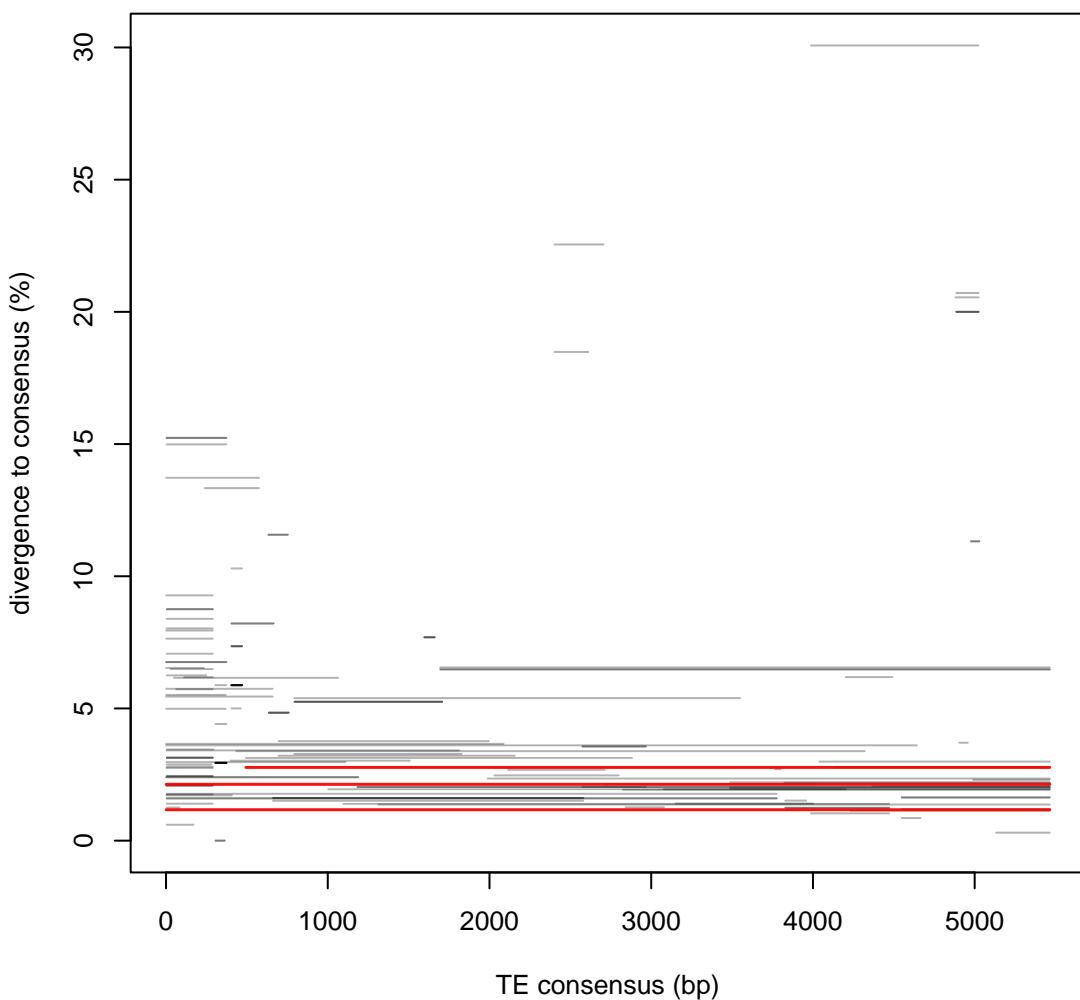
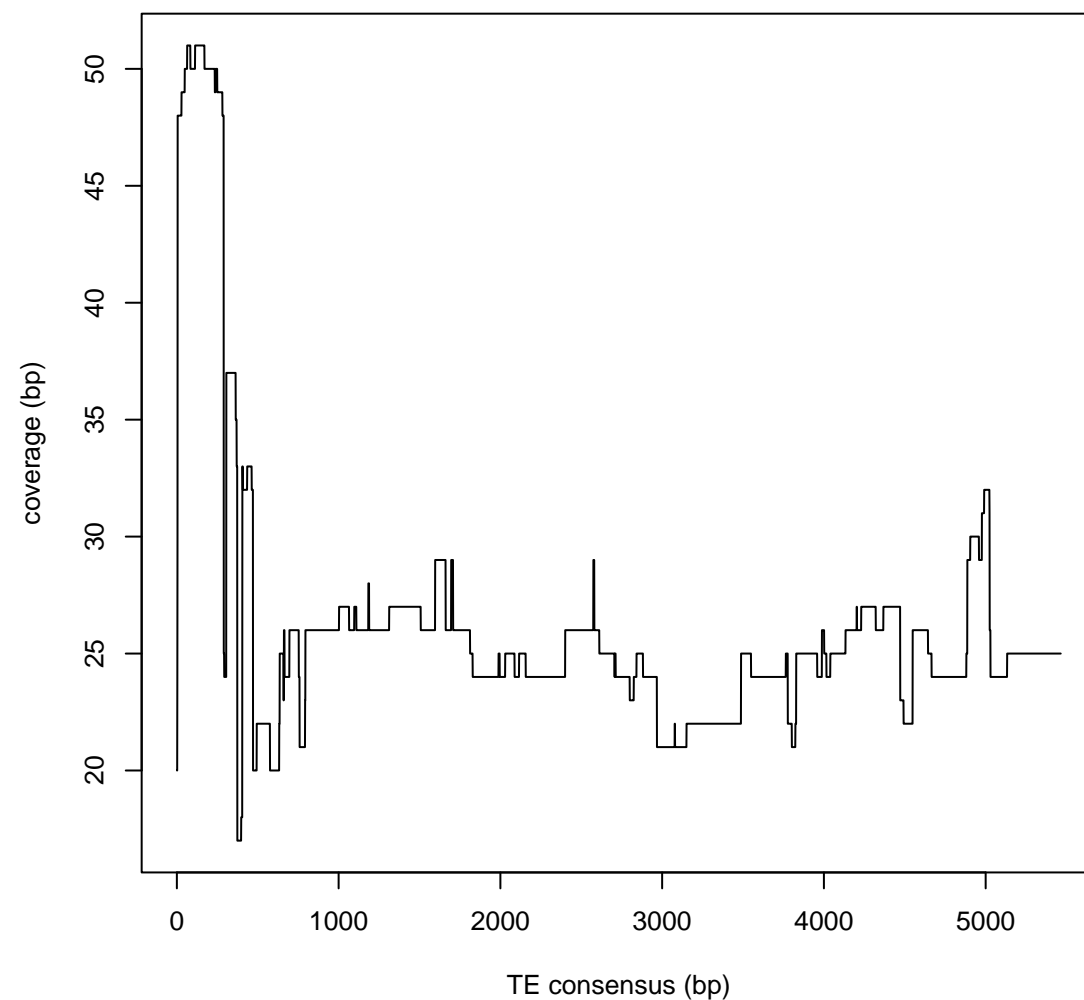


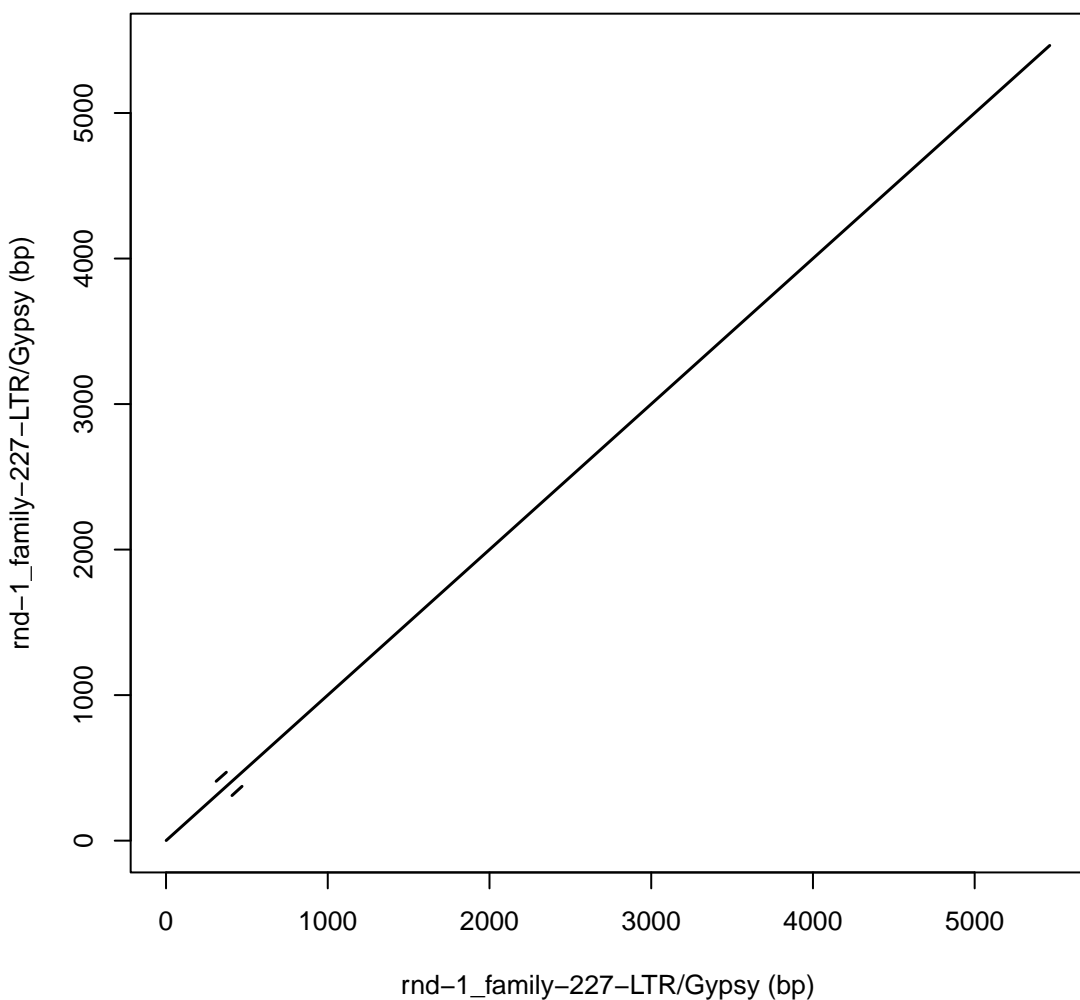
TE: rnd-1\_family-227-LTR/Gypsy  
consensus size: 5464bp; fragments: 153; full length: 3 (>=4917.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

