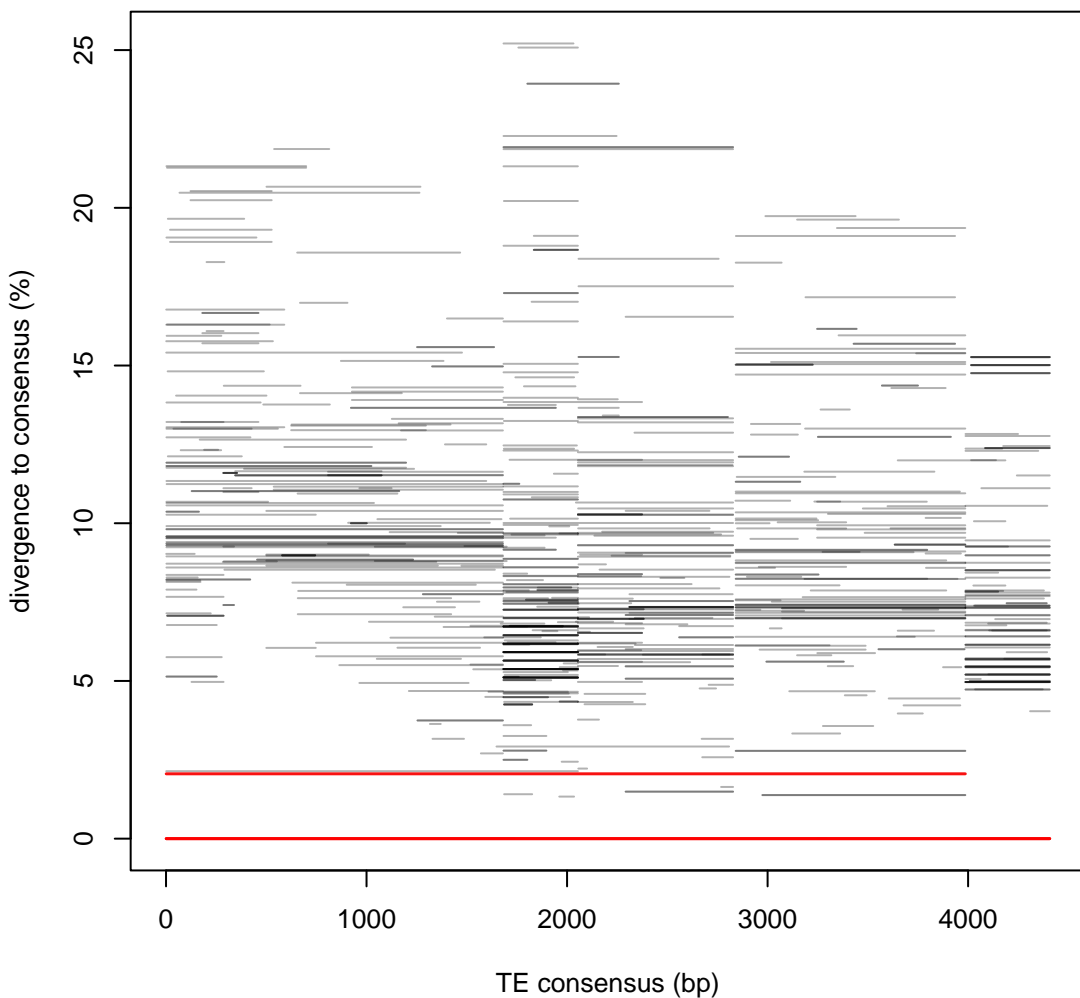
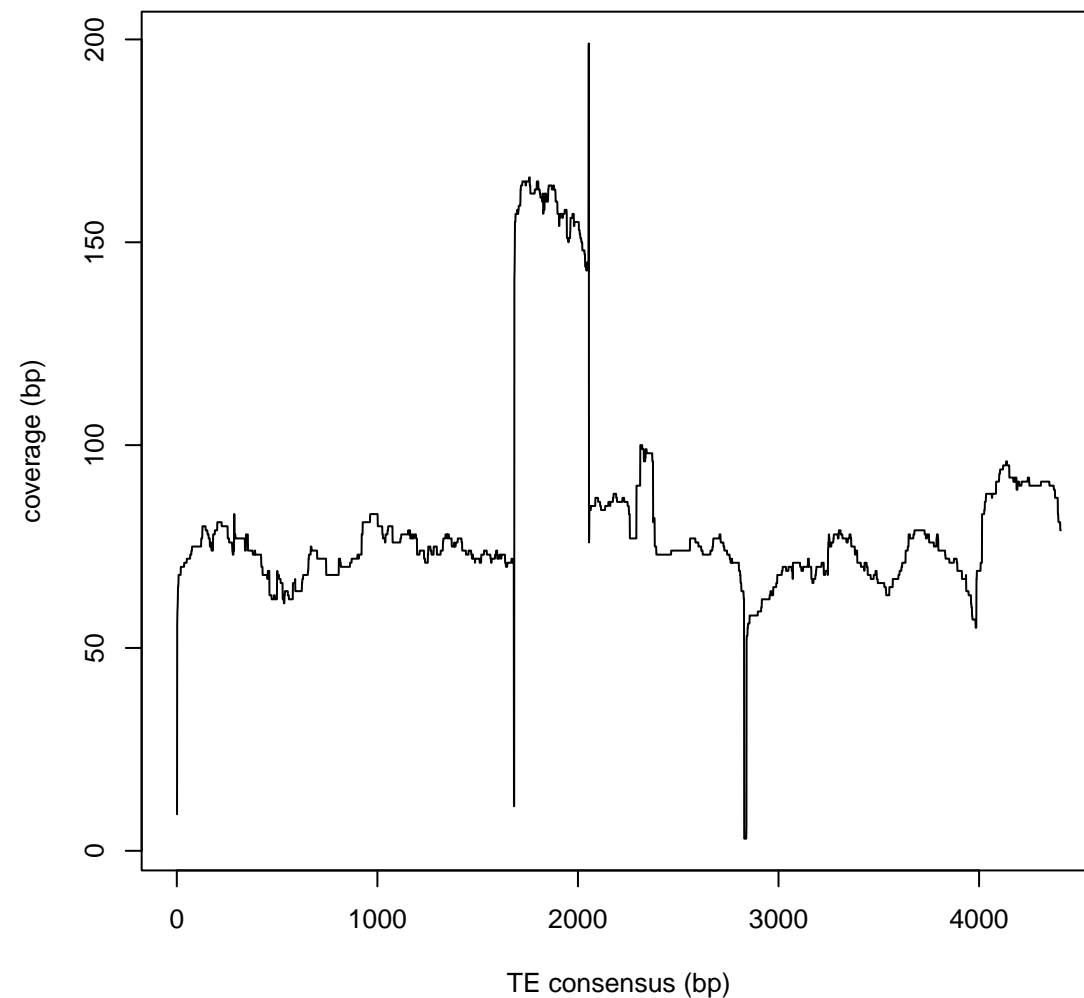


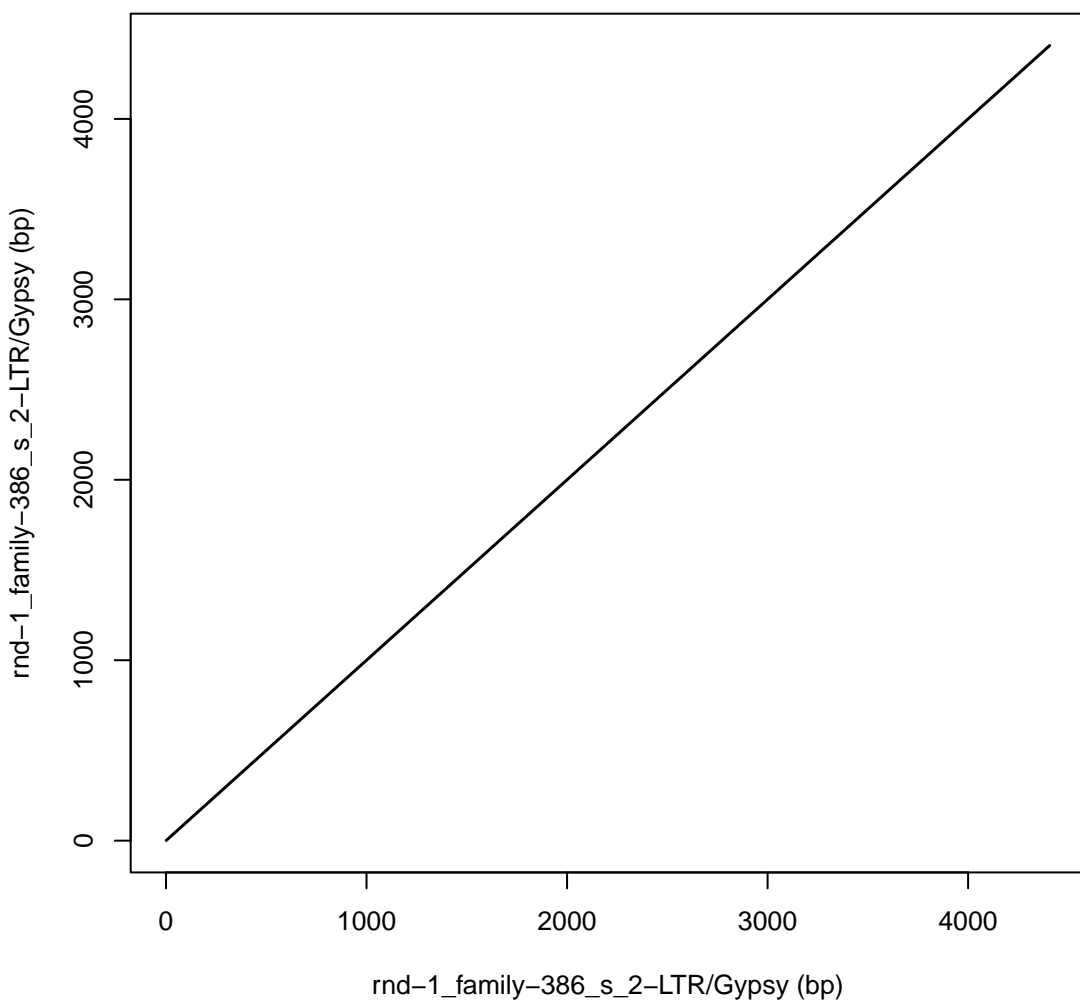
TE: rnd-1\_family-386\_s\_2-LTR/Gypsy  
consensus size: 4407bp; fragments: 732; full length: 3 (>=3966.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

