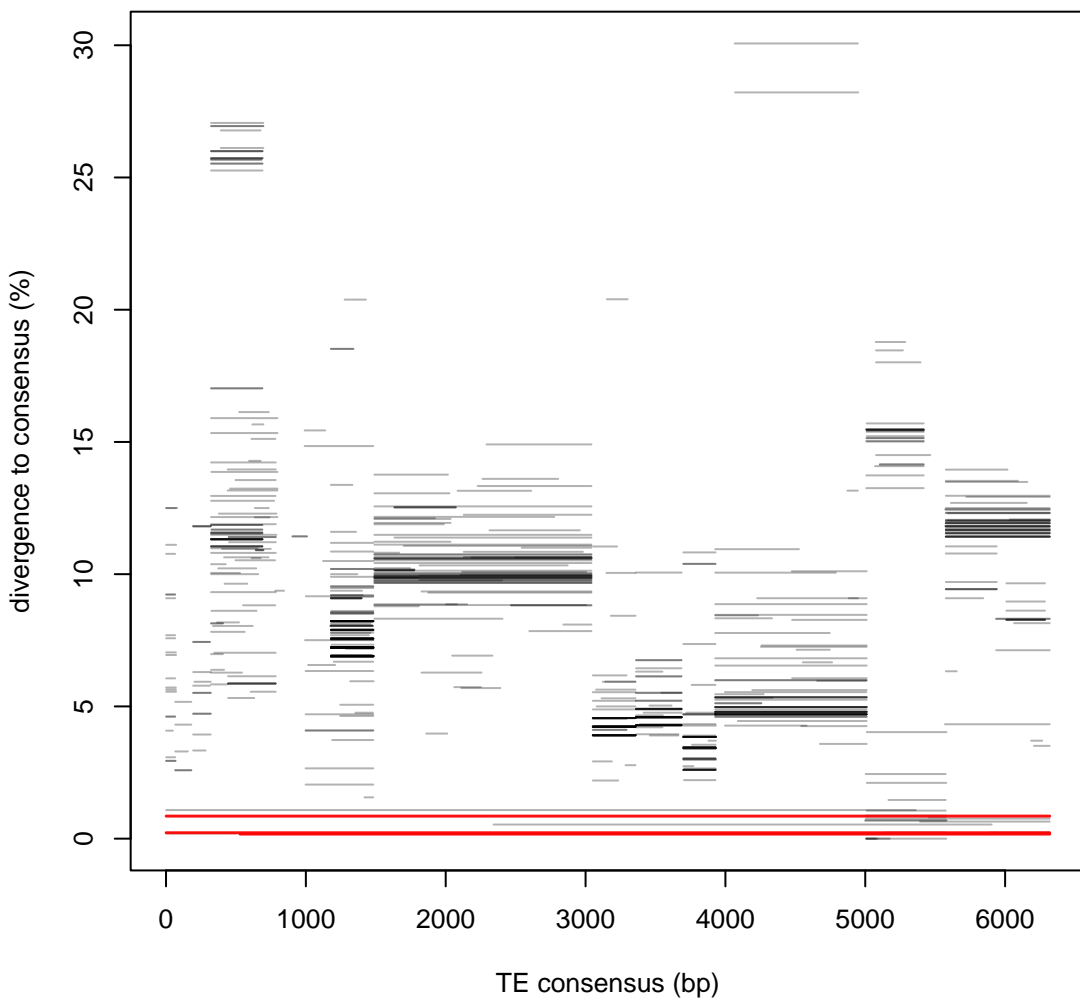
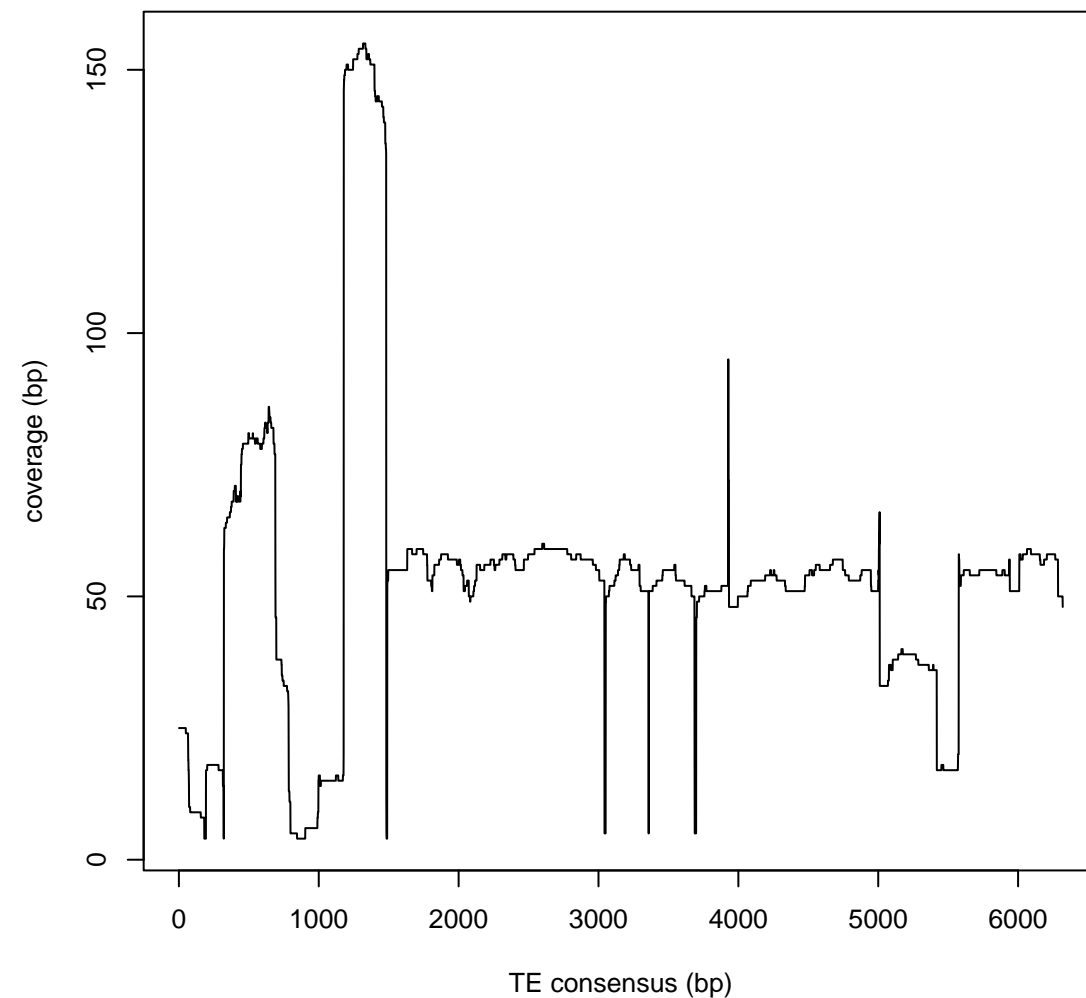


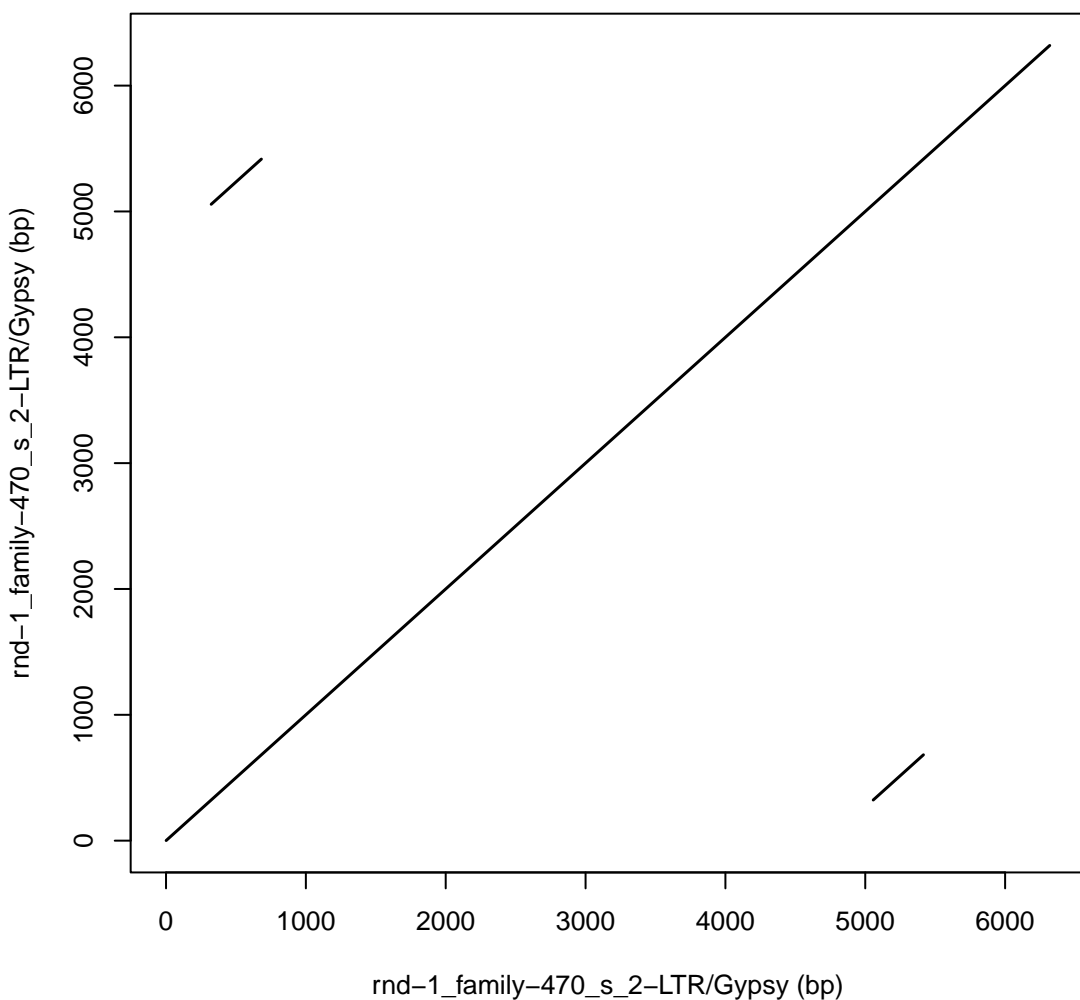
TE: rnd-1_family-470_s_2-LTR/Gypsy
consensus size: 6319bp; fragments: 702; full length: 3 (>=5687.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

