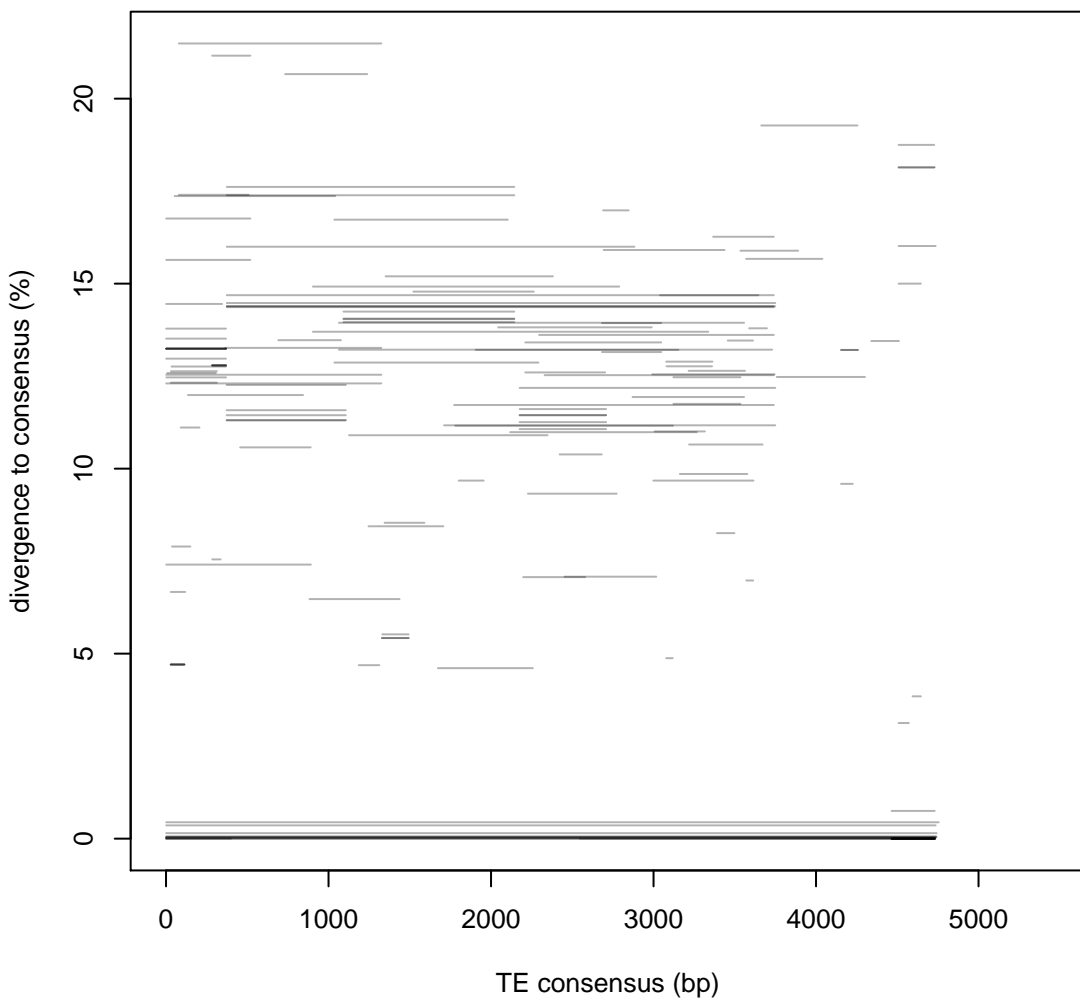
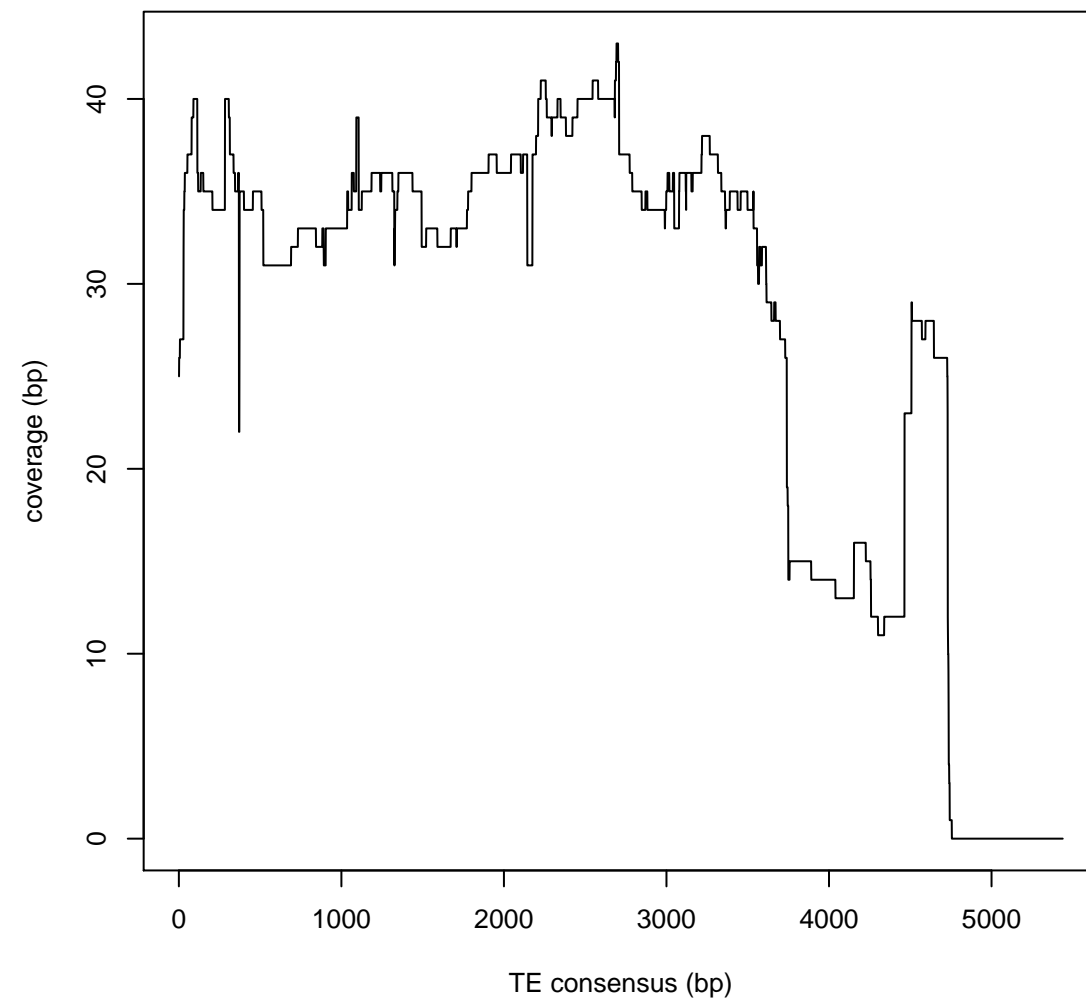


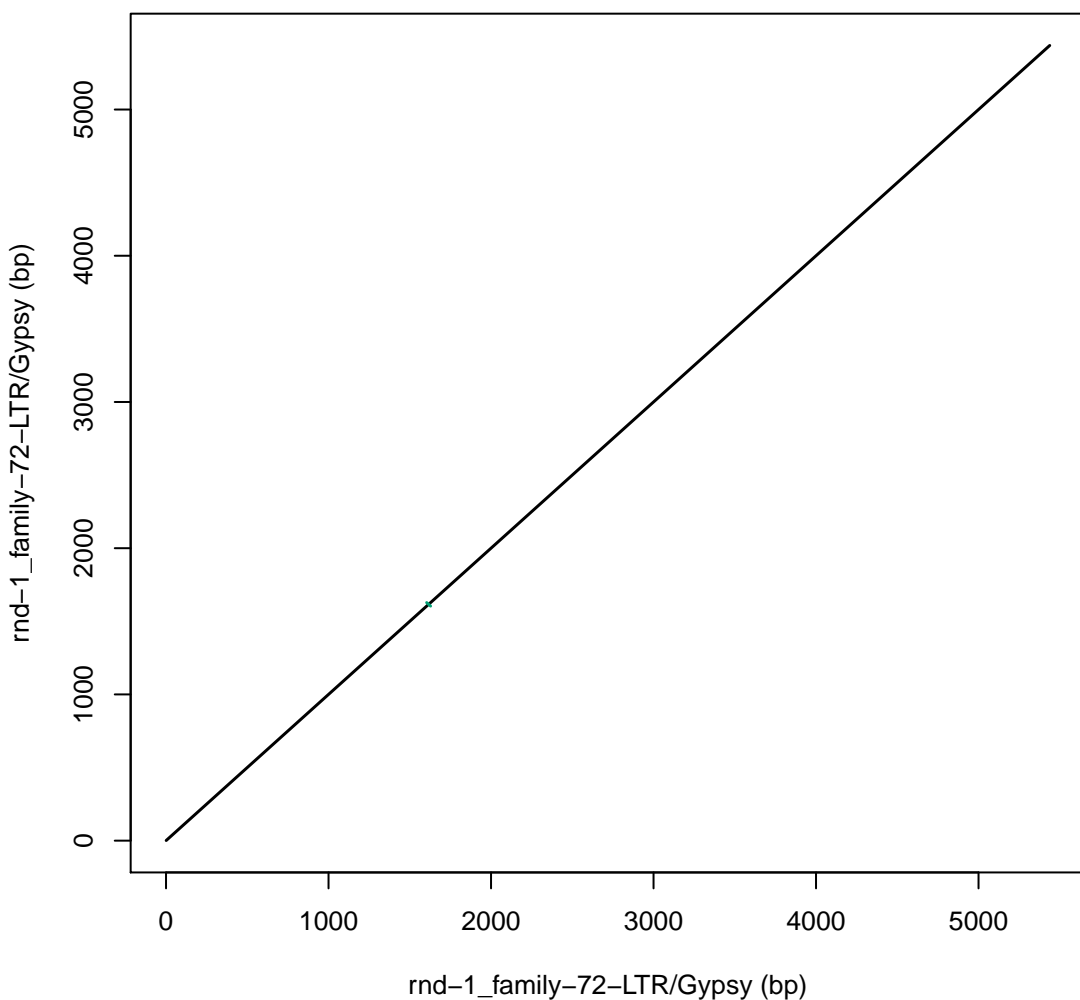
TE: rnd-1\_family-72-LTR/Gypsy  
consensus size: 5438bp; fragments: 155; full length: 0 (>=4894.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

