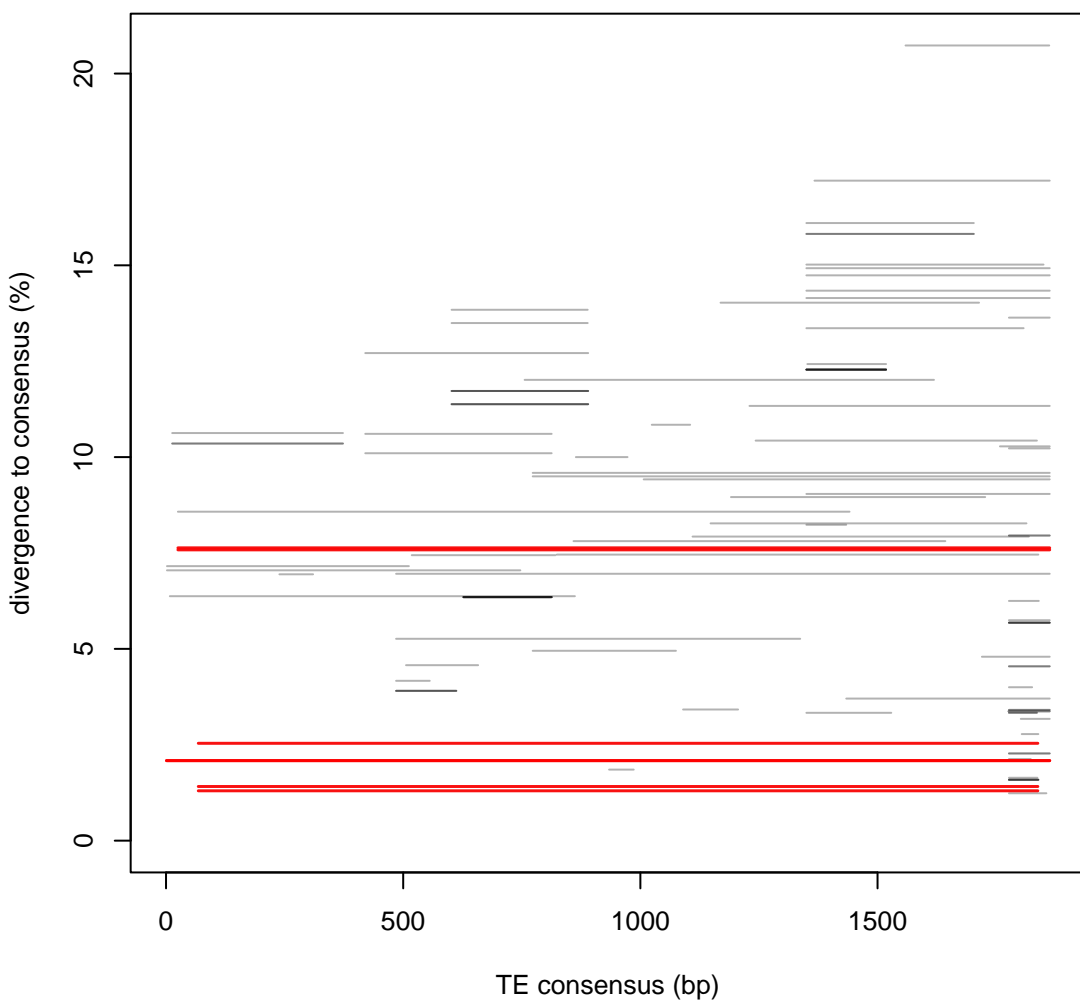
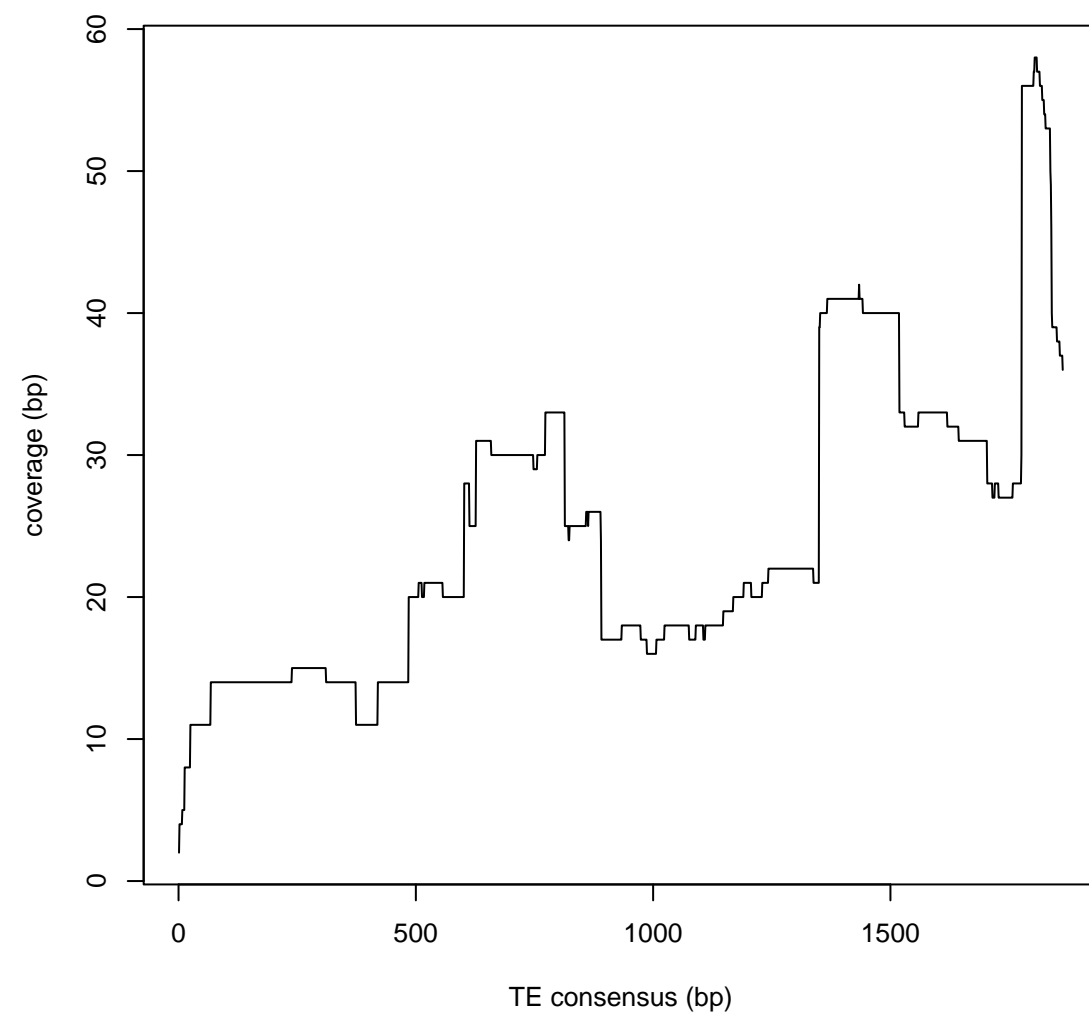


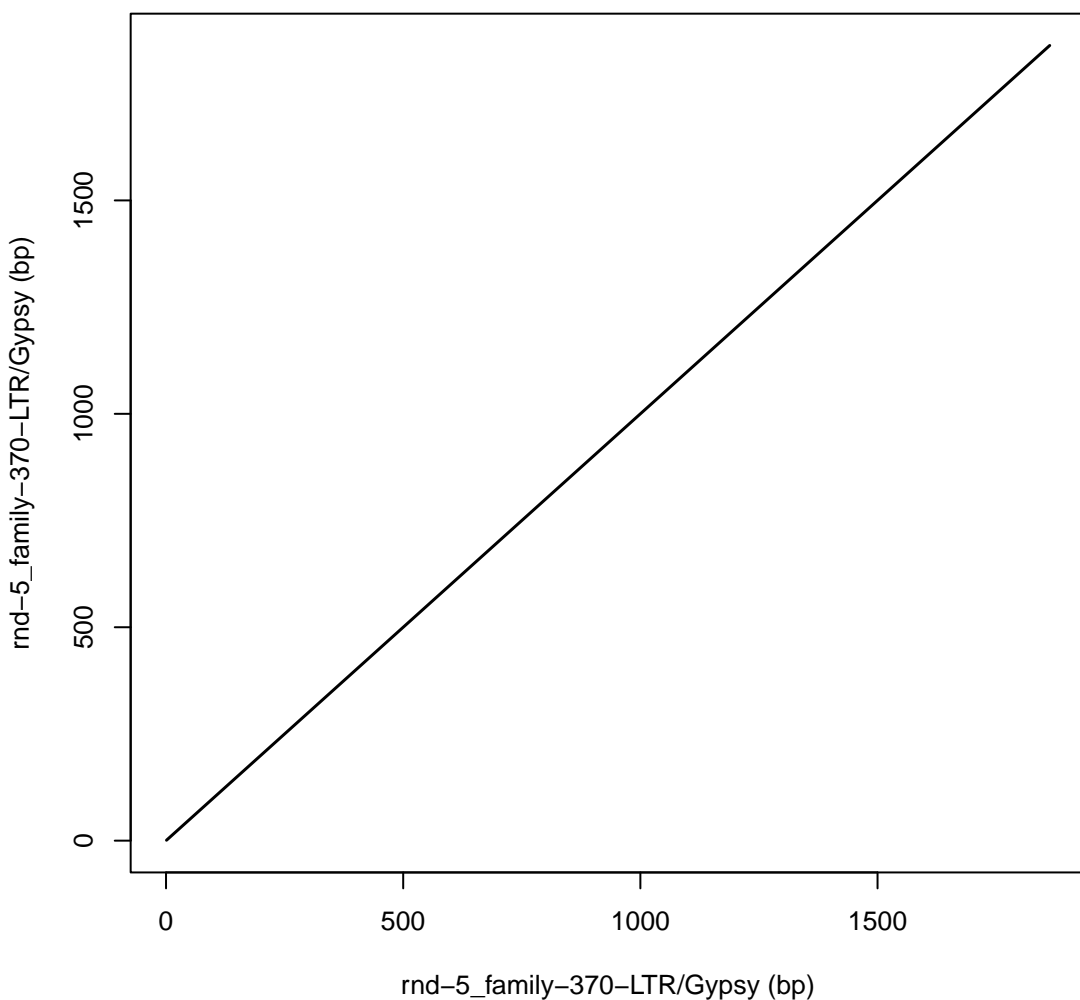
TE: rnd-5\_family-370-LTR/Gypsy  
consensus size: 1863bp; fragments: 111; full length: 7 (>=1676.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

