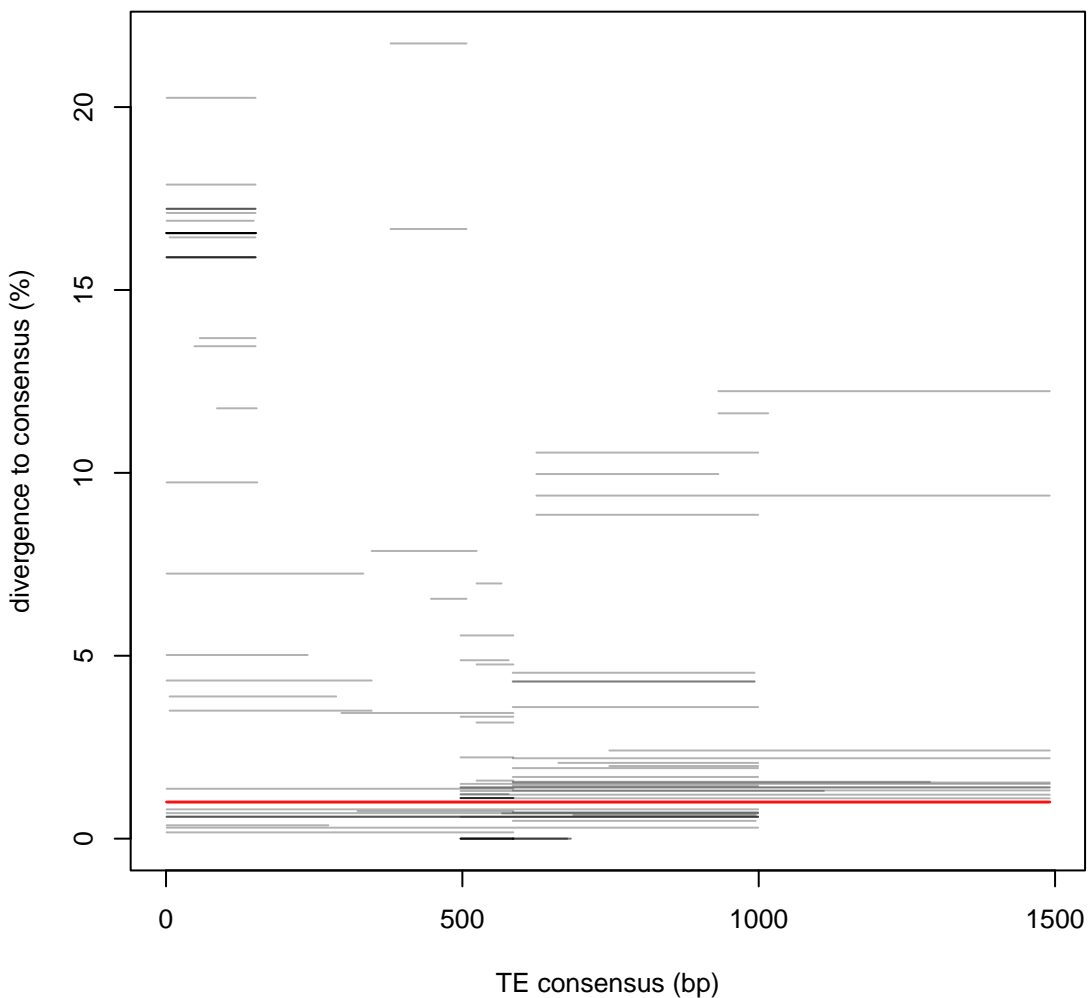
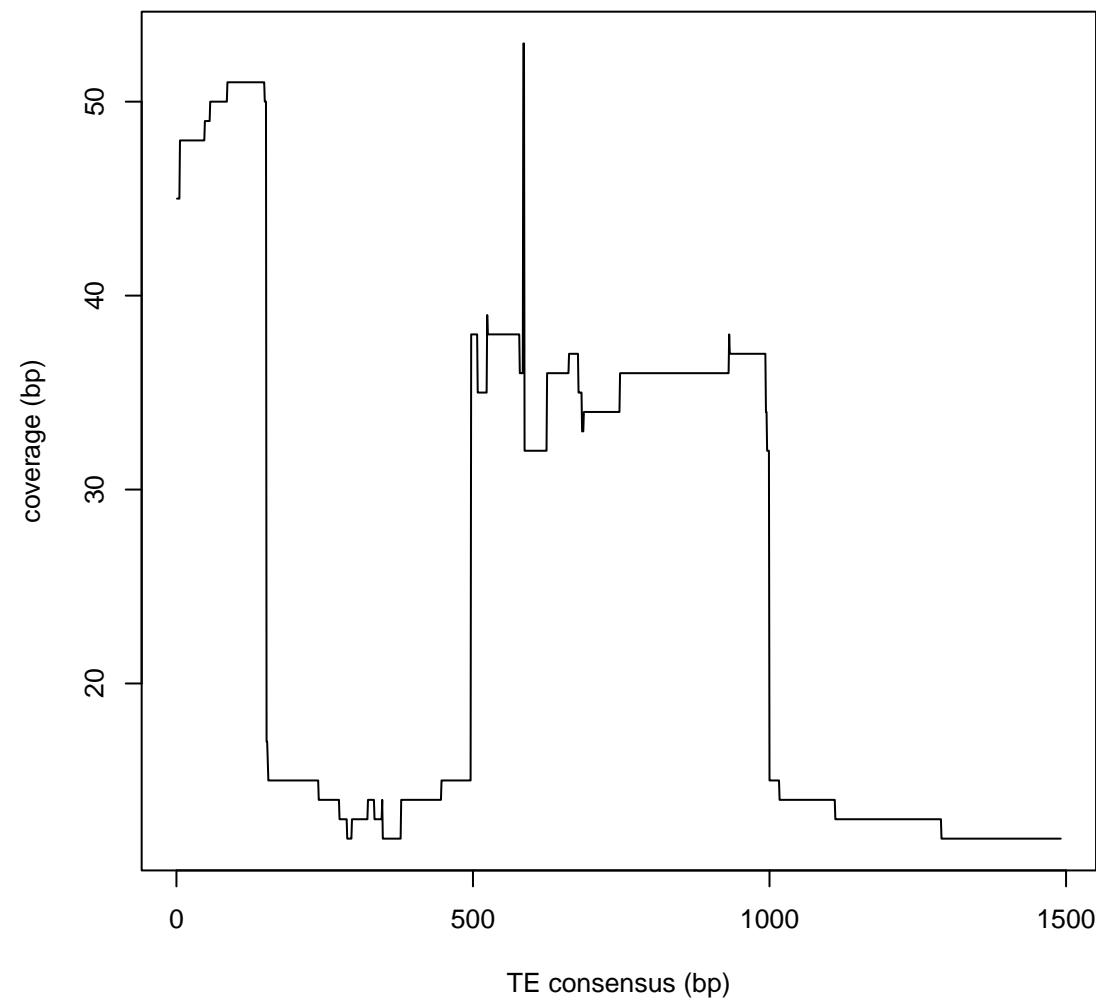


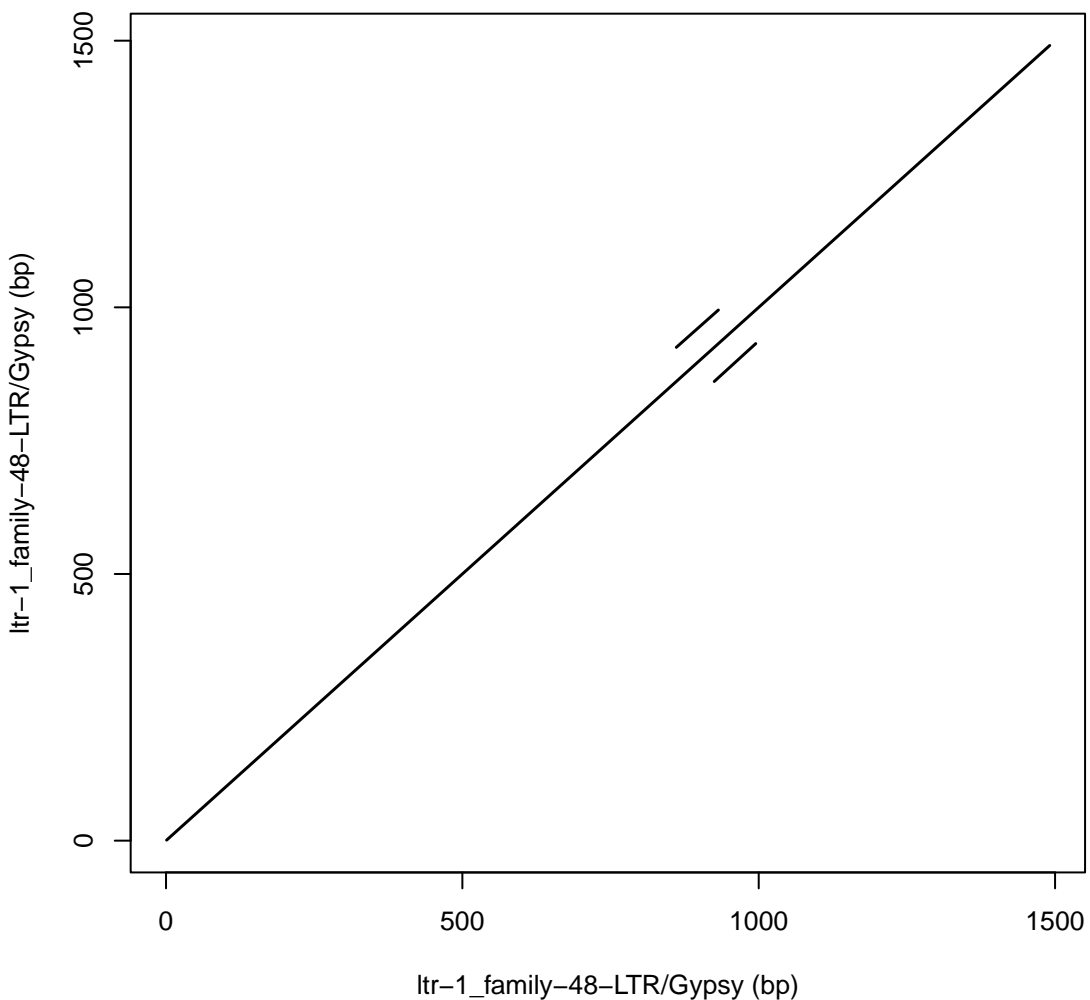
TE: ltr-1_family-48-LTR/Gypsy
consensus size: 1491bp; fragments: 112; full length: 1 (>=1341.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

