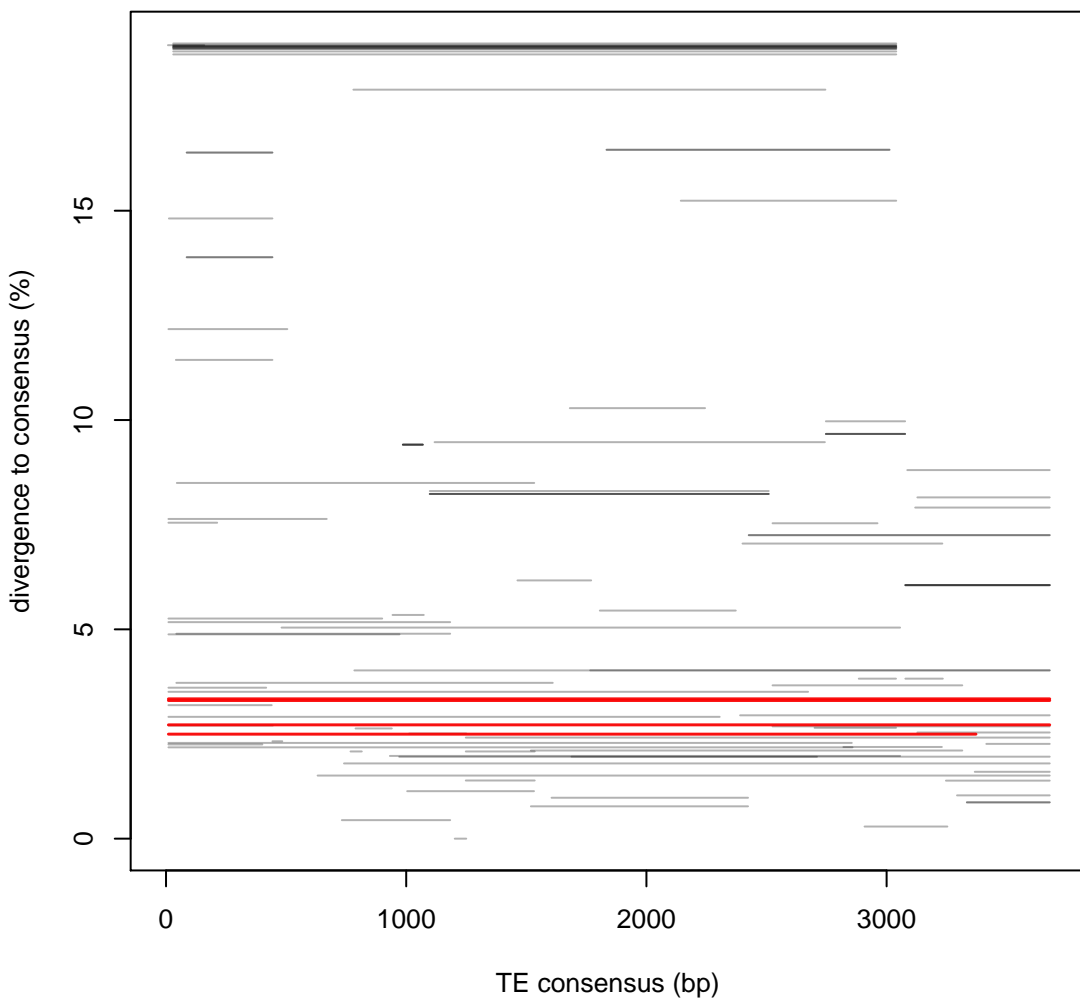
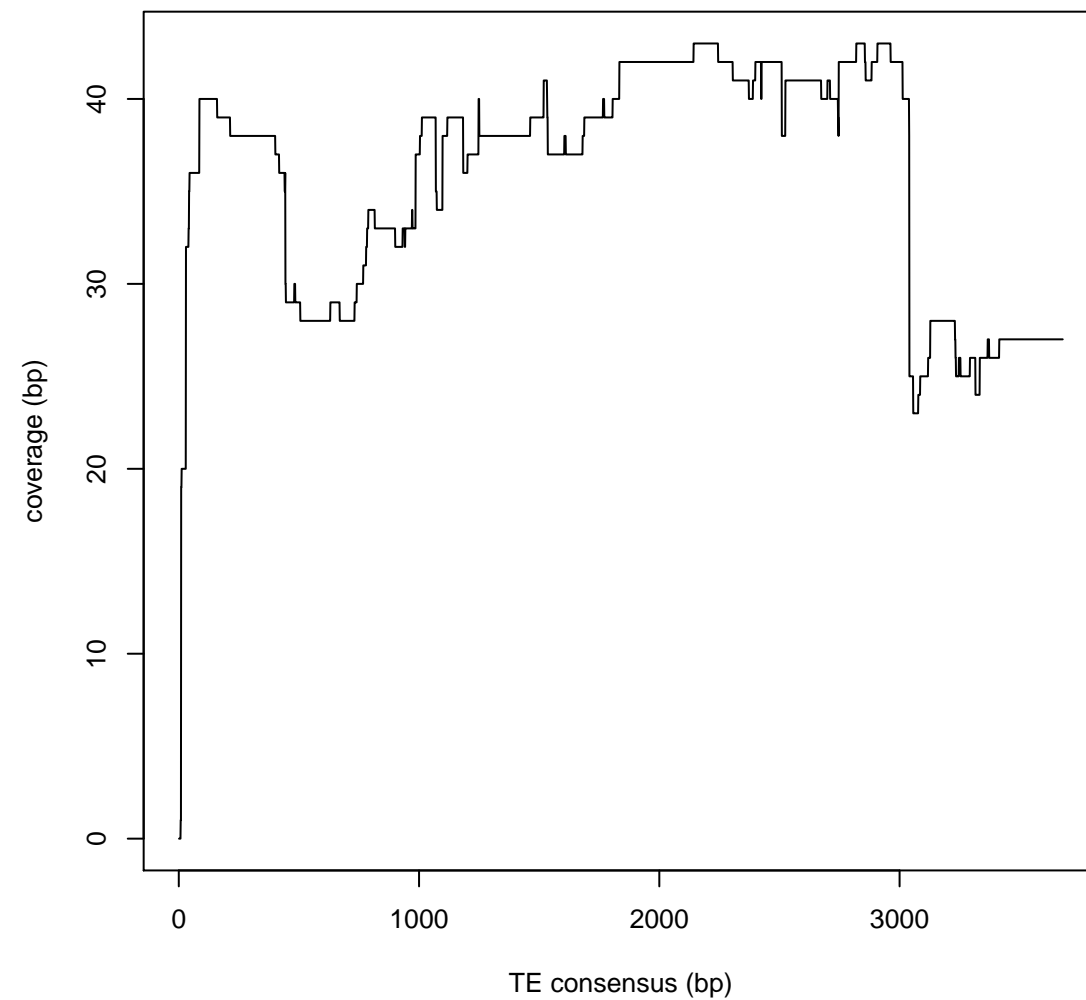


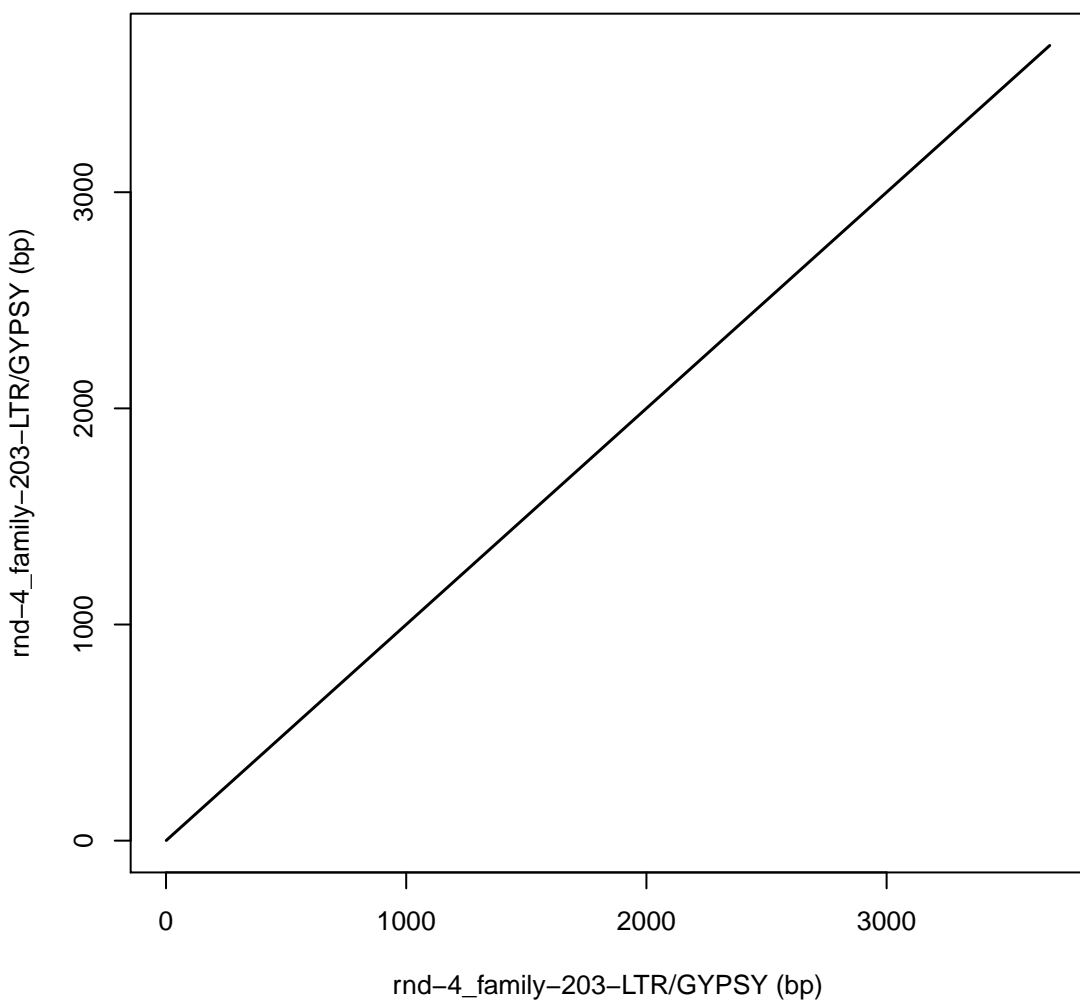
TE: rnd-4\_family-203-LTR/GYPSY  
consensus size: 3679bp; fragments: 108; full length: 4 (>=3311.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

