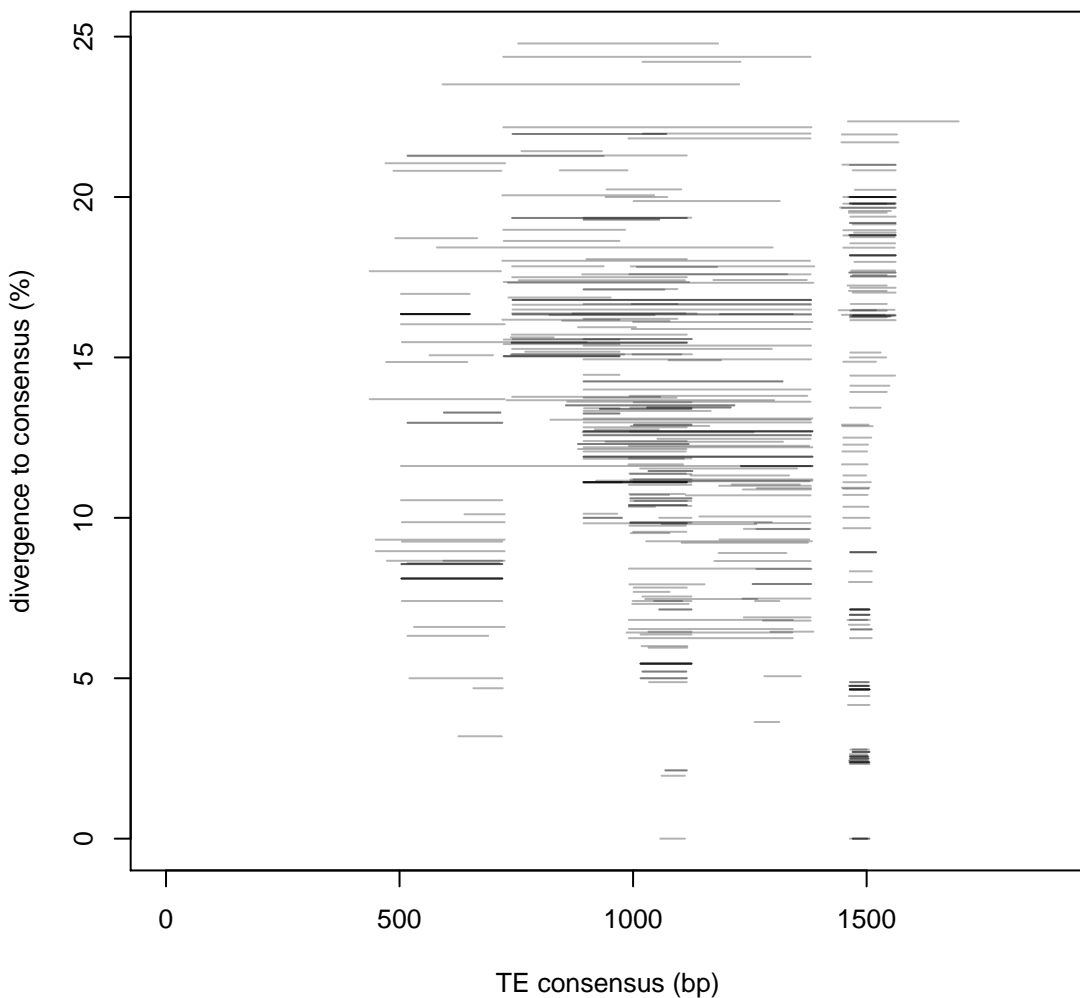
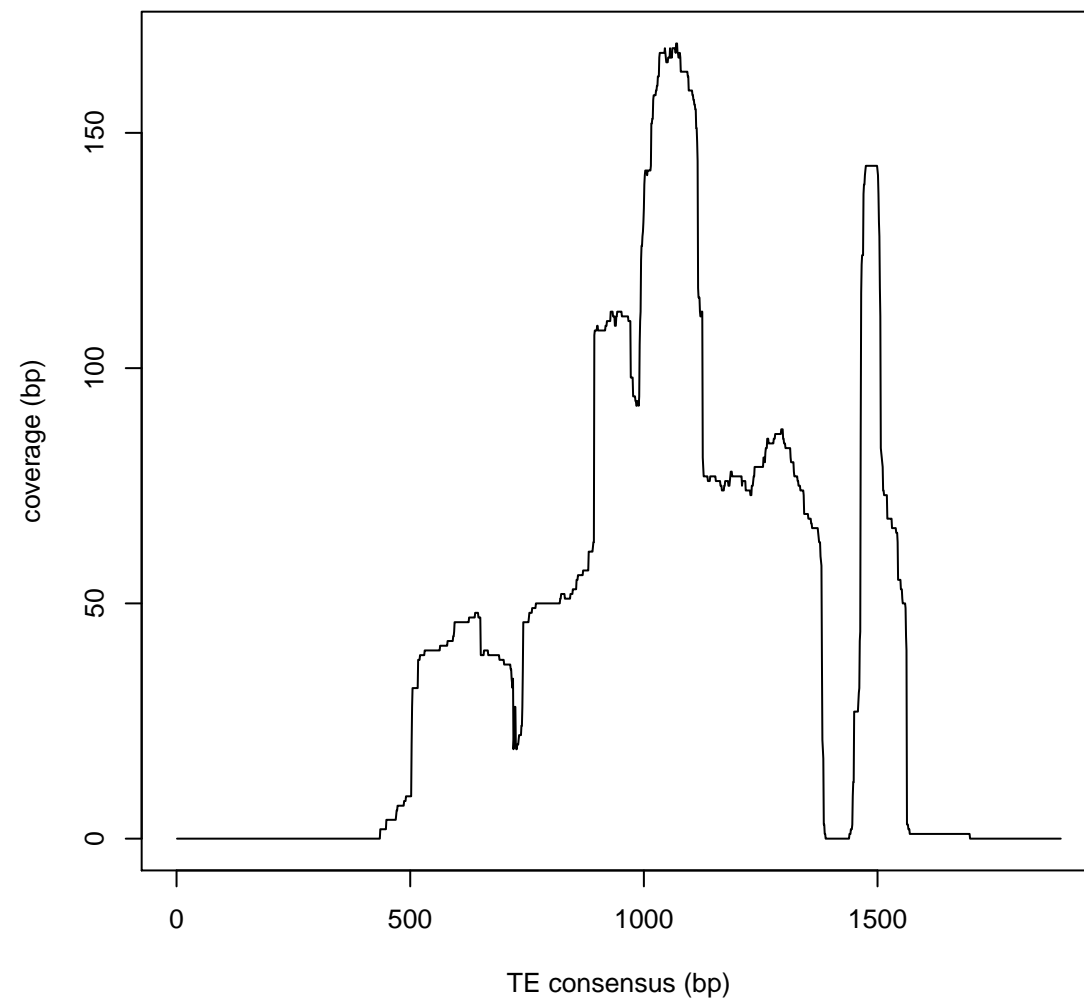


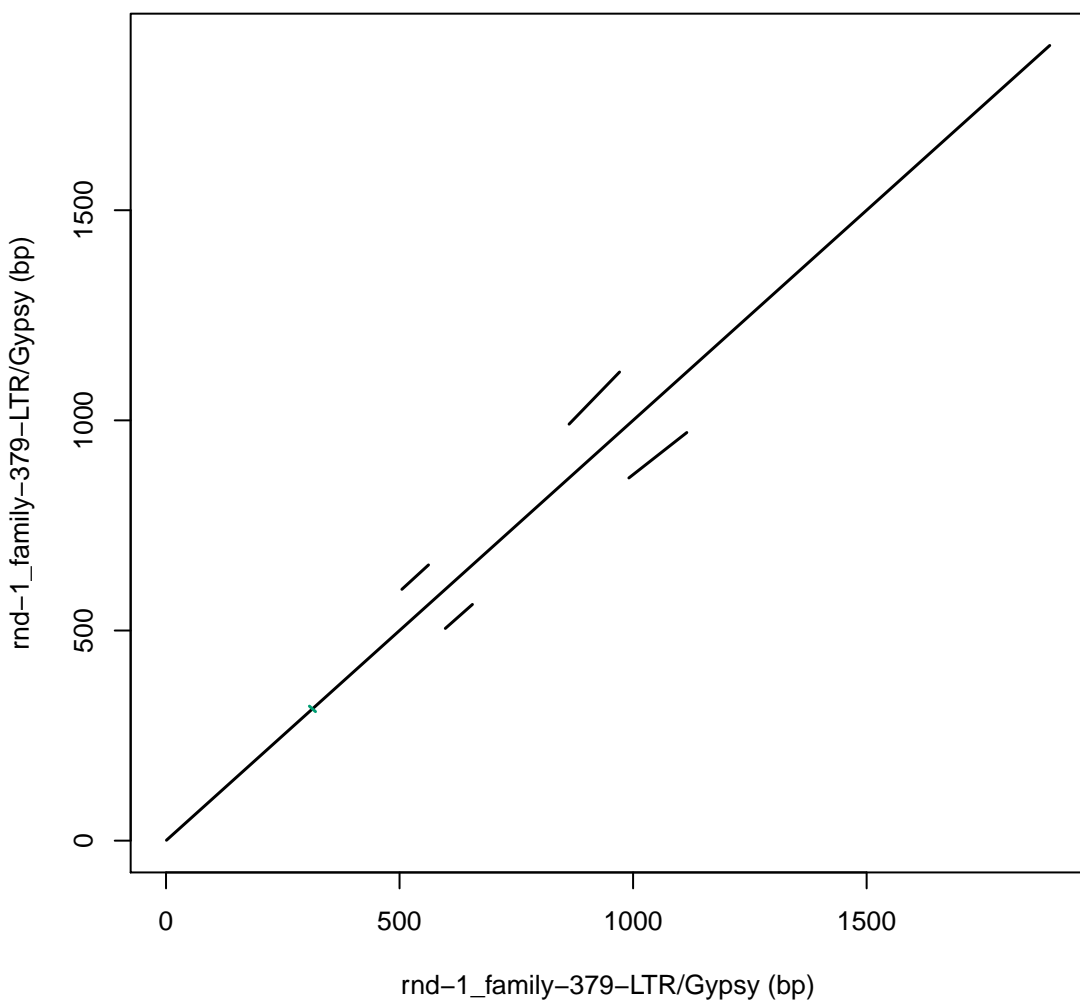
TE: rnd-1_family-379-LTR/Gypsy
consensus size: 1892bp; fragments: 420; full length: 0 (>=1702.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

