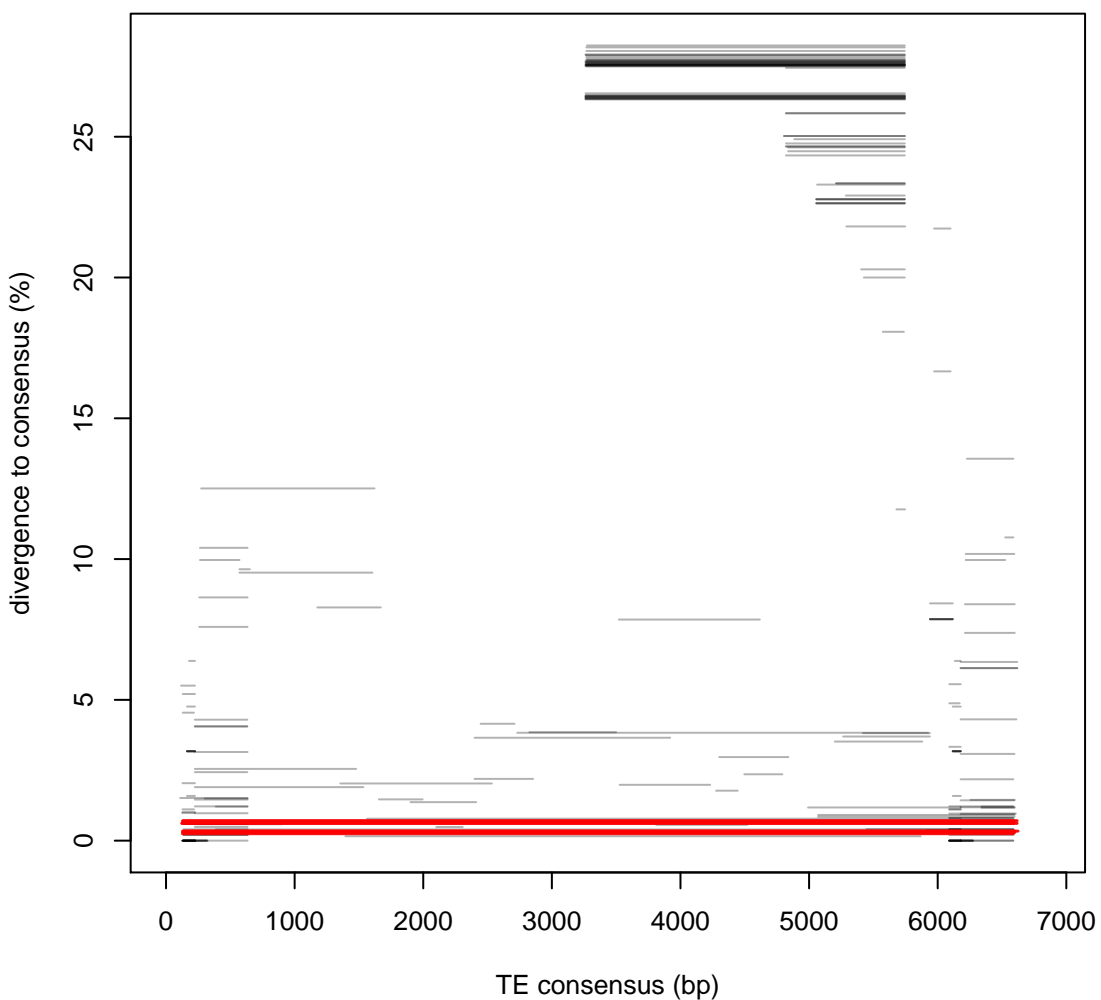
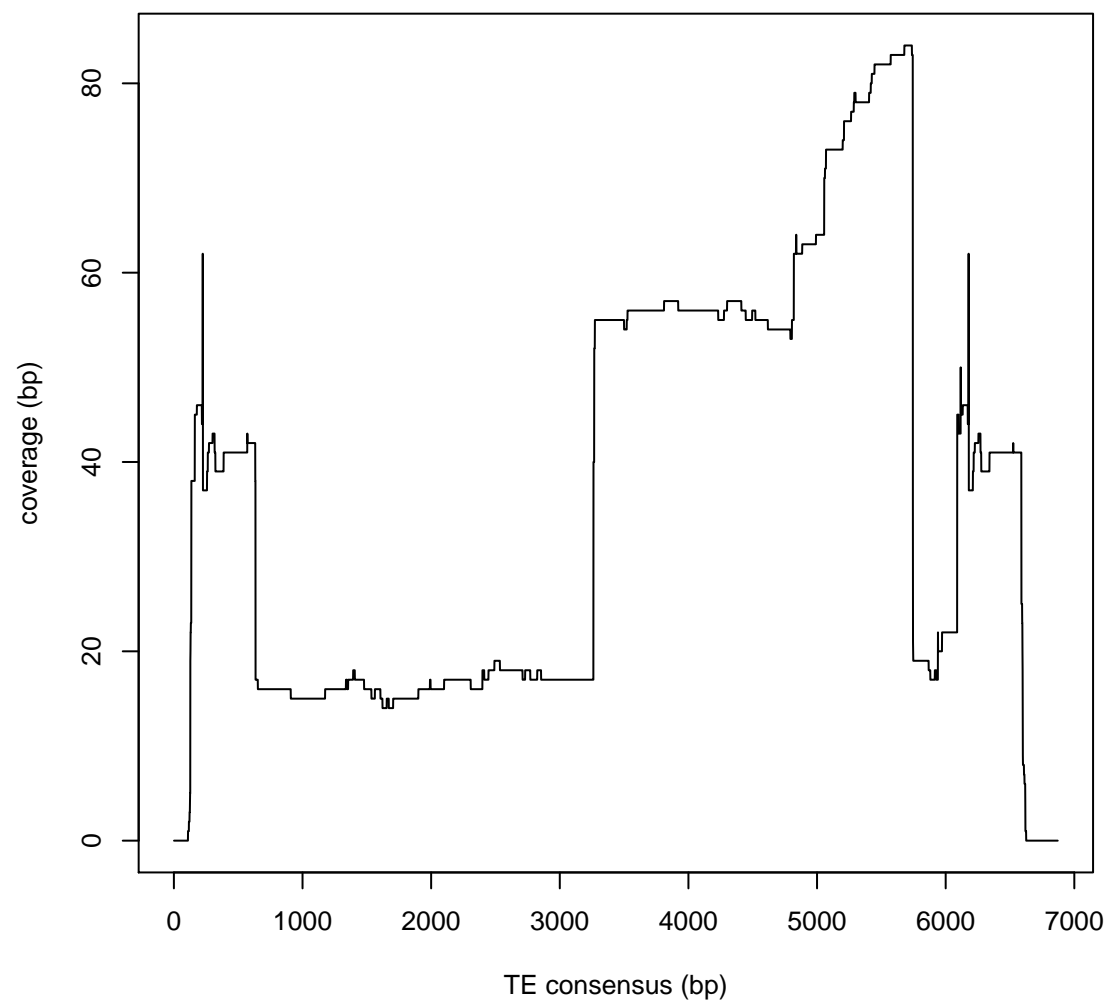


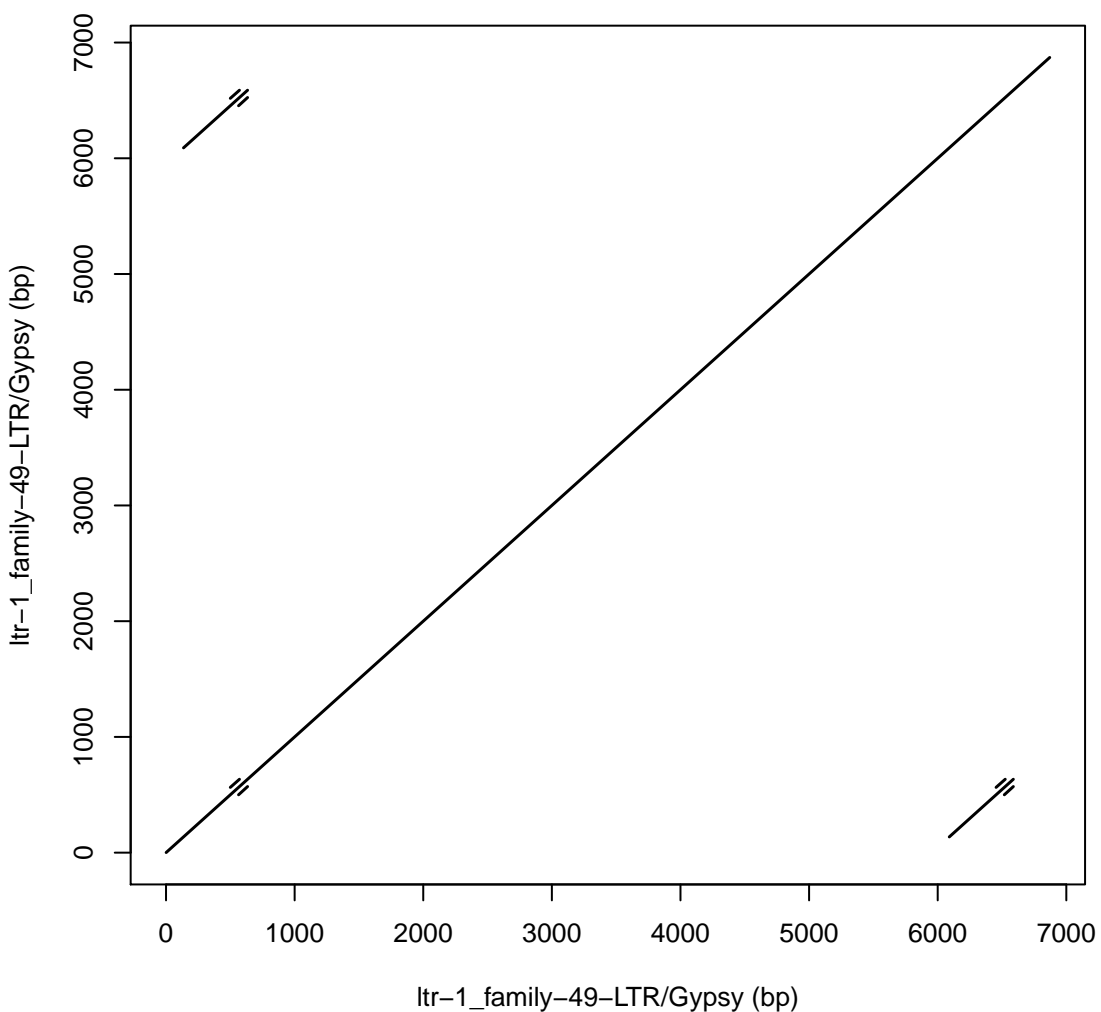
TE: ltr-1_family-49-LTR/Gypsy
consensus size: 6871bp; fragments: 234; full length: 8 (>=6183.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

