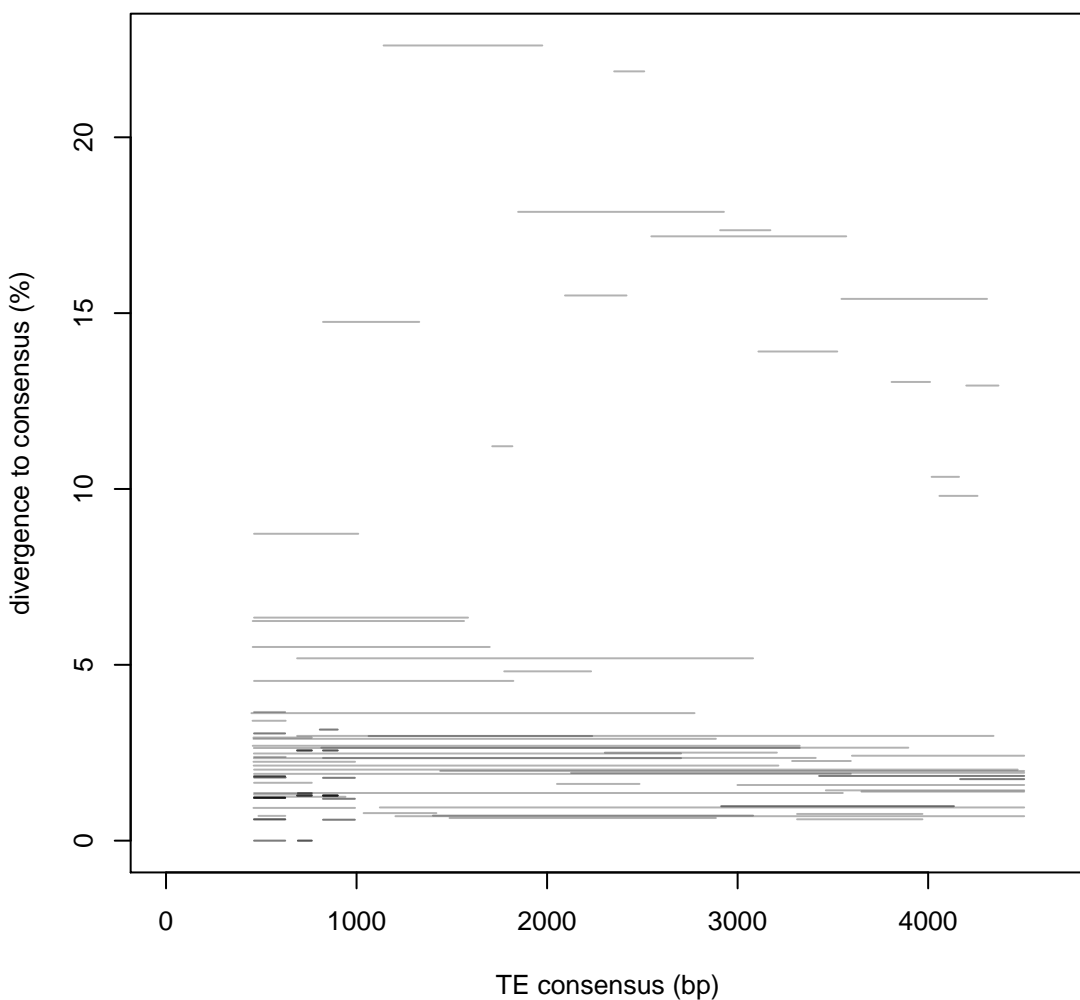
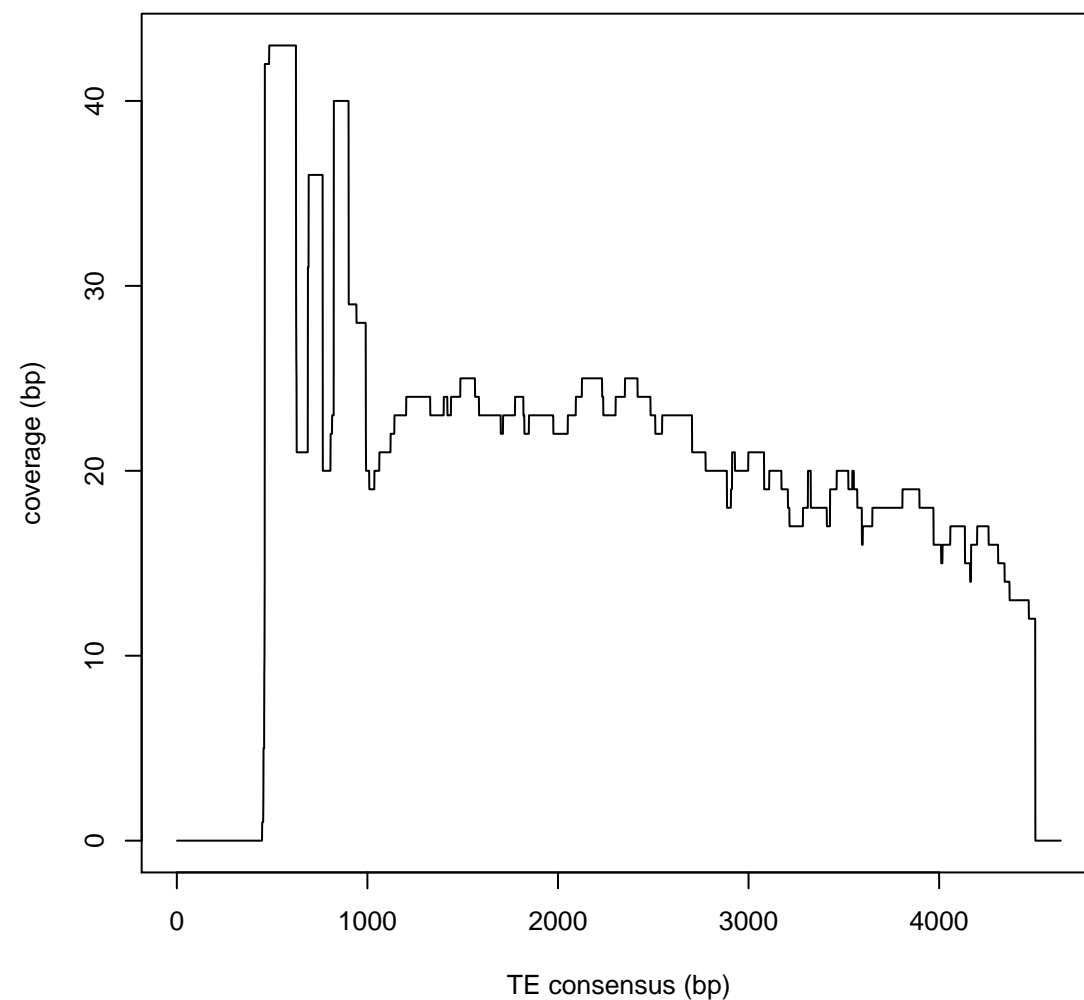


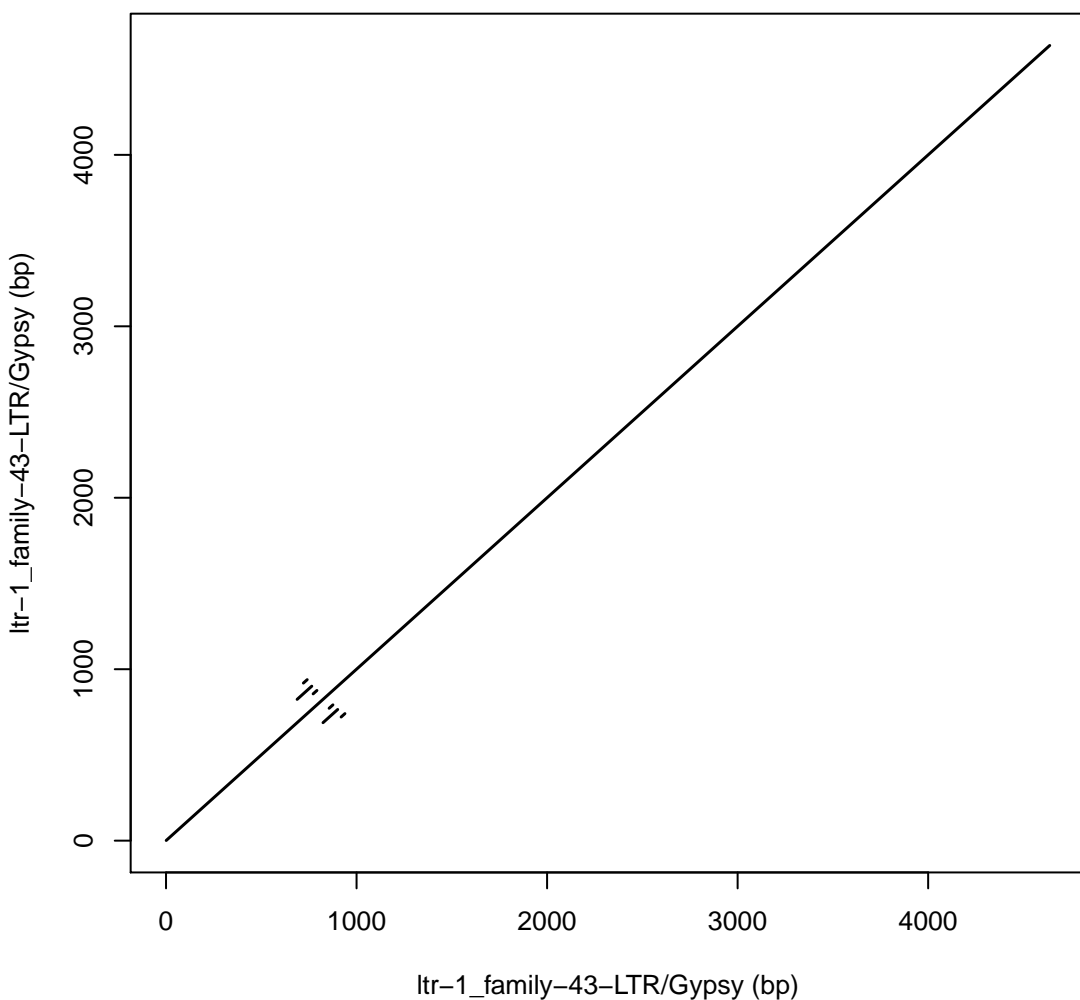
TE: ltr-1_family-43-LTR/Gypsy
consensus size: 4638bp; fragments: 115; full length: 0 (>=4174.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

