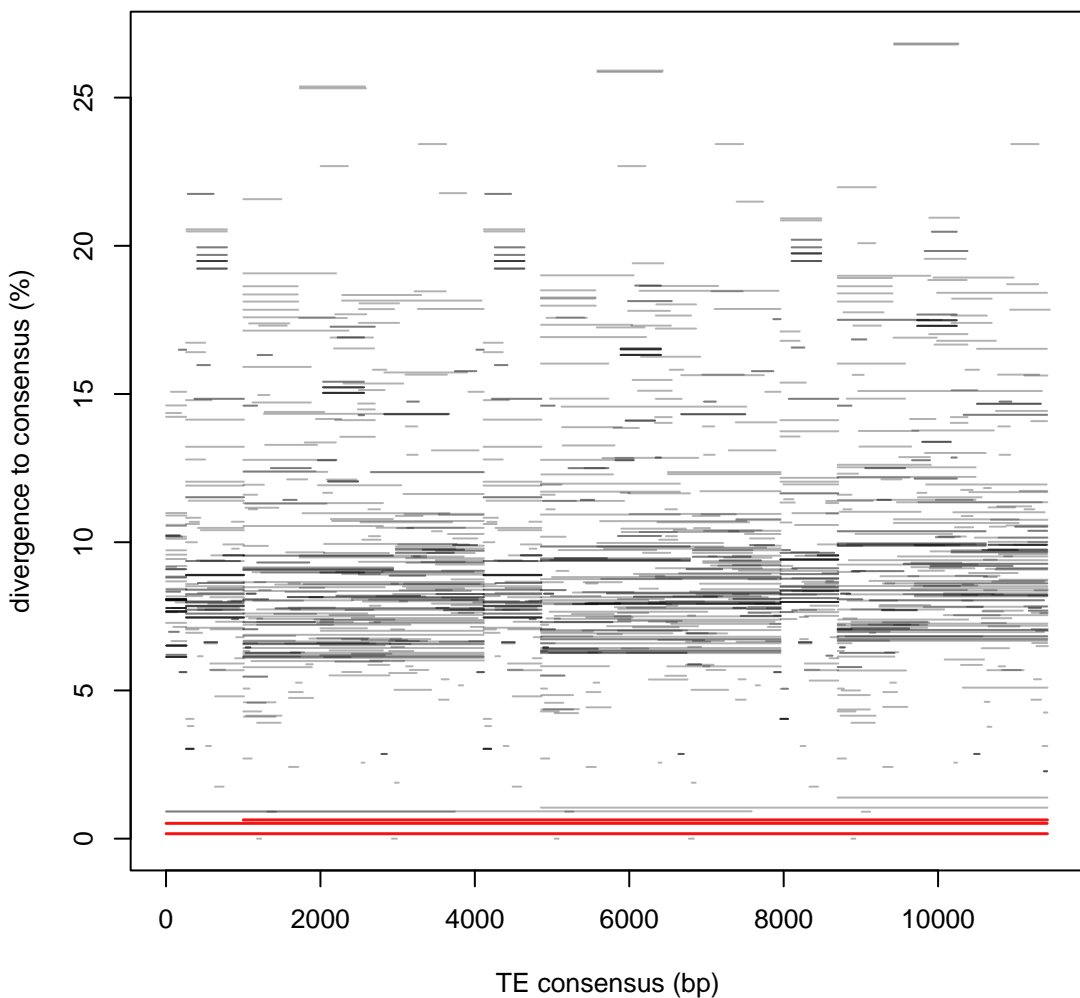
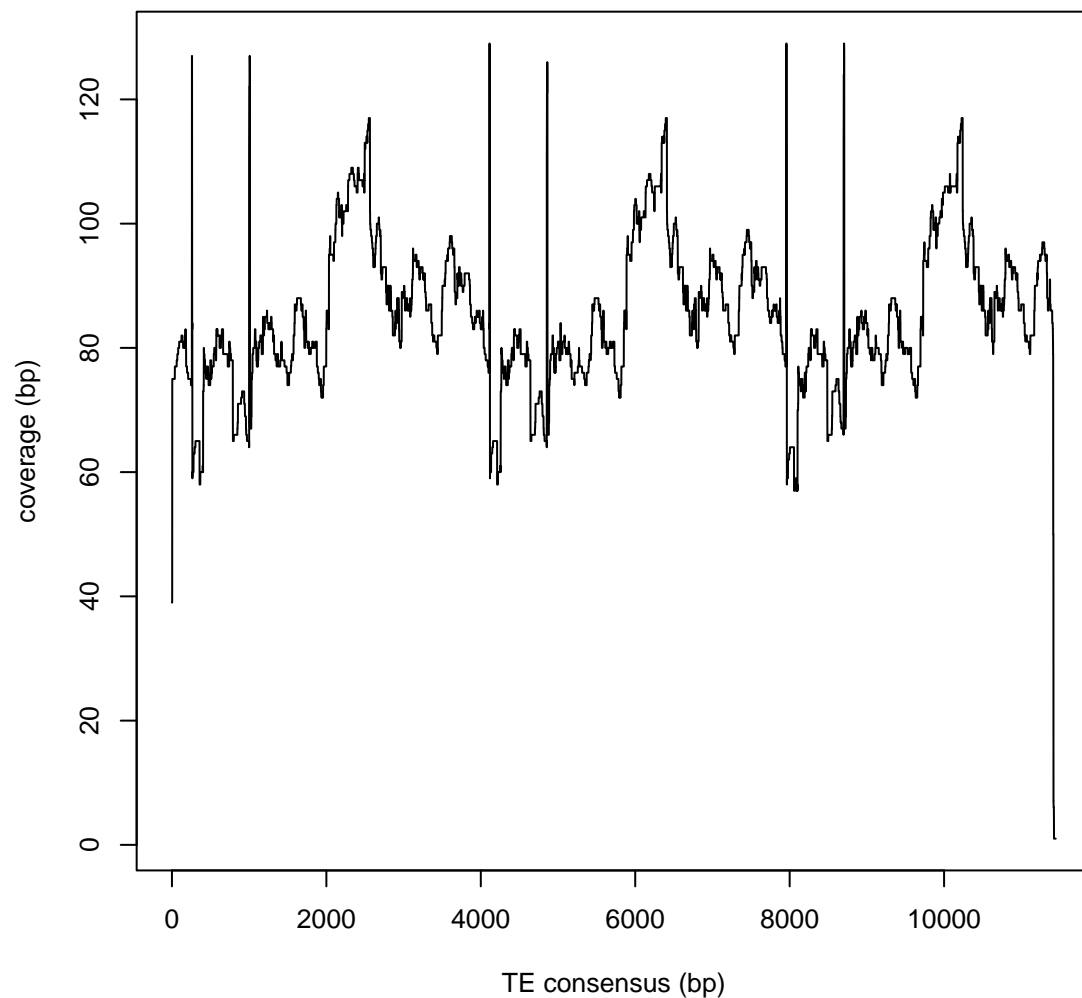


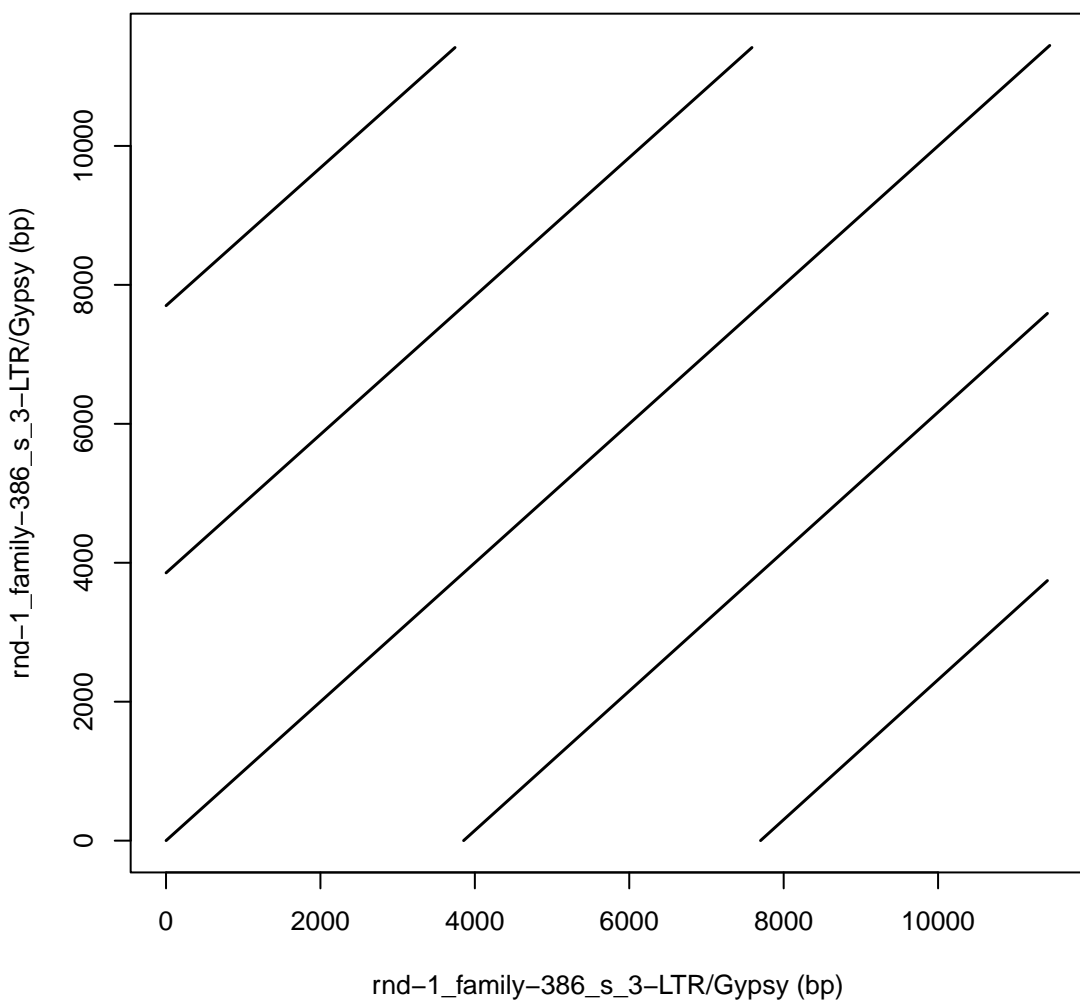
TE: rnd-1\_family-386\_s\_3-LTR/Gypsy  
consensus size: 11446bp; fragments: 1423; full length: 3 (>=10301.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

