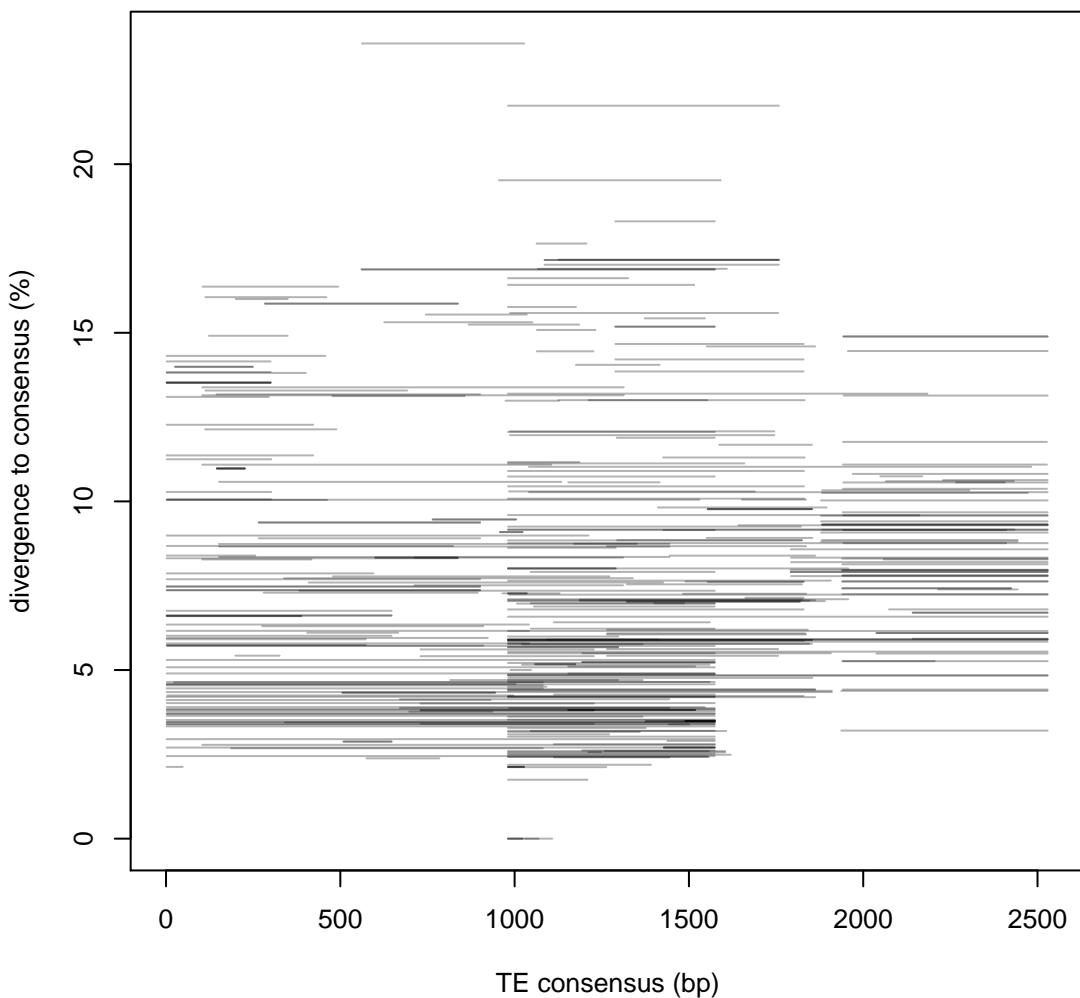
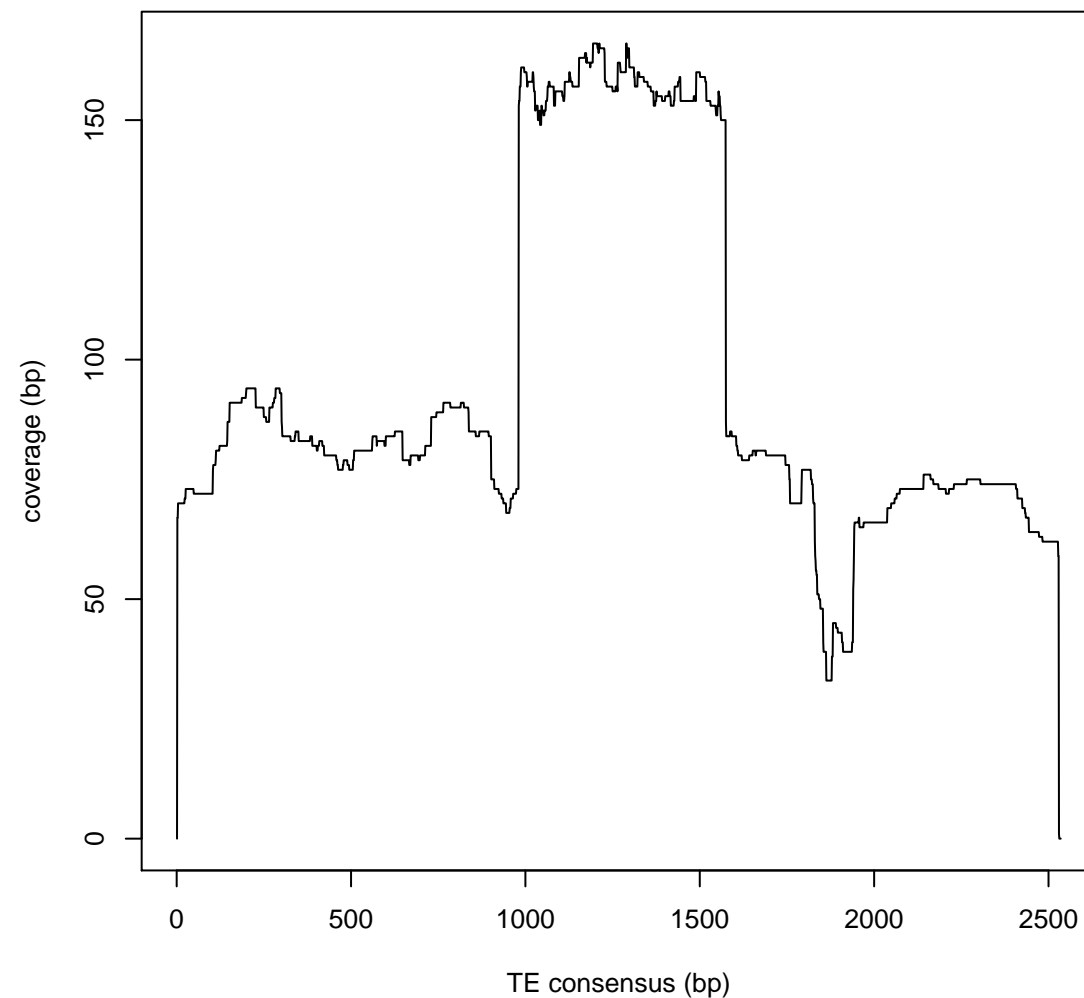


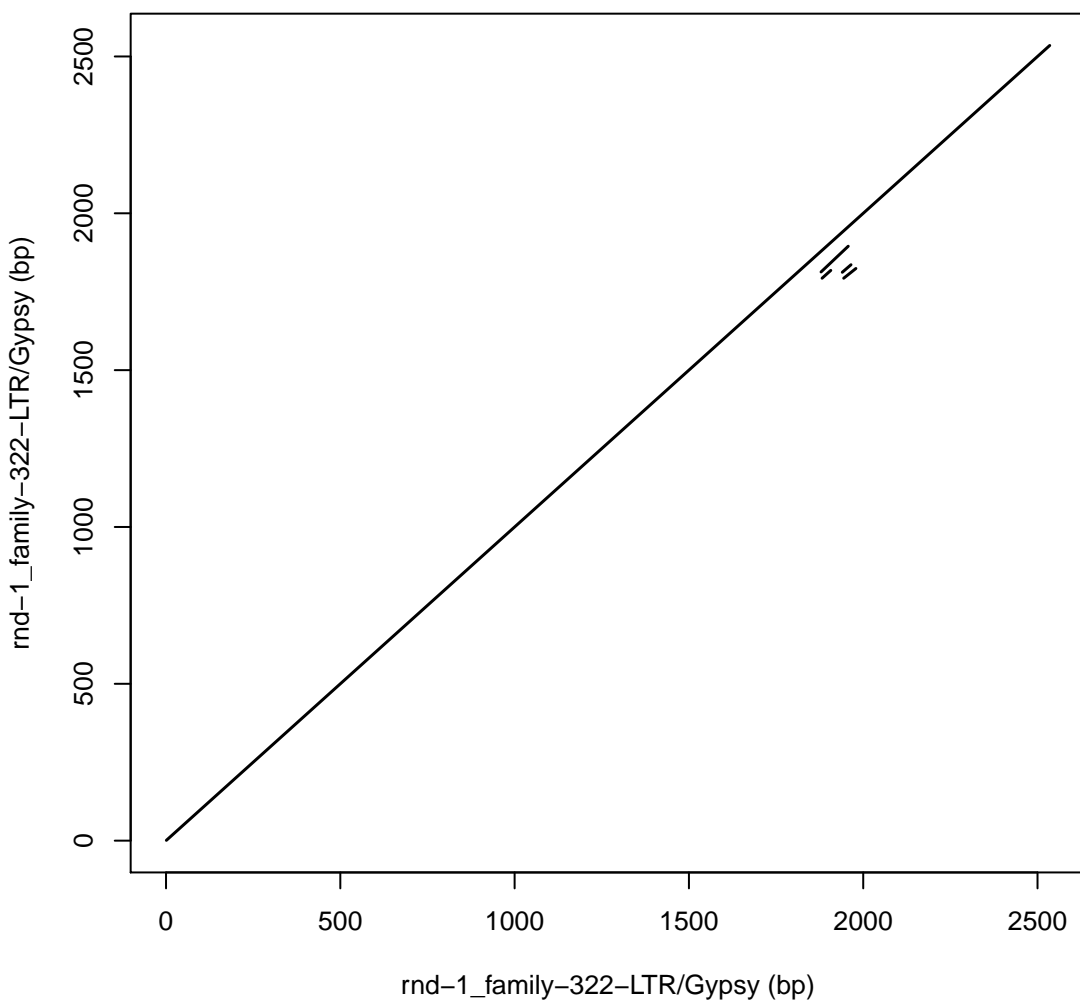
TE: rnd-1\_family-322-LTR/Gypsy  
consensus size: 2535bp; fragments: 381; full length: 0 (>=2281.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

