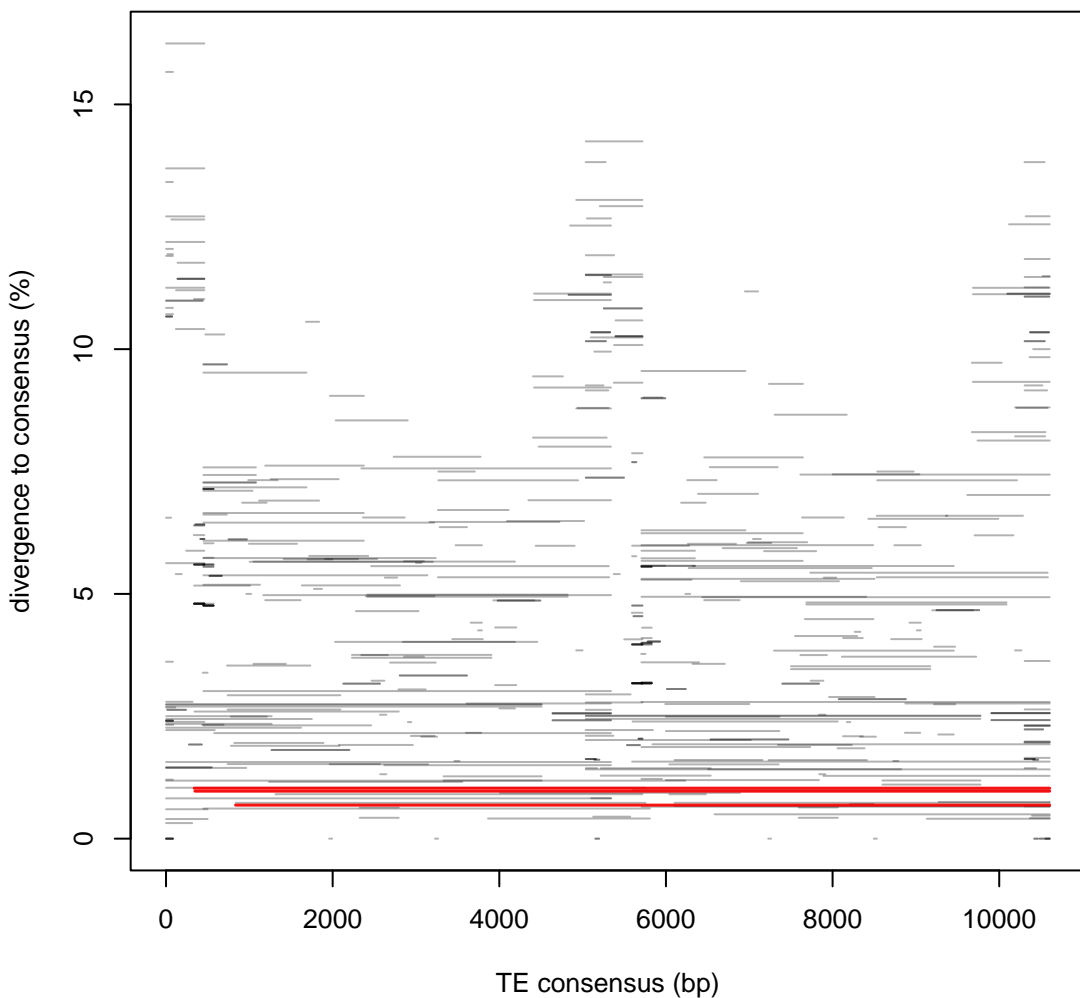
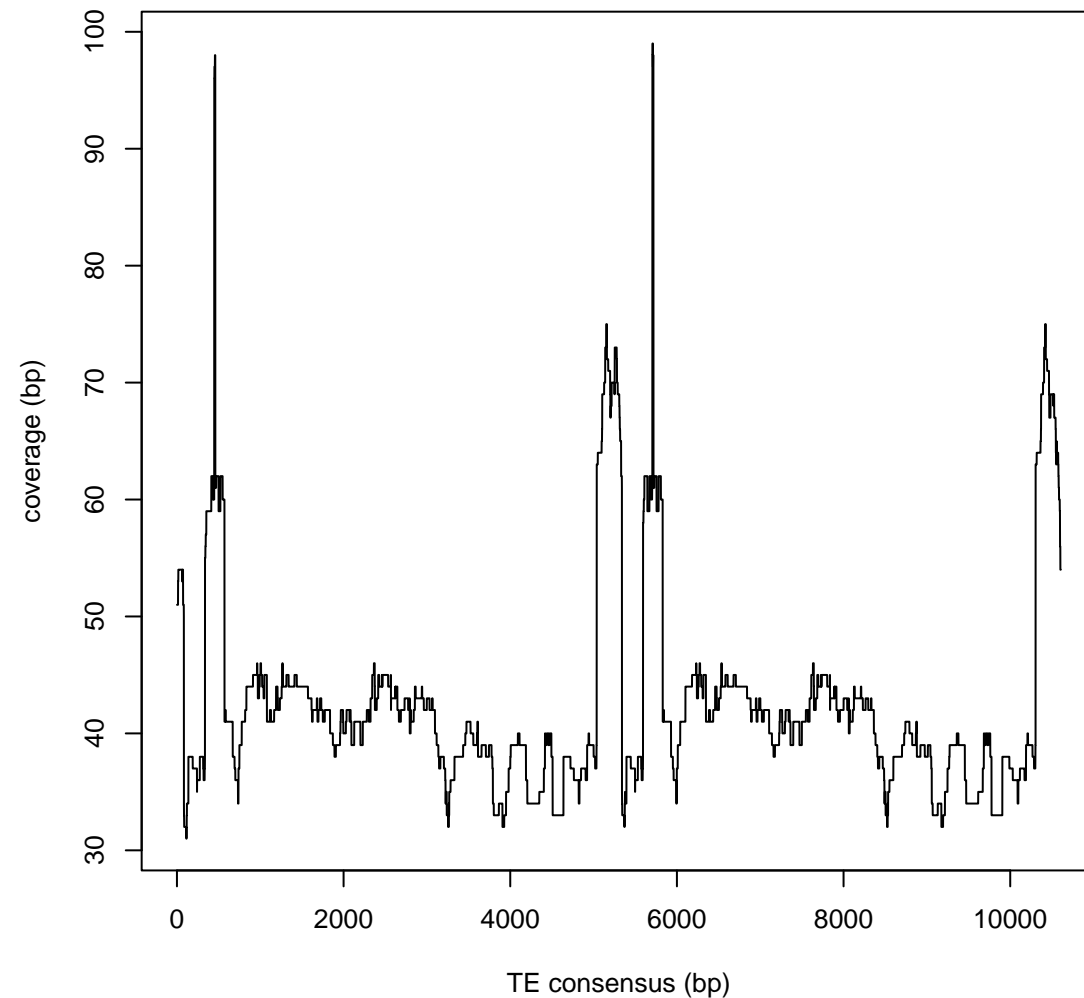


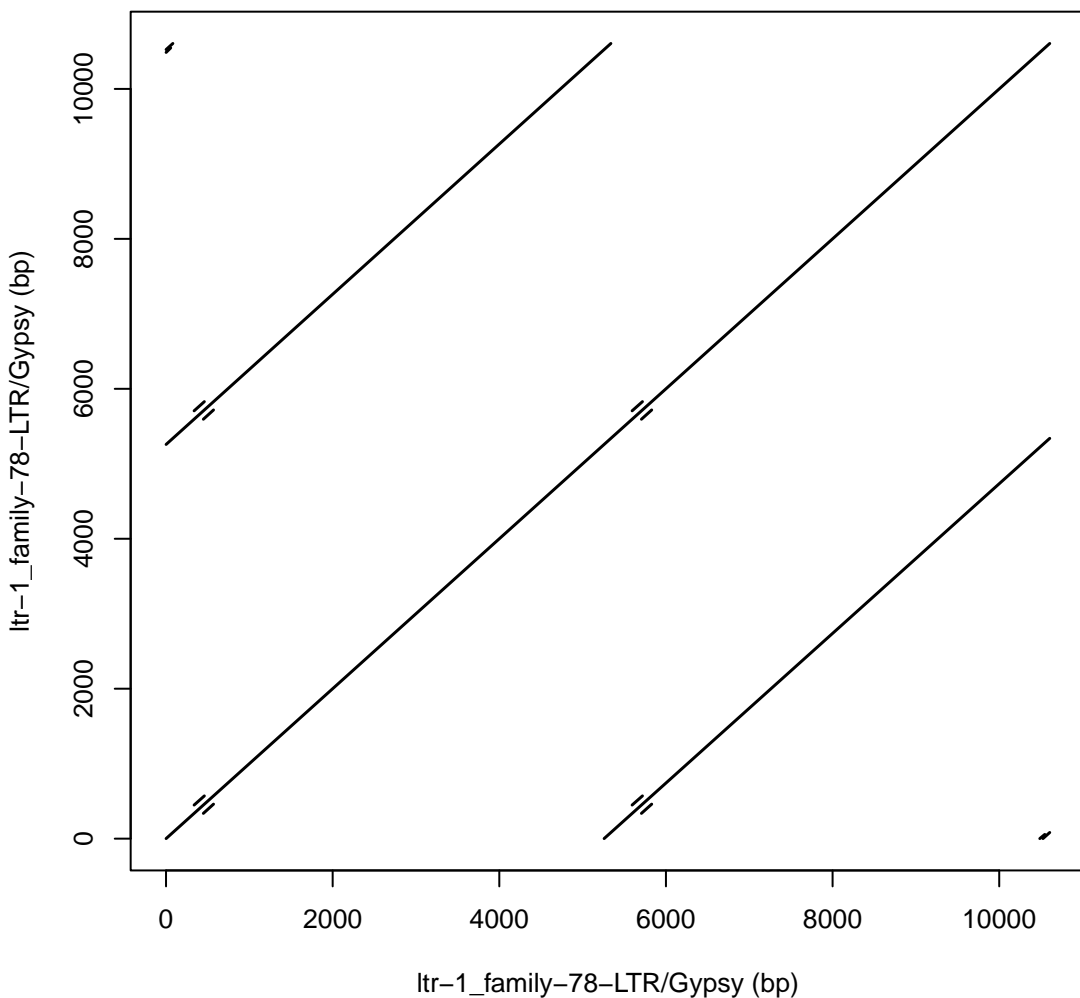
TE: ltr-1_family-78-LTR/Gypsy
consensus size: 10606bp; fragments: 540; full length: 3 (>=9545.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

