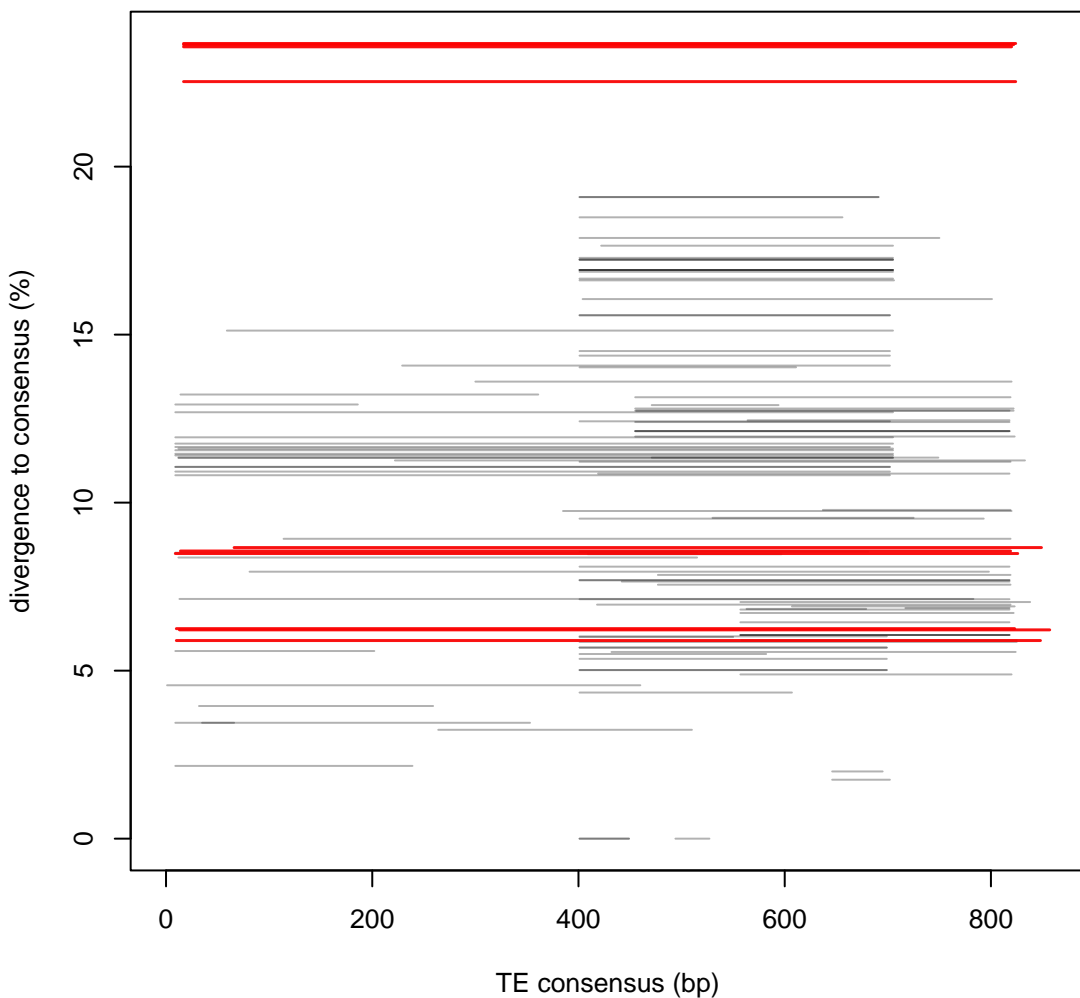
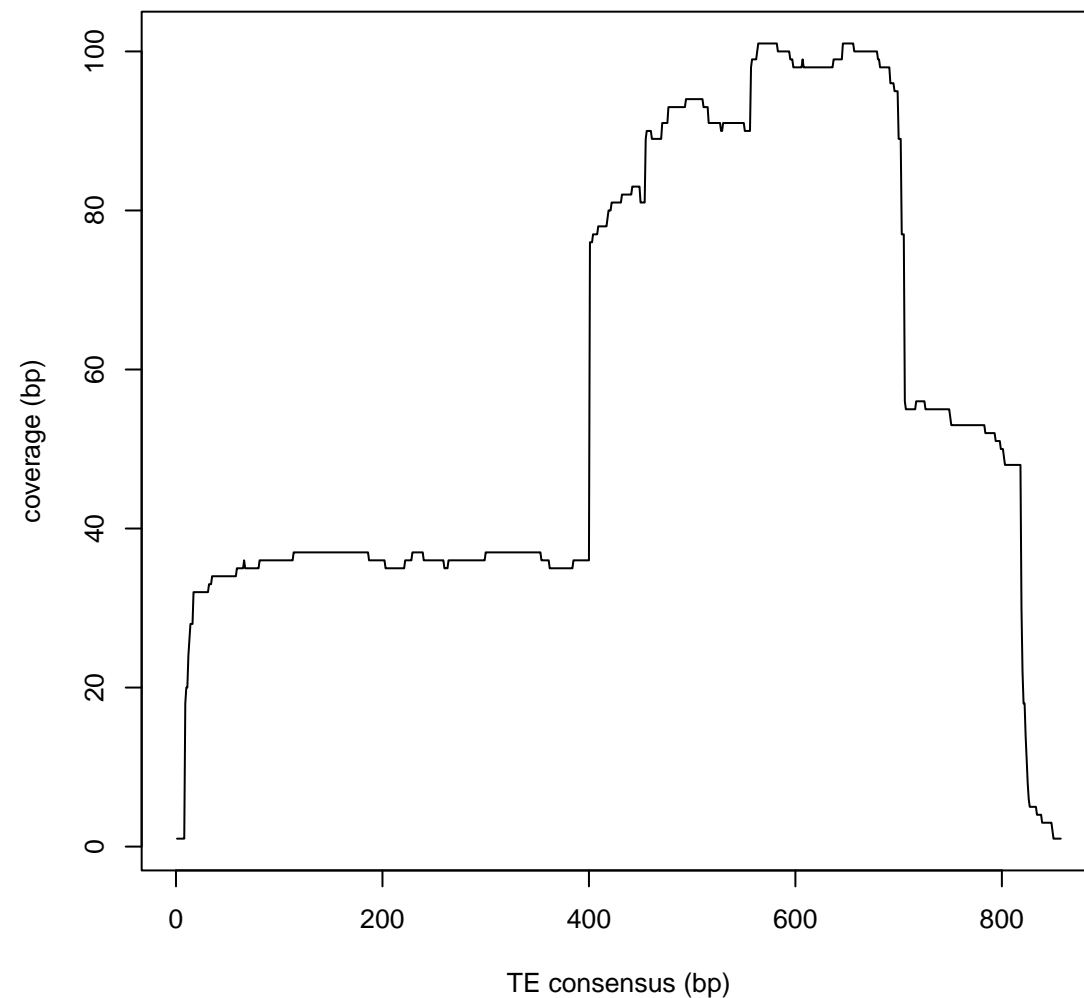


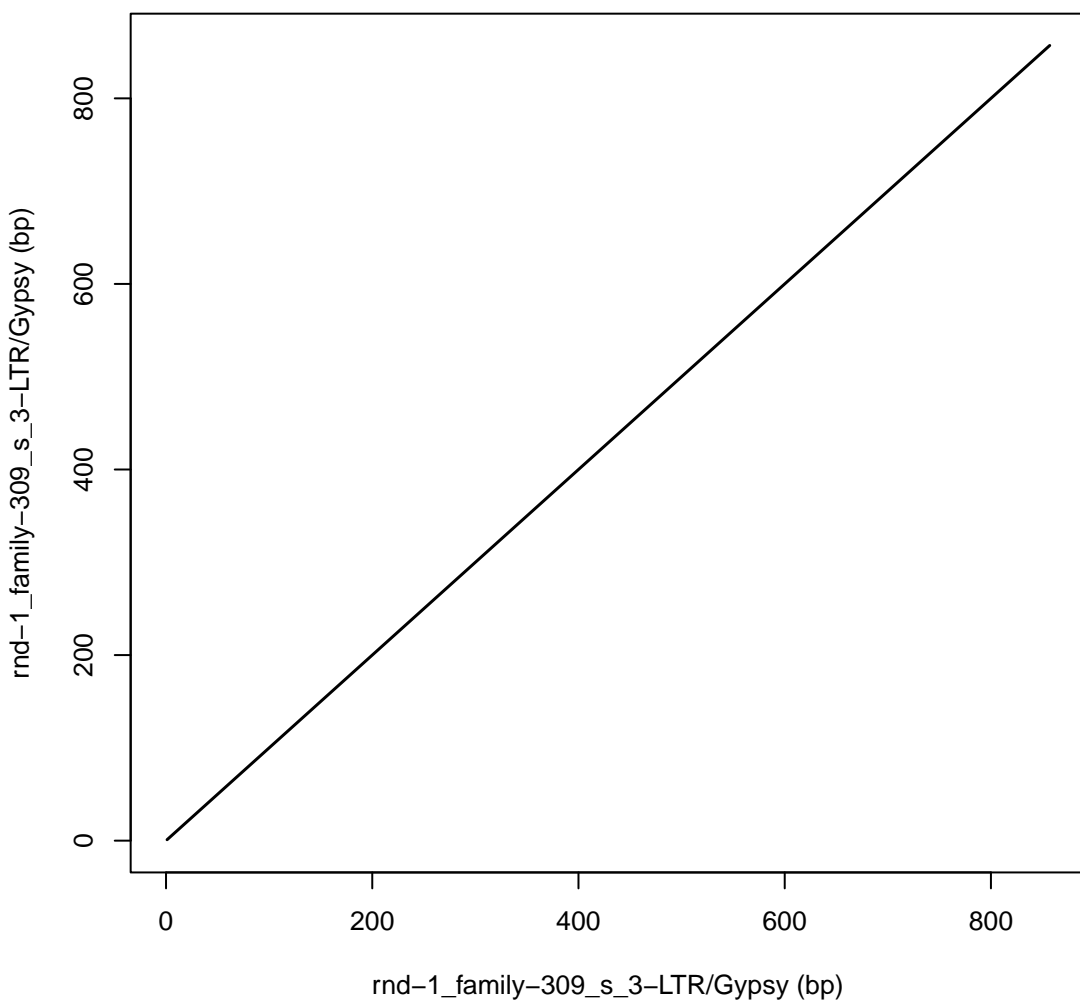
TE: rnd-1\_family-309\_s\_3-LTR/Gypsy  
consensus size: 857bp; fragments: 122; full length: 10 (>=771.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

