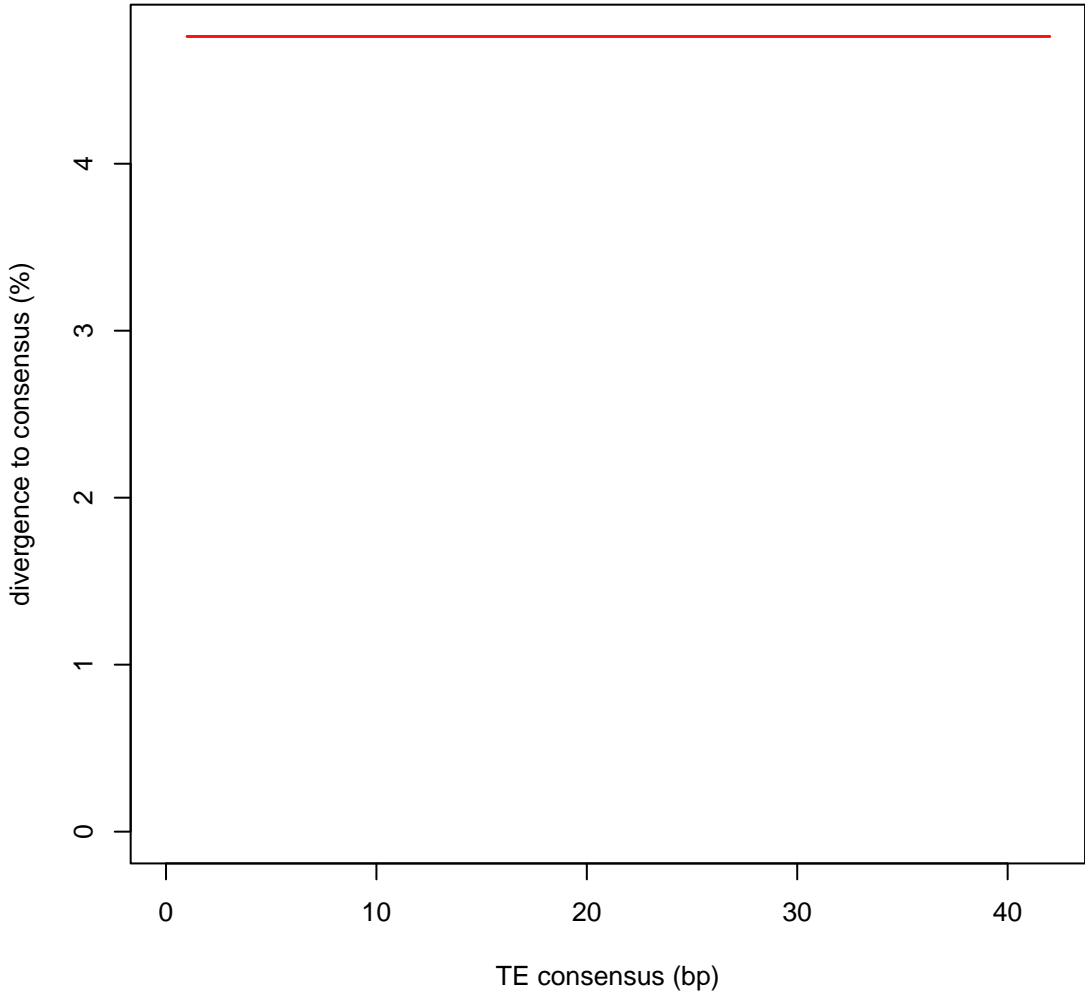
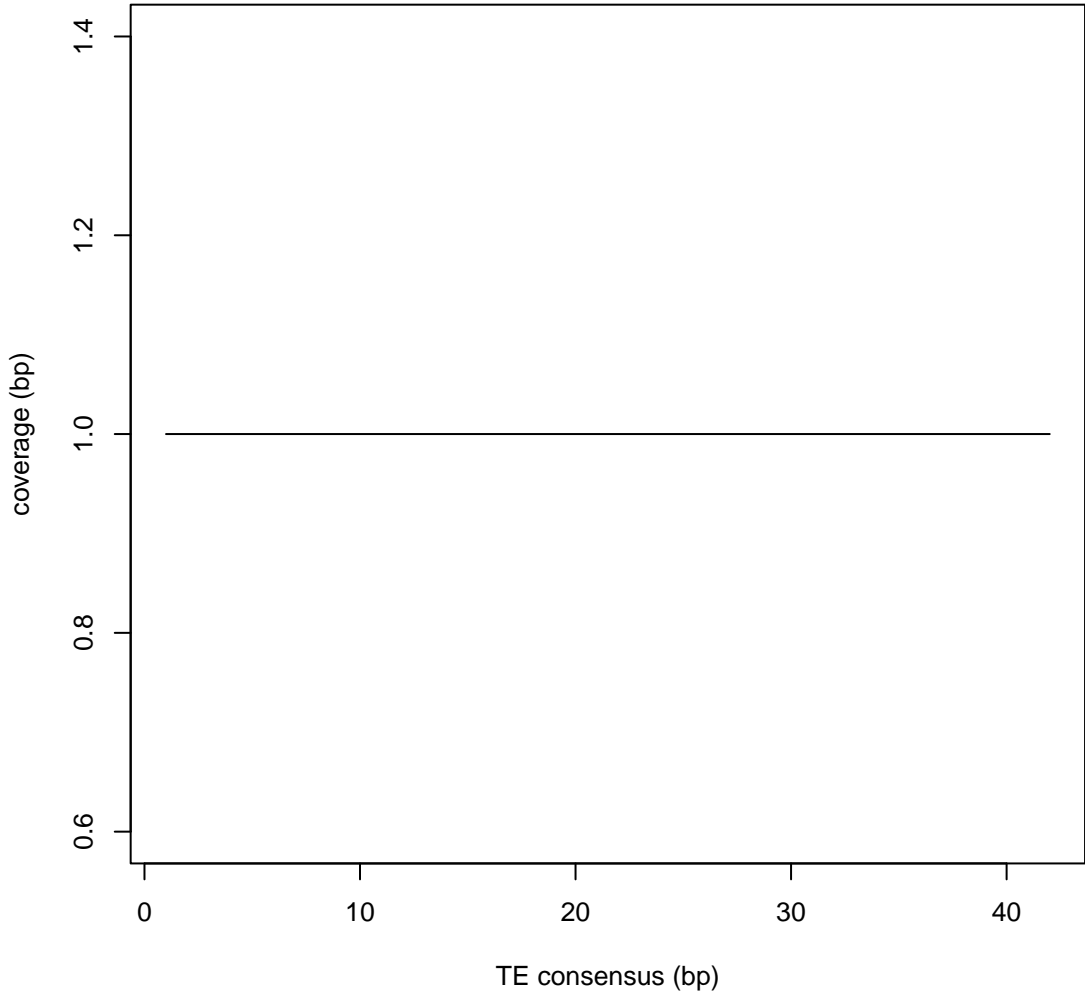


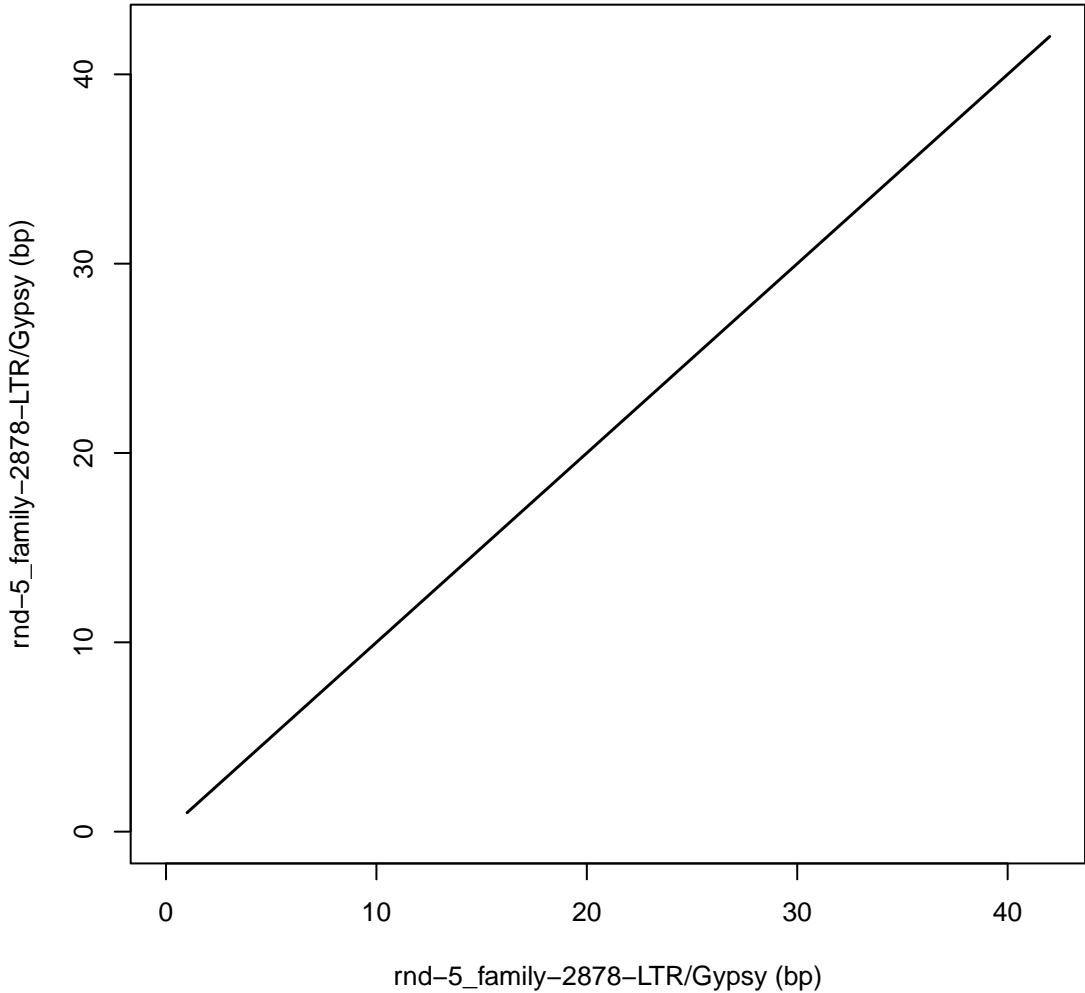
TE: rnd-5_family-2878-LTR/Gypsy
consensus size: 42bp; fragments: 1; full length: 1 (>=37.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

