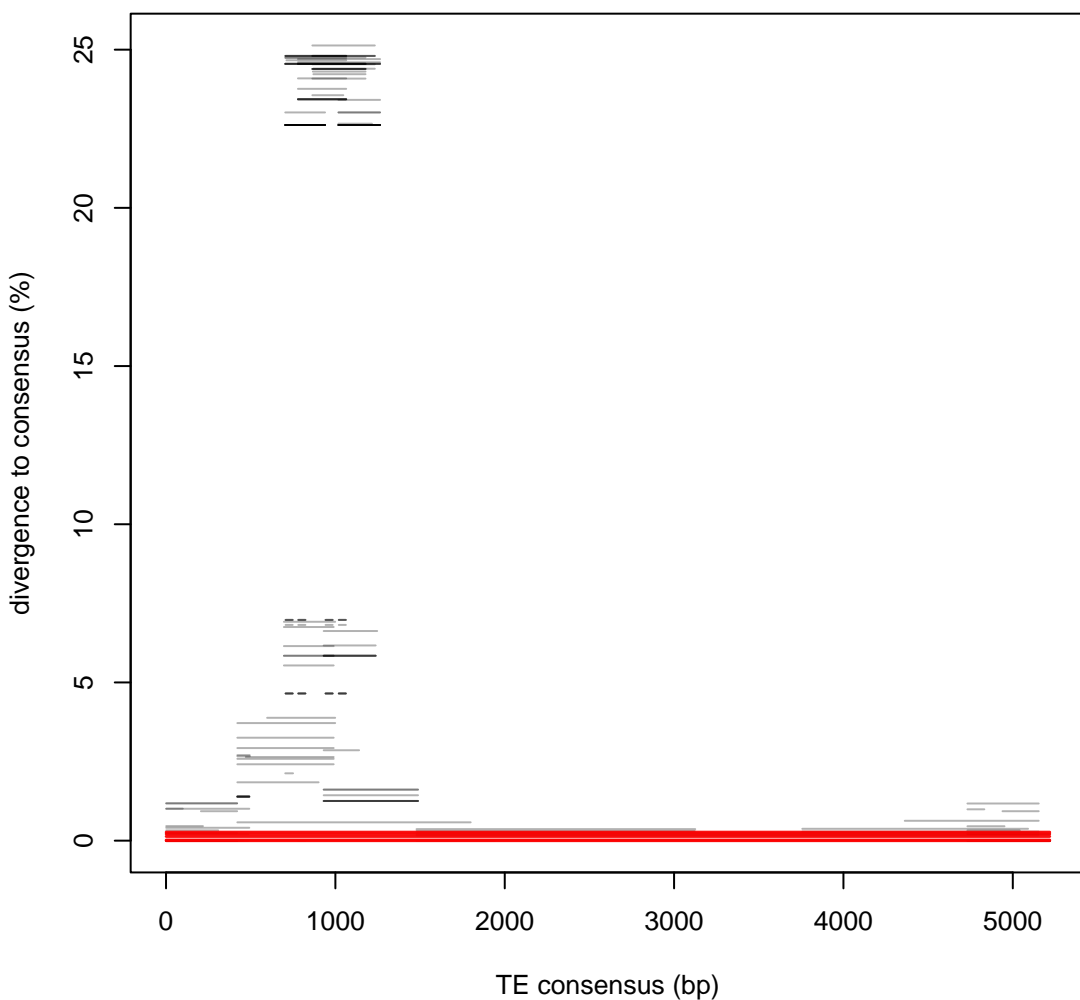
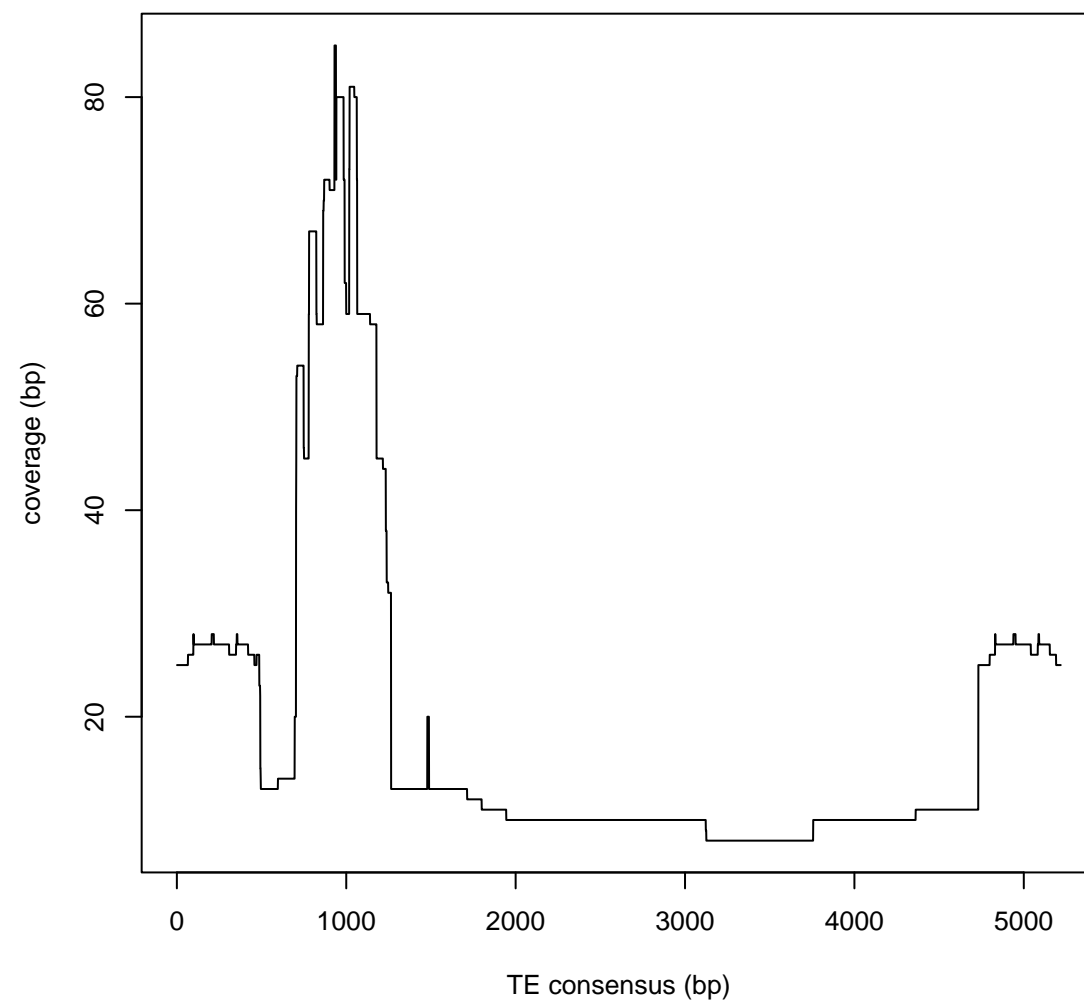


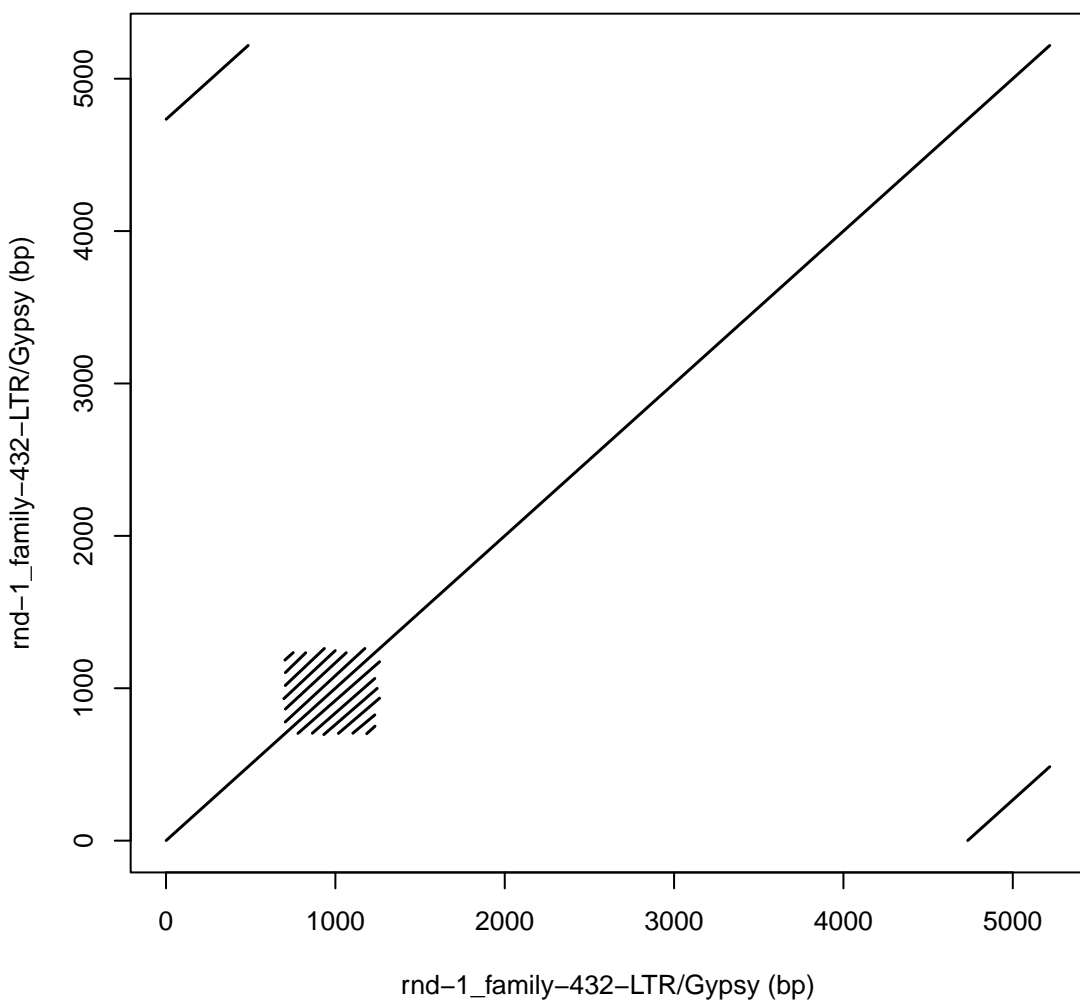
TE: rnd-1_family-432-LTR/Gypsy
consensus size: 5218bp; fragments: 219; full length: 5 (≥ 4696.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

