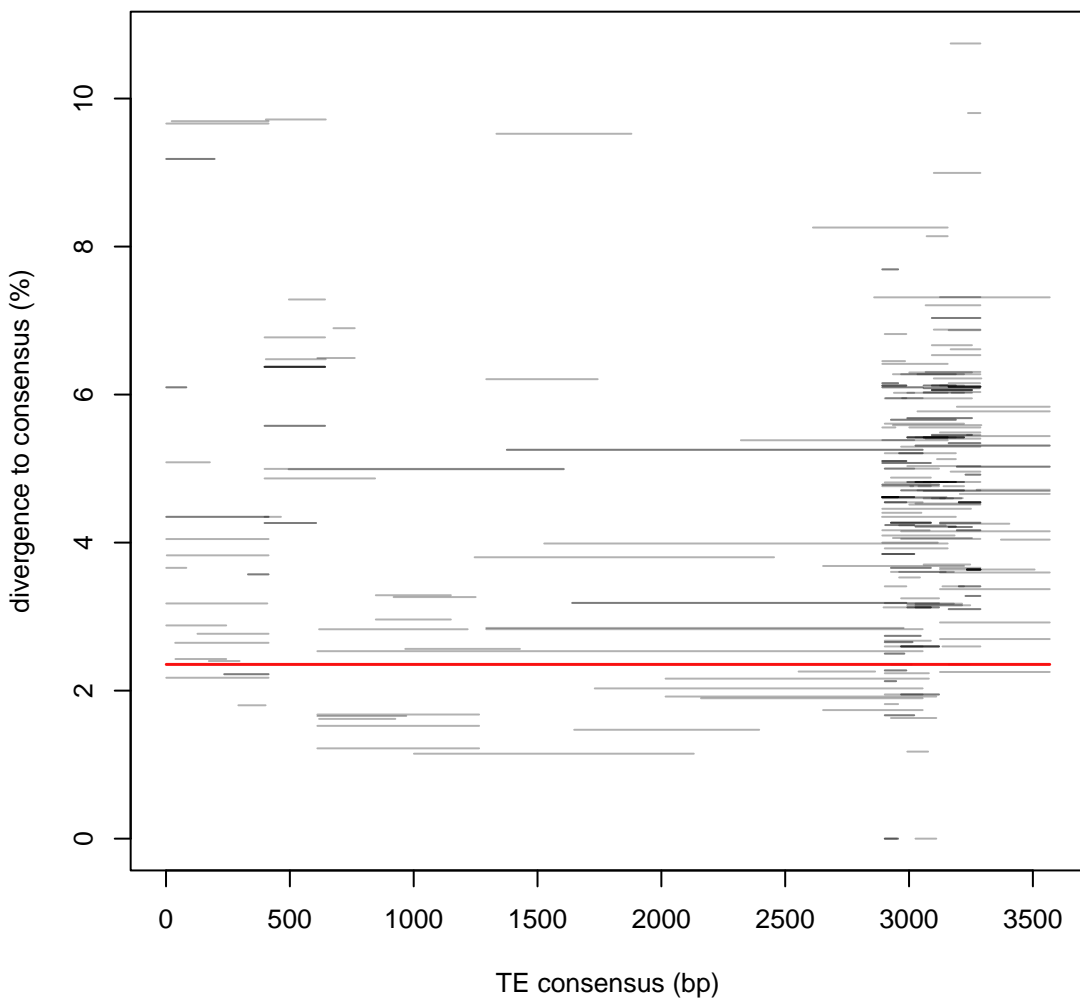
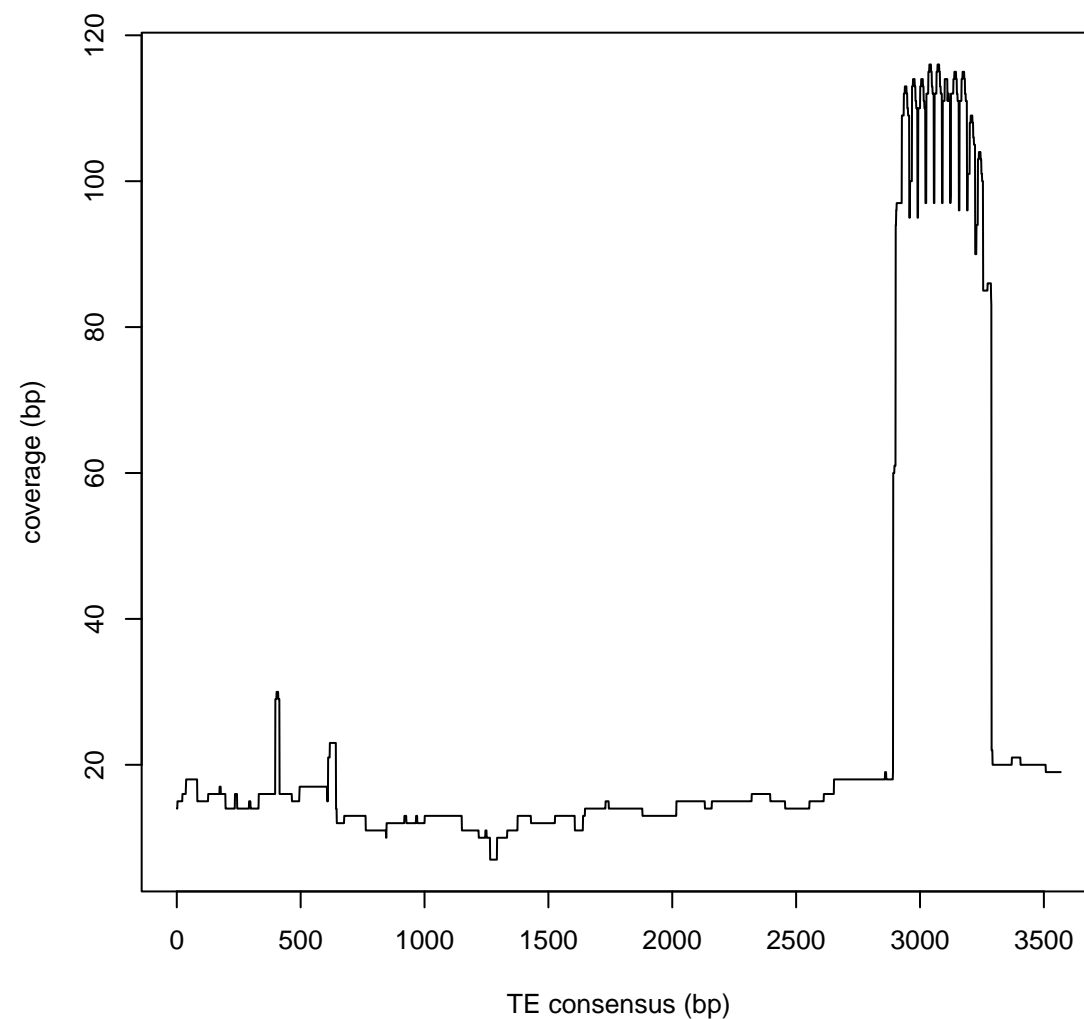


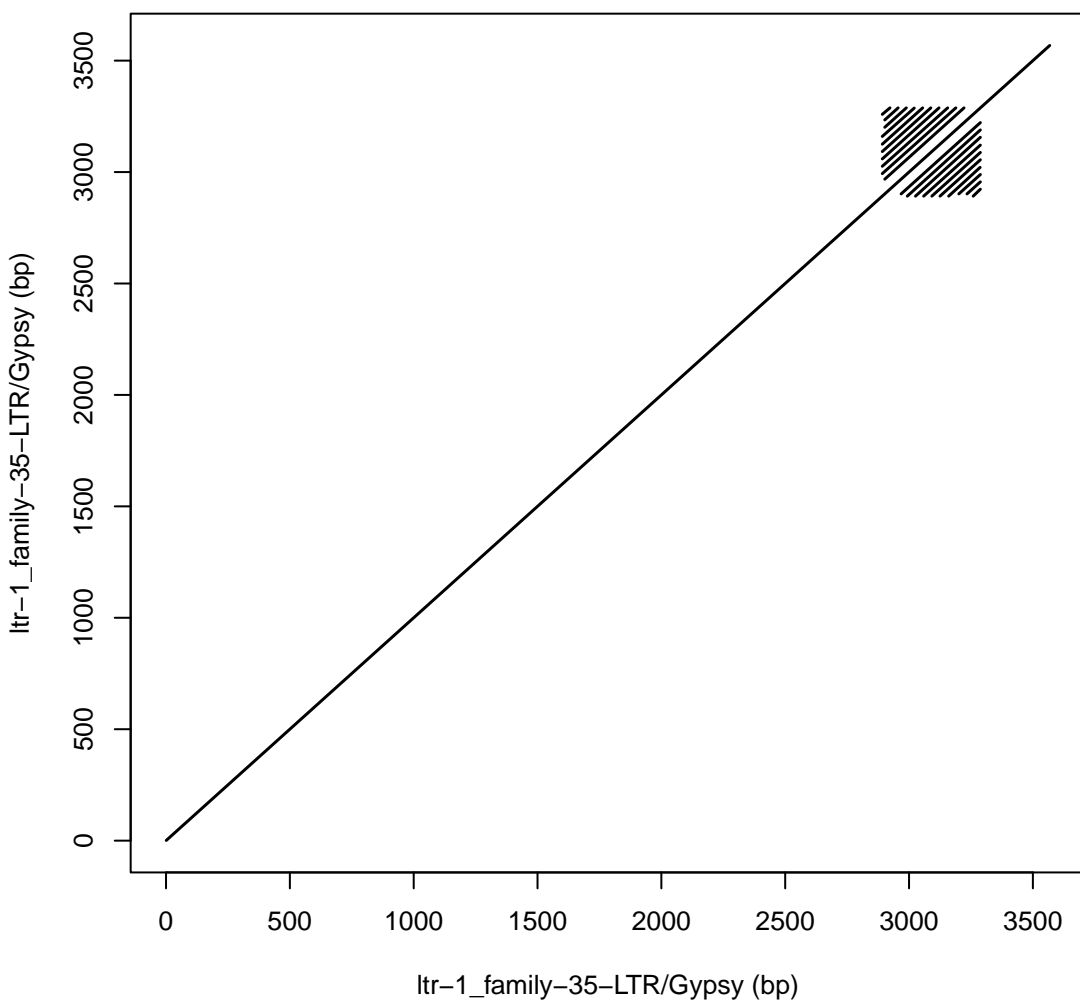
TE: ltr-1_family-35-LTR/Gypsy
consensus size: 3568bp; fragments: 332; full length: 1 (>=3211.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

