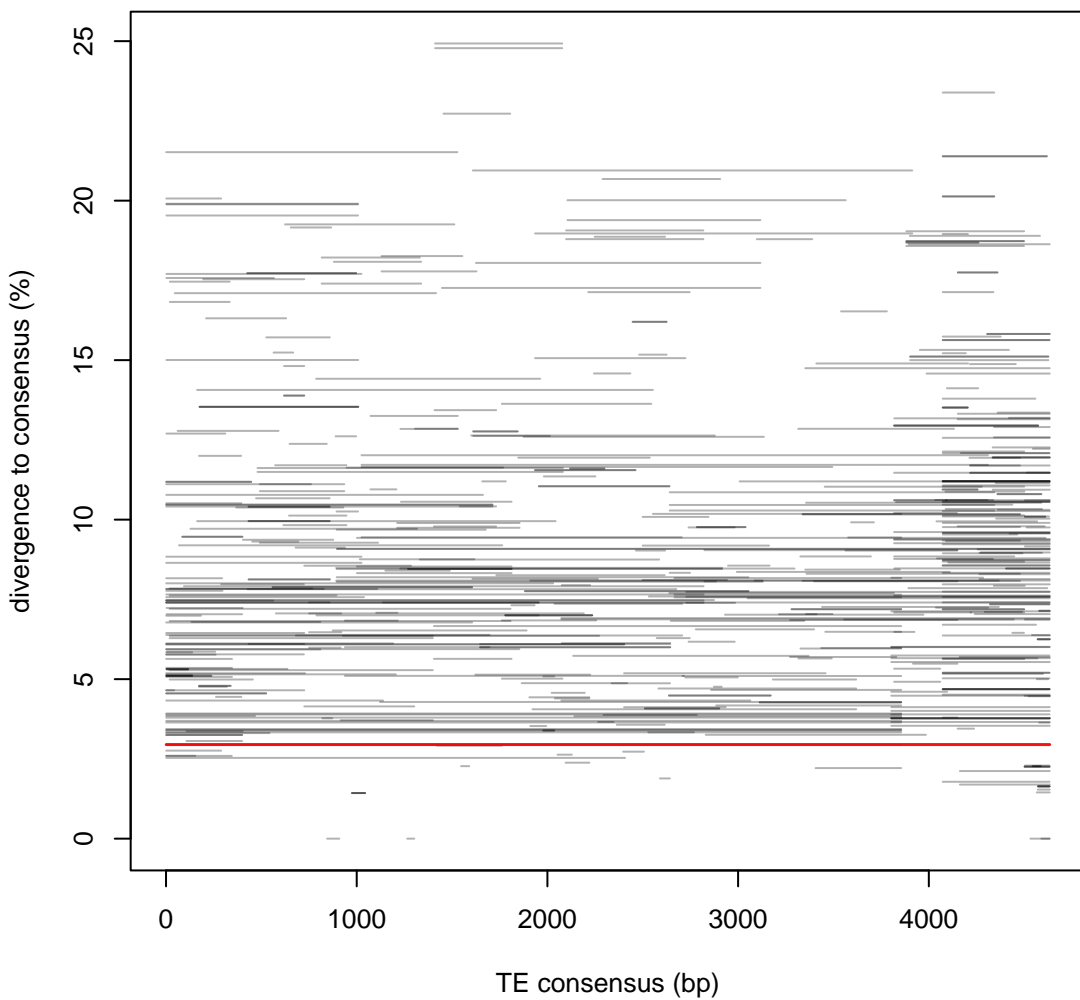
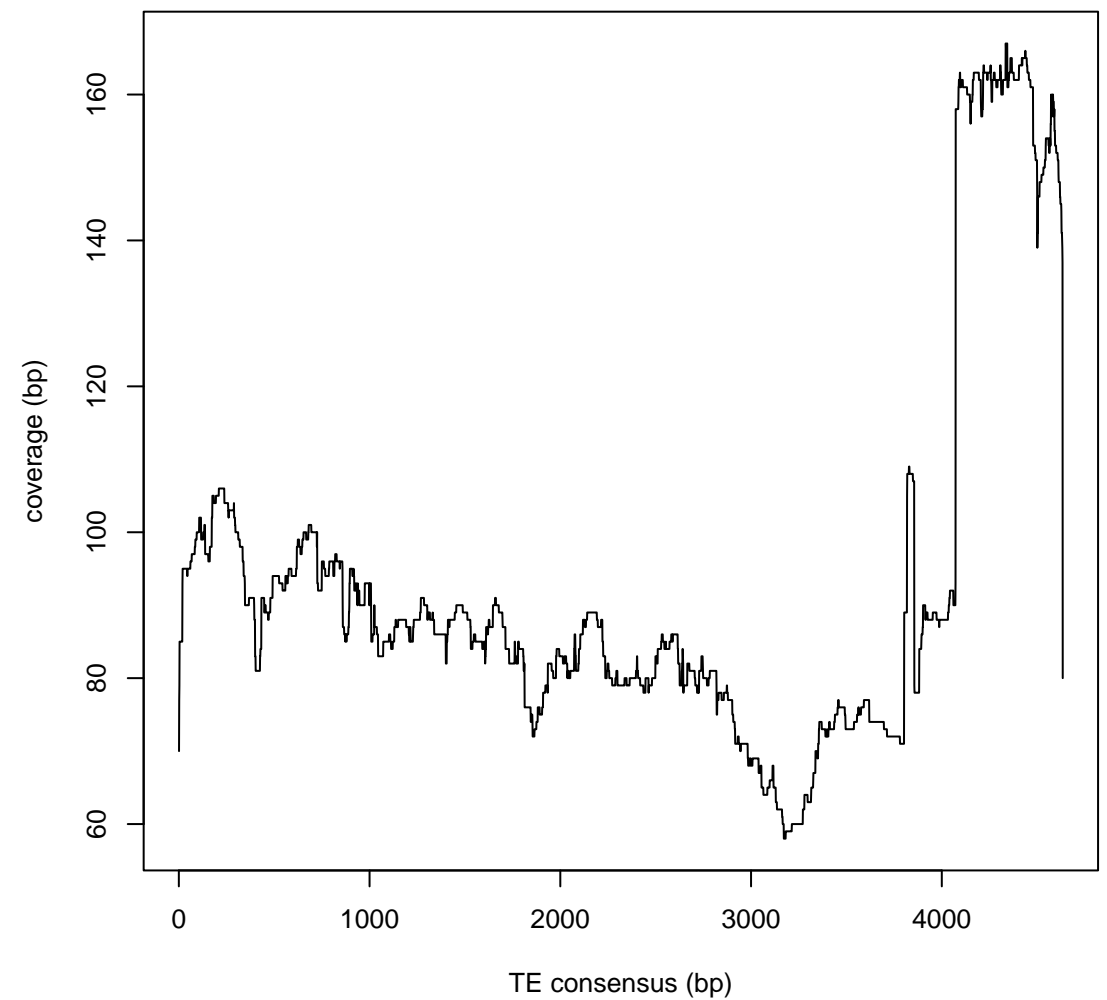


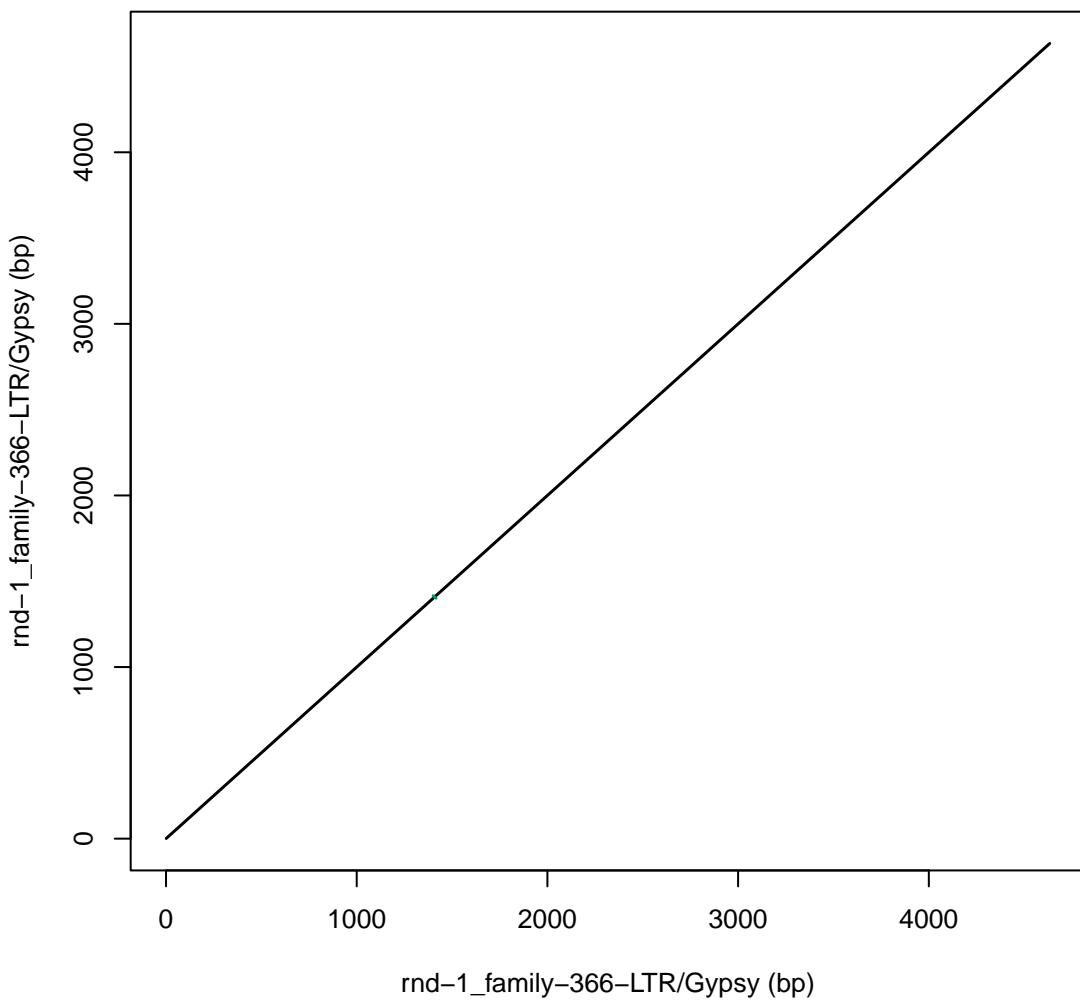
TE: rnd-1_family-366-LTR/Gypsy
consensus size: 4634bp; fragments: 583; full length: 1 (≥ 4170.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

