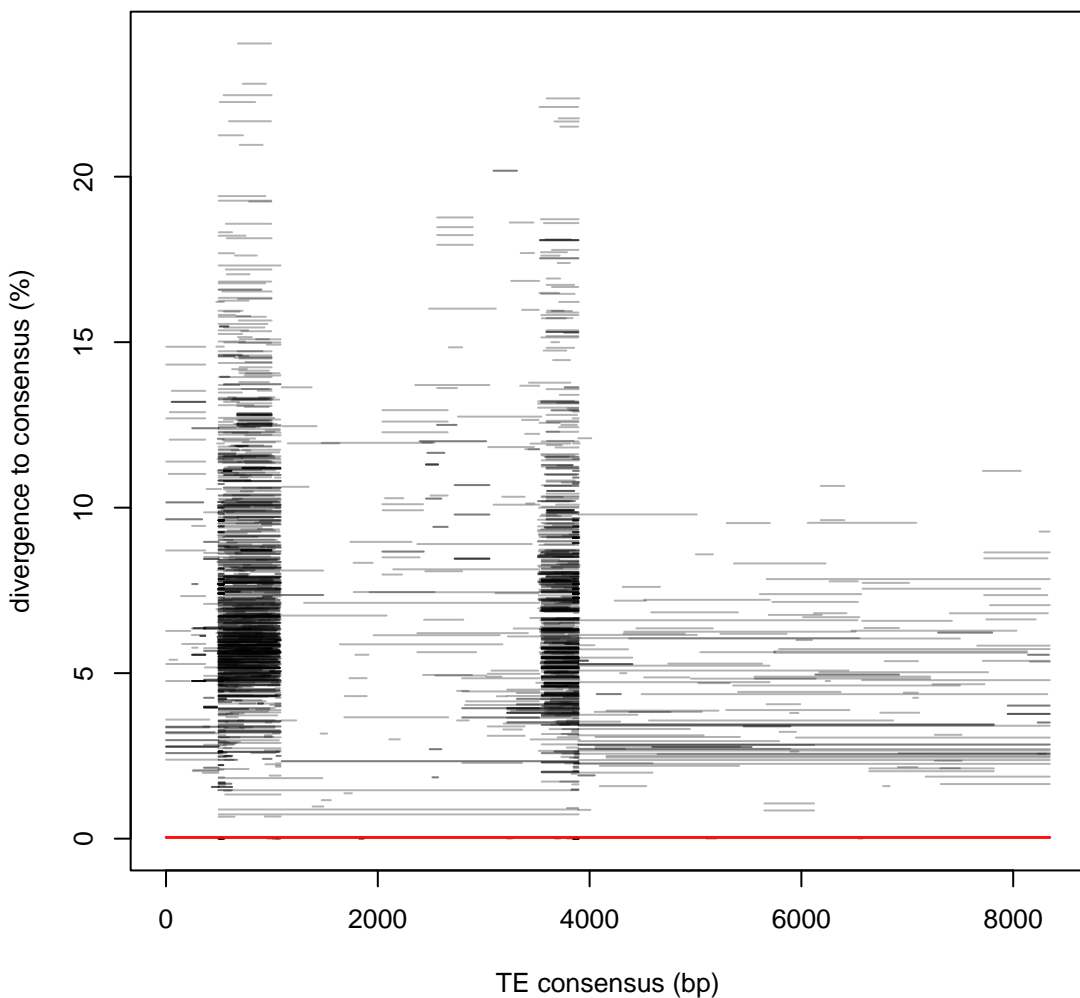
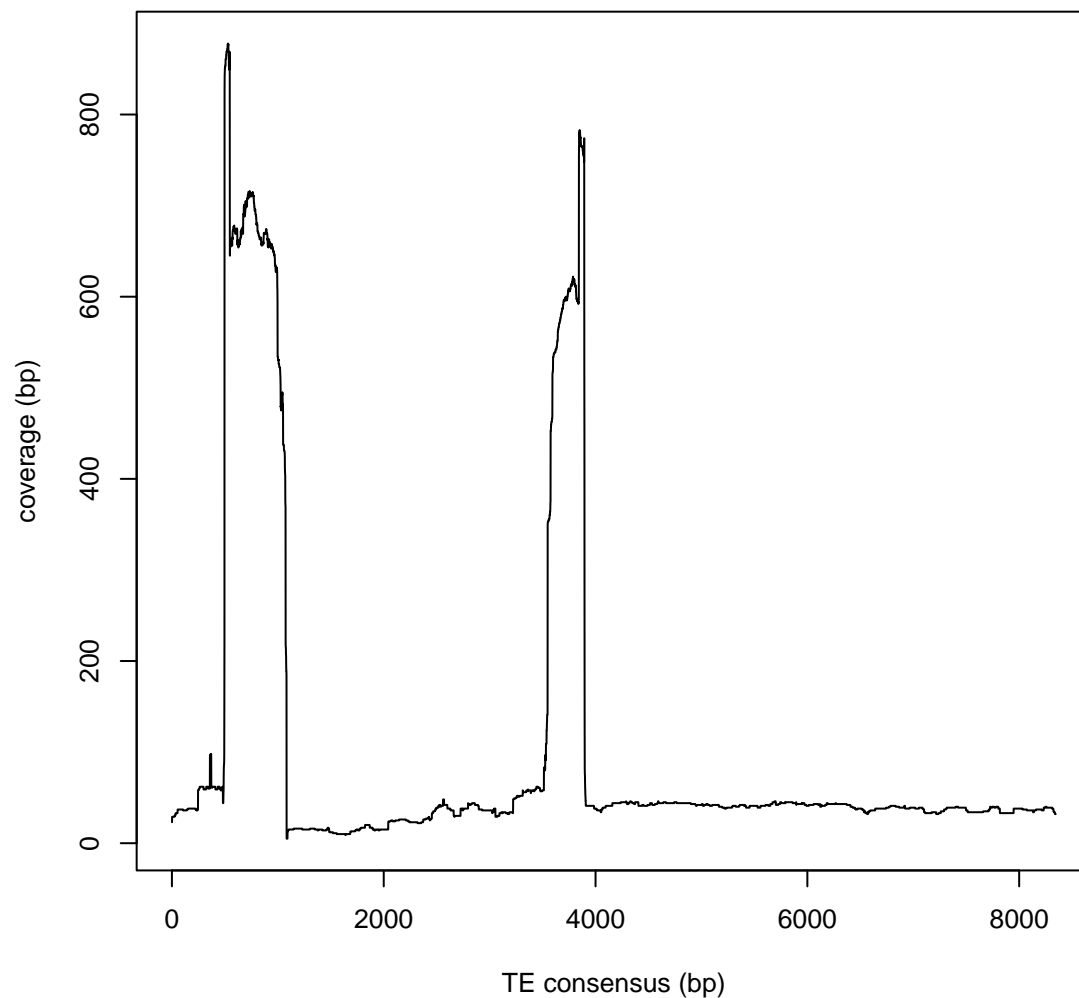


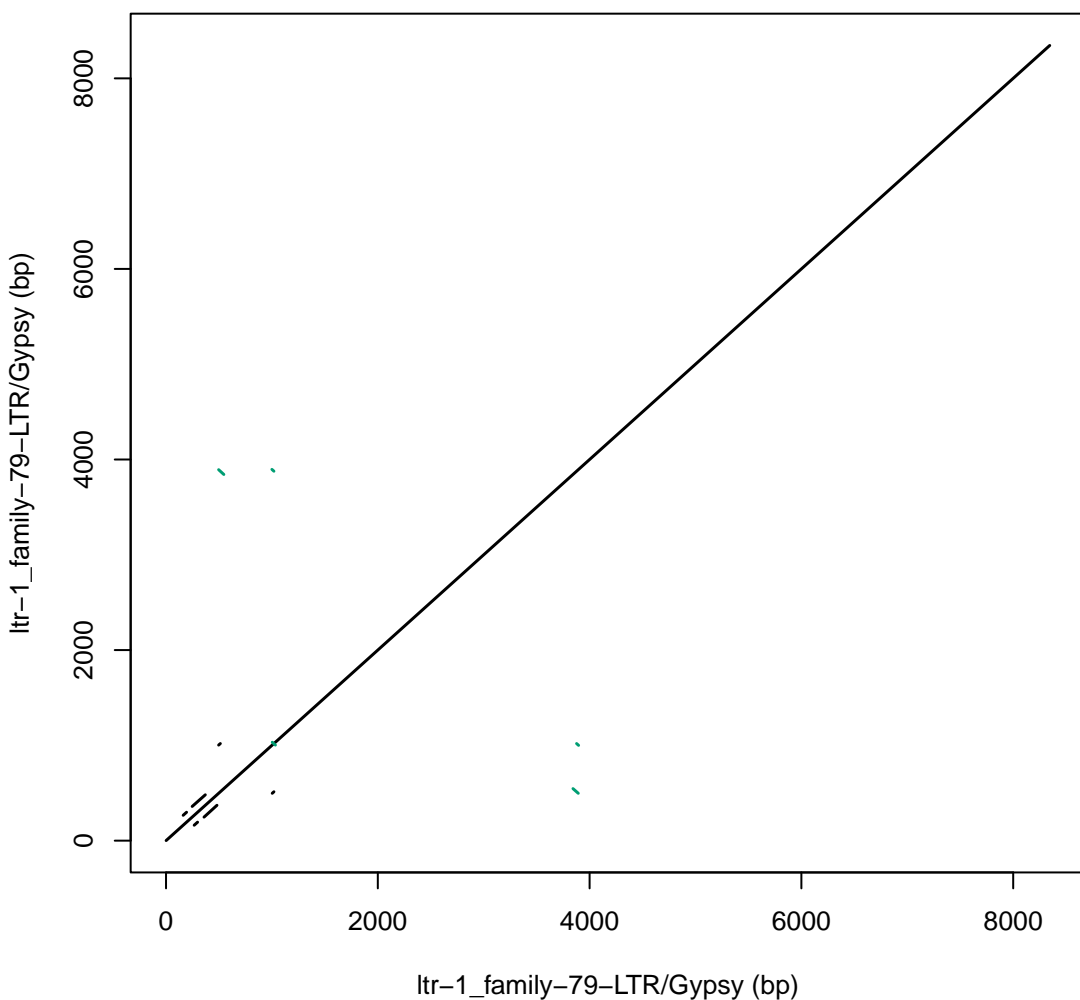
TE: ltr-1_family-79-LTR/Gypsy
consensus size: 8345bp; fragments: 2648; full length: 1 (>=7510.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

