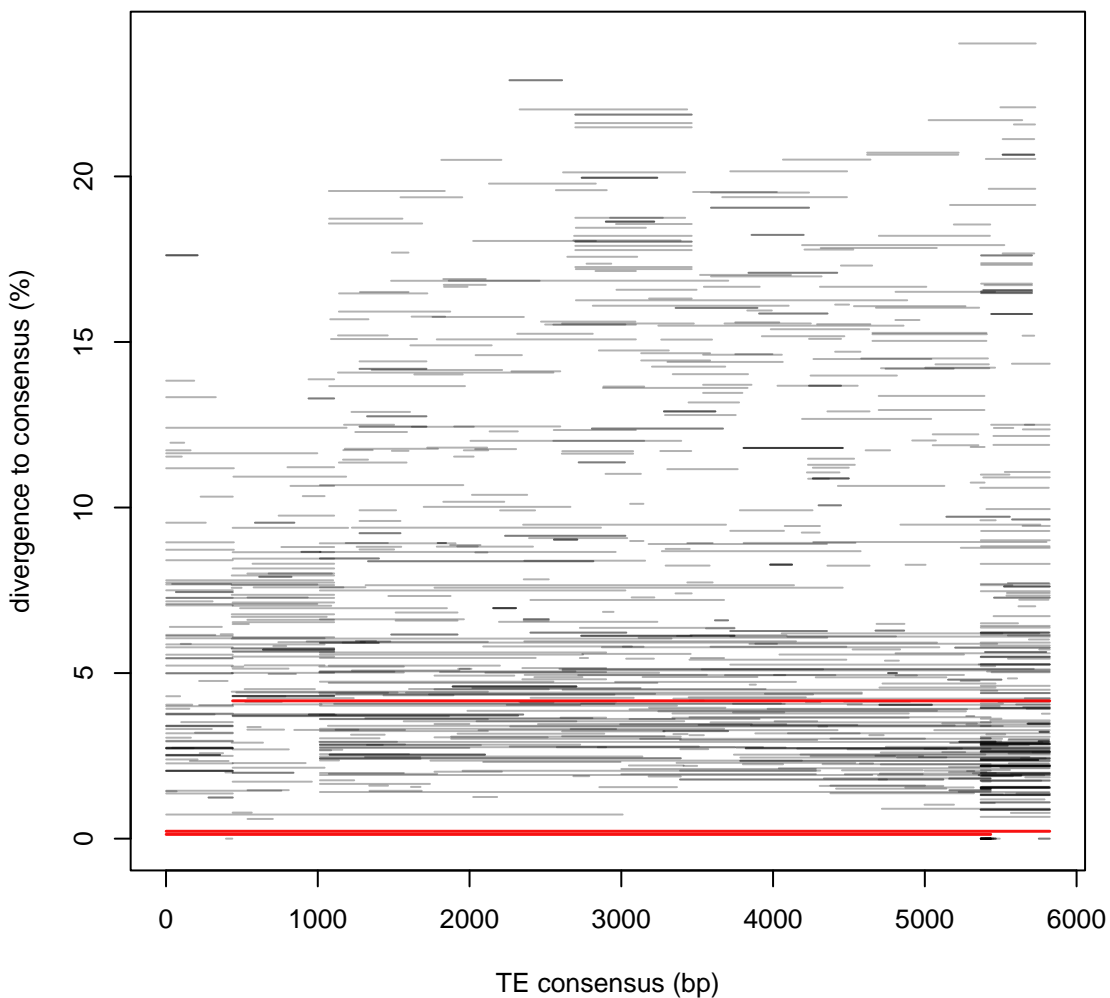
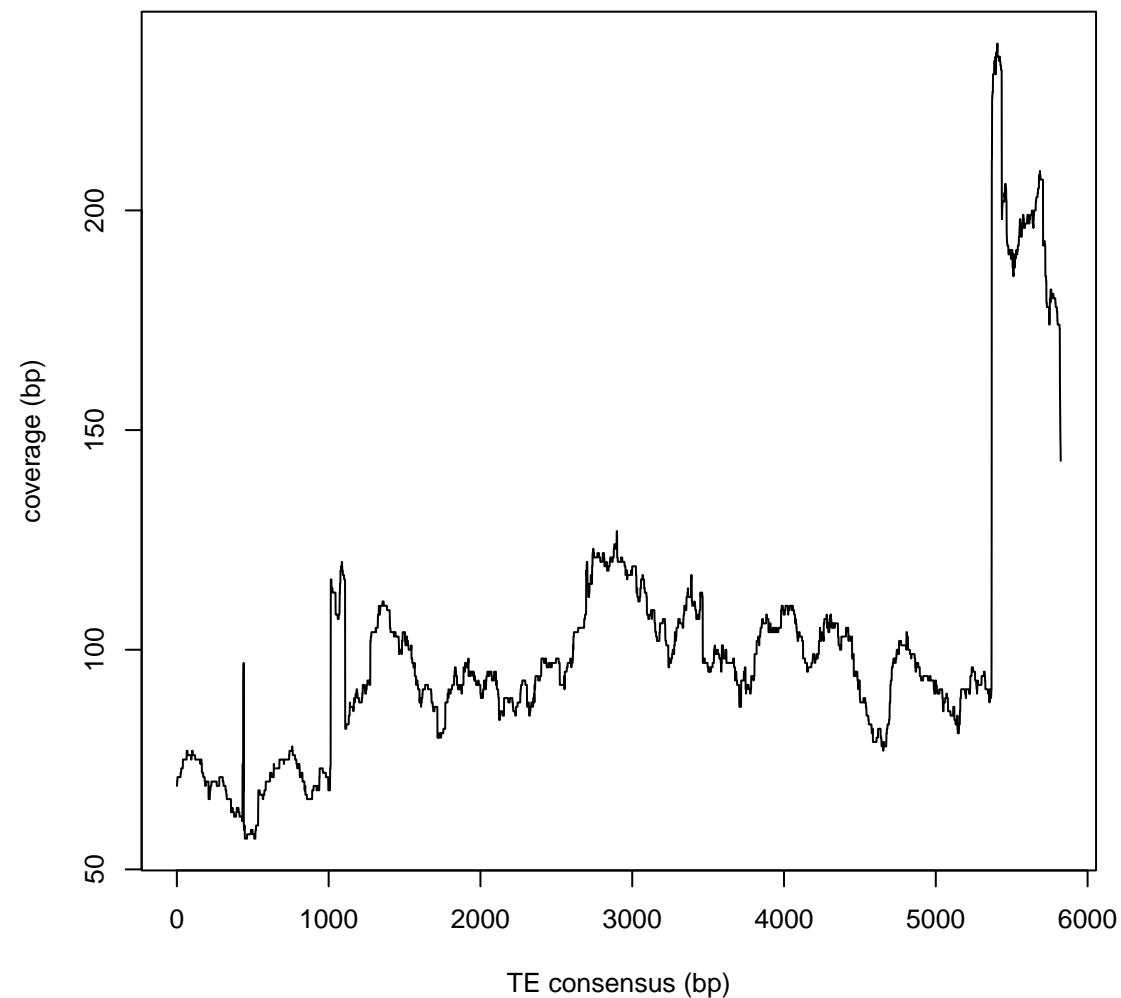


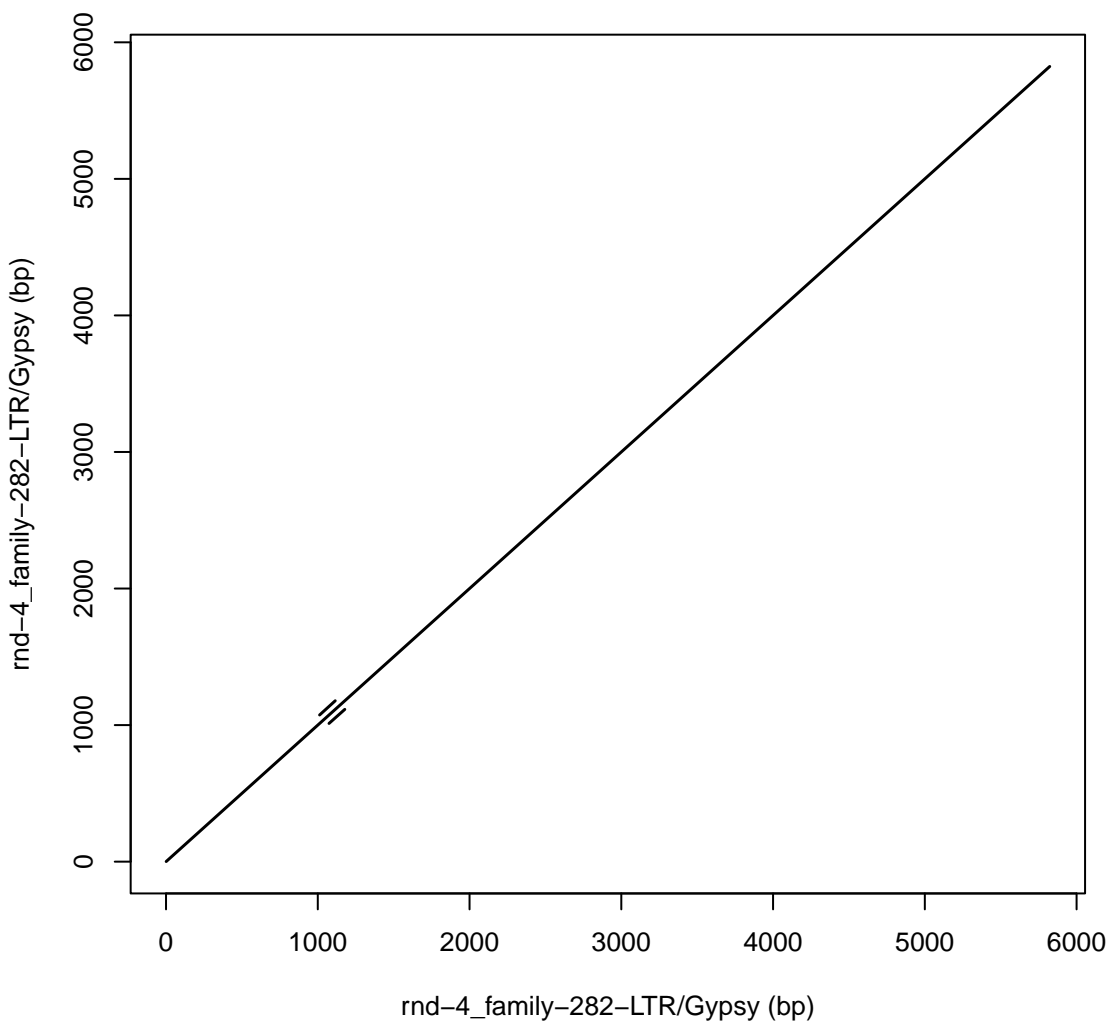
TE: rnd-4_family-282-LTR/Gypsy
consensus size: 5823bp; fragments: 956; full length: 3 (>=5240.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

