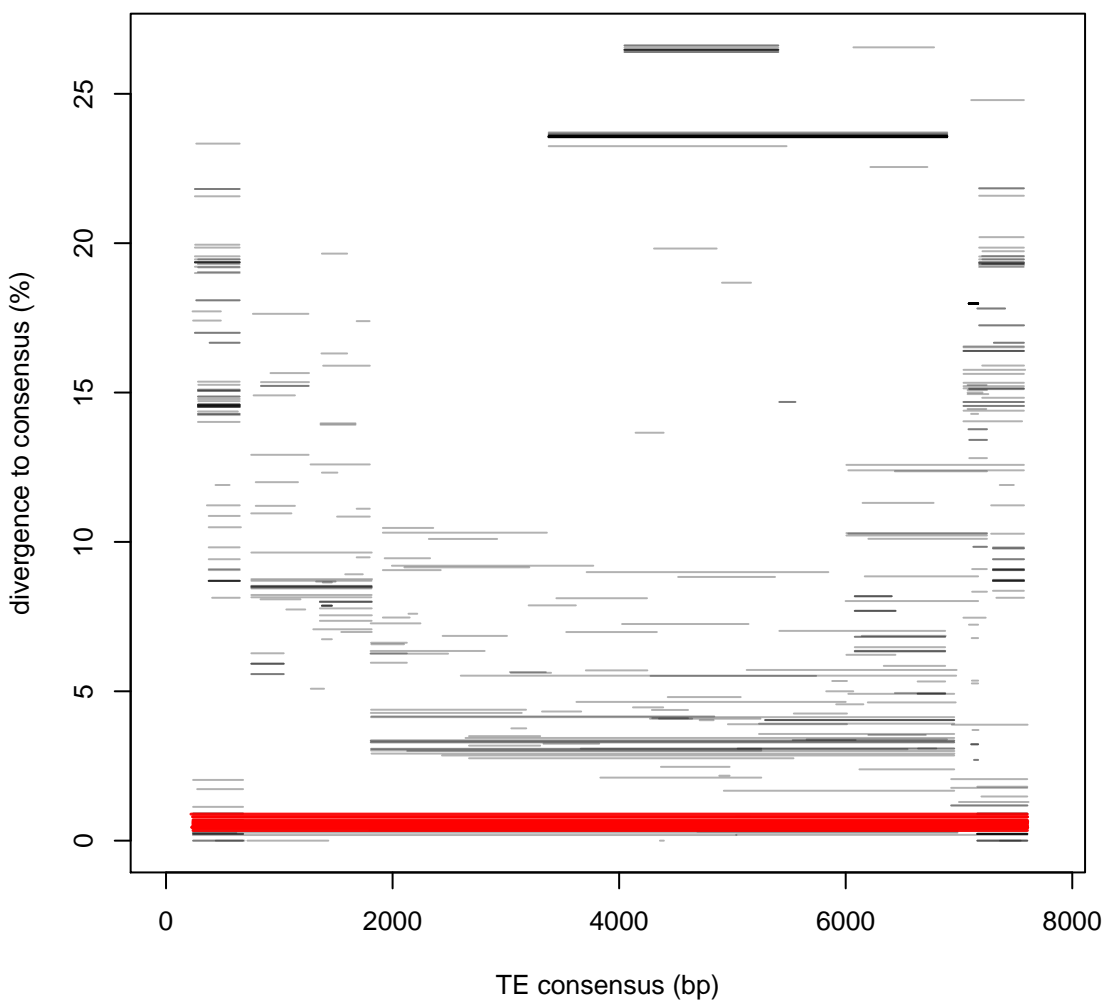
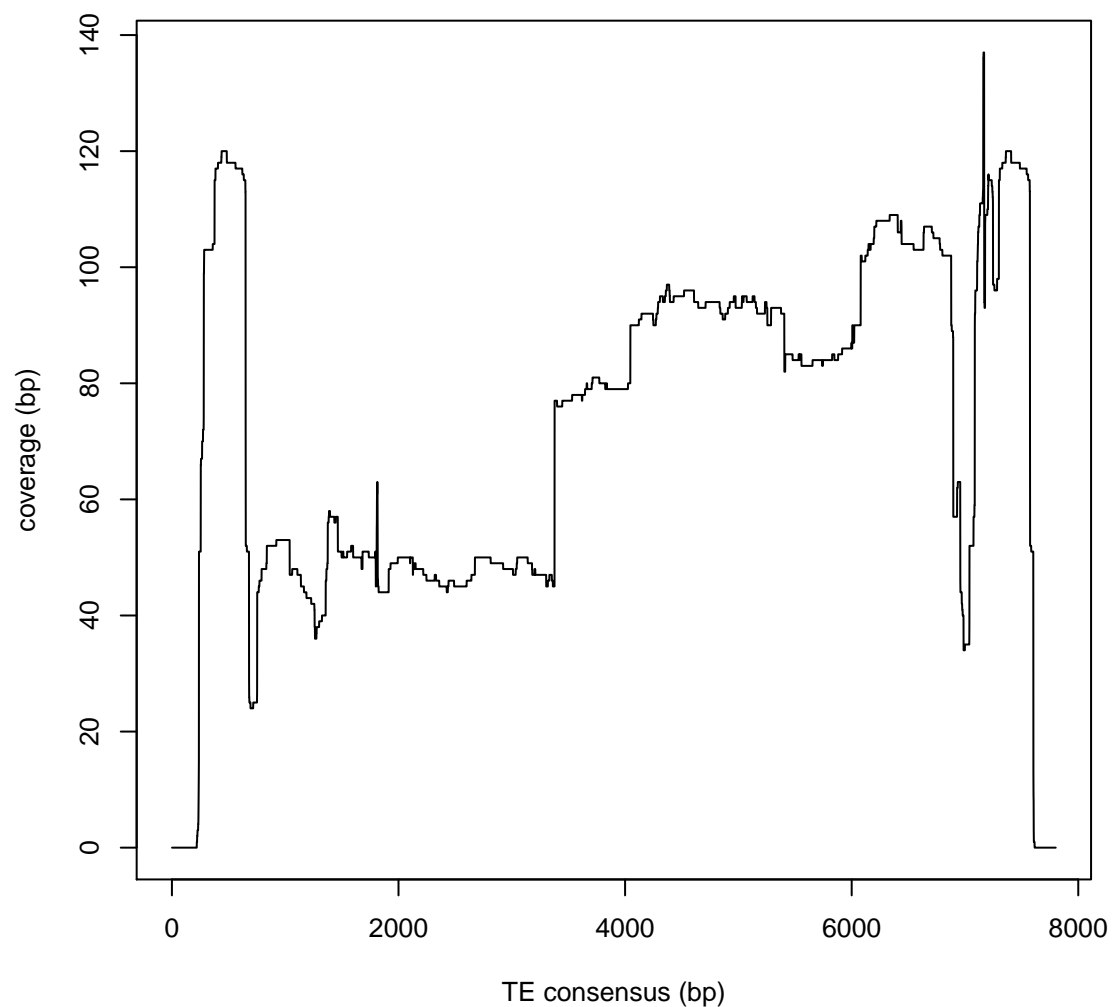


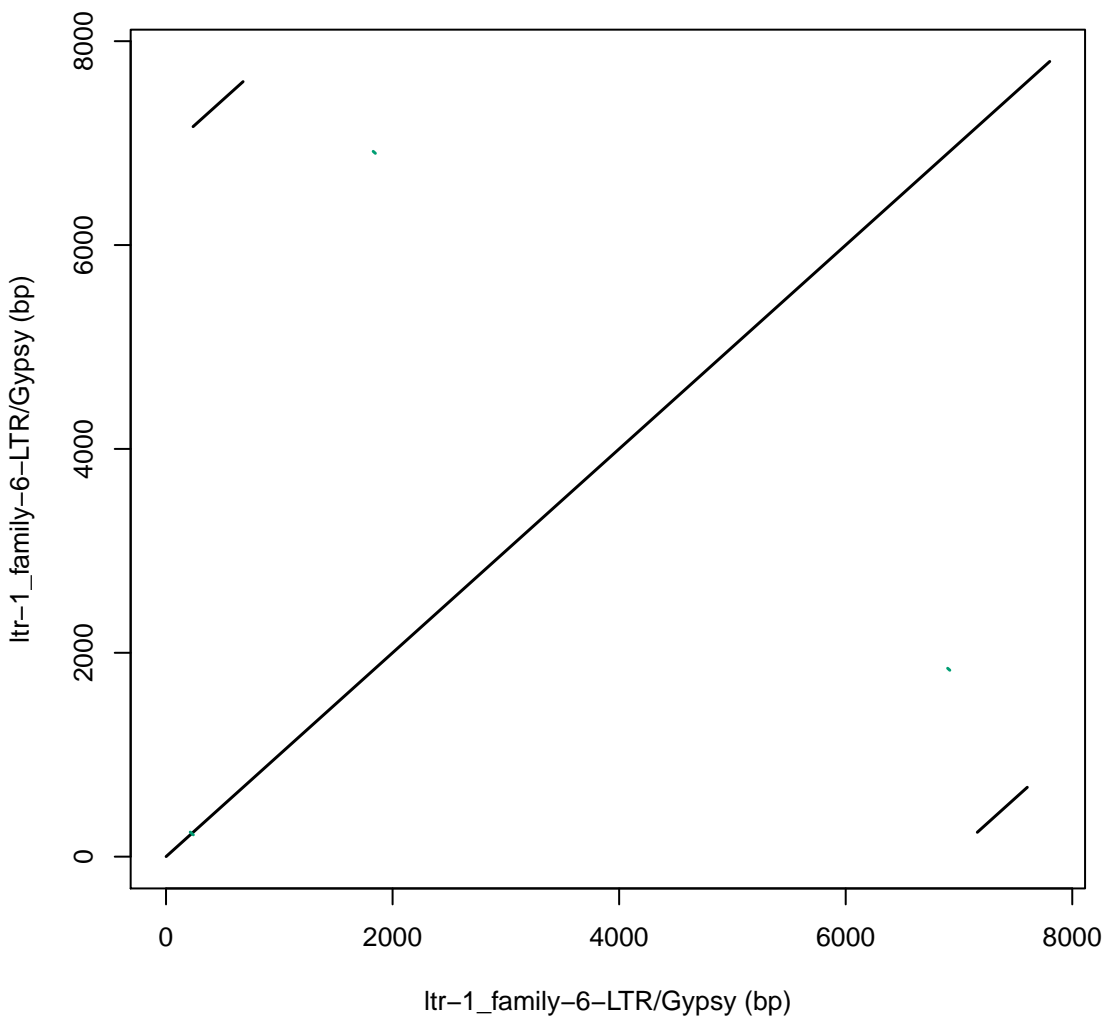
TE: ltr-1_family-6-LTR/Gypsy
consensus size: 7801bp; fragments: 502; full length: 17 (>=7020.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

