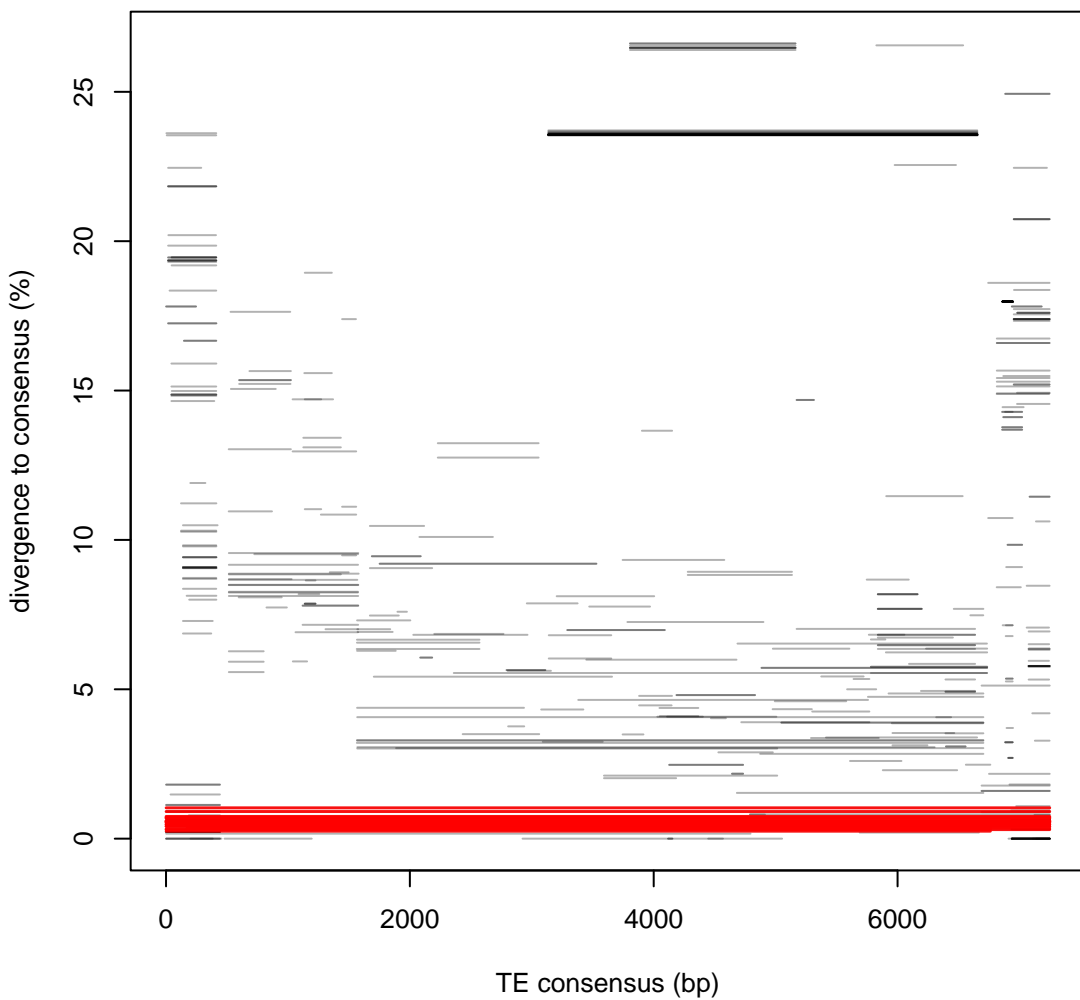
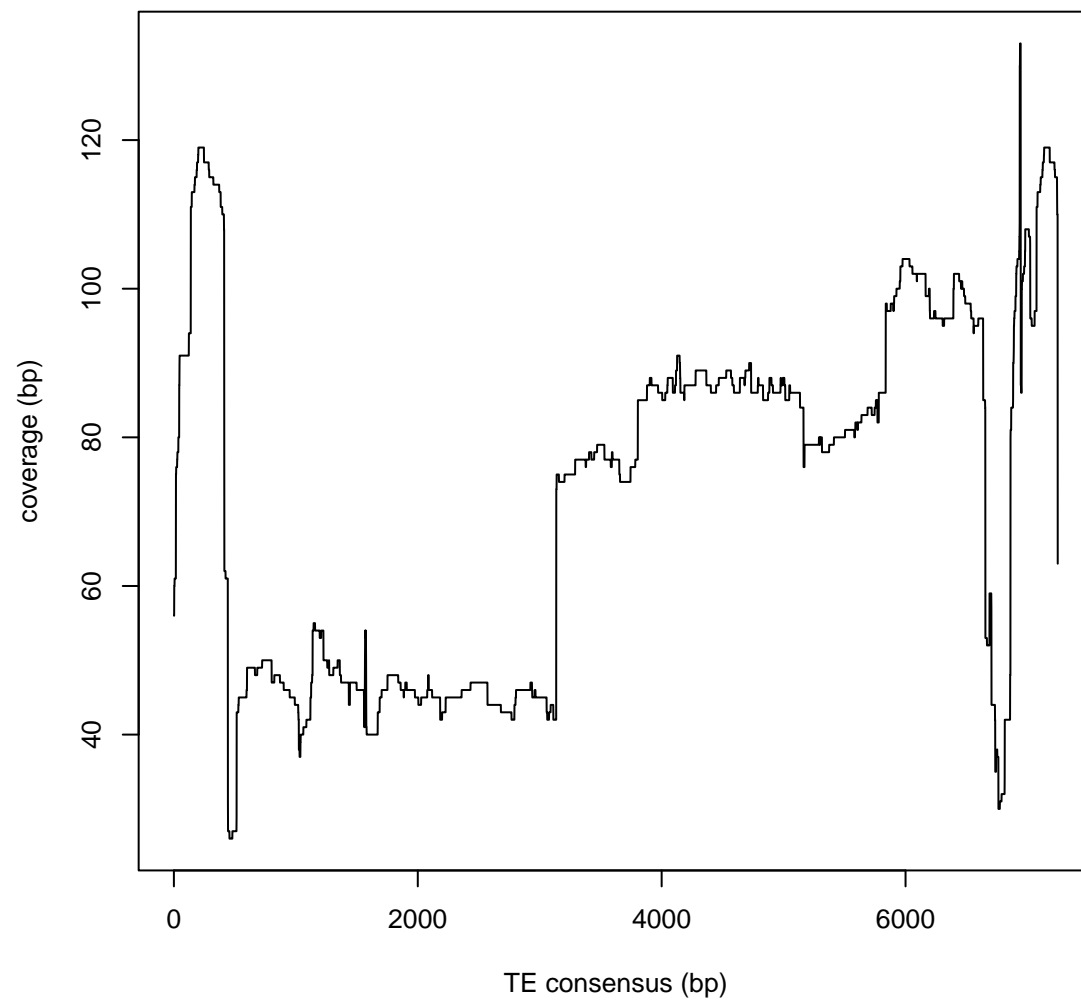


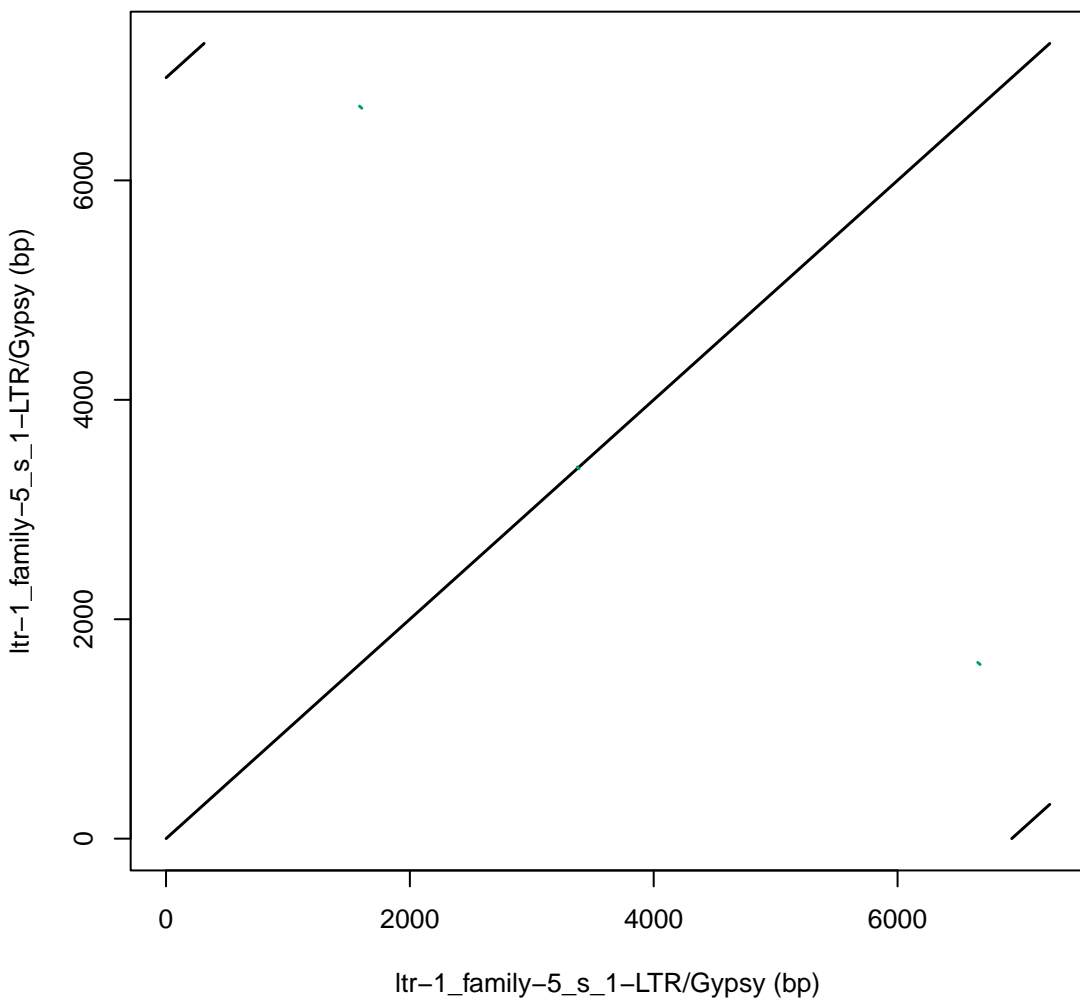
TE: ltr-1_family-5_s_1-LTR/Gypsy
consensus size: 7248bp; fragments: 512; full length: 22 (>=6523.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

