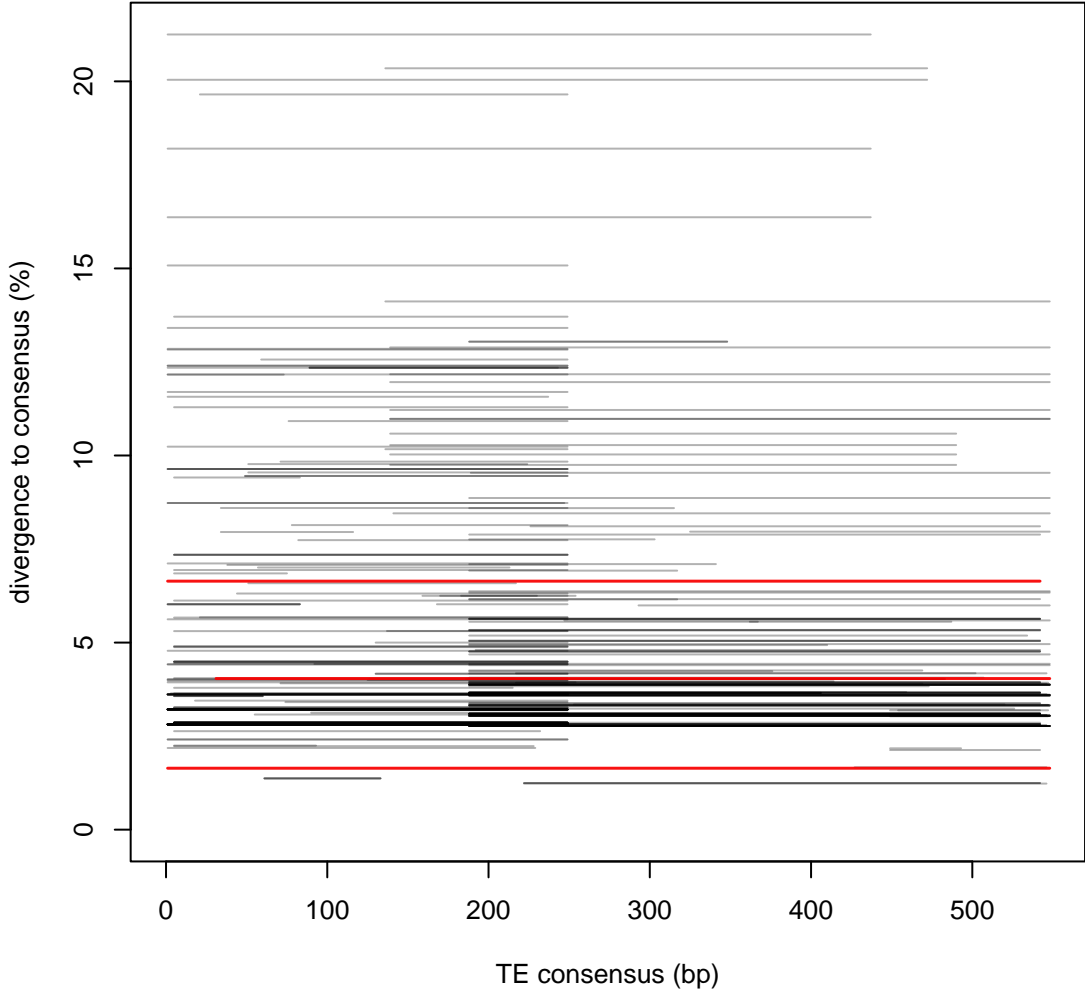
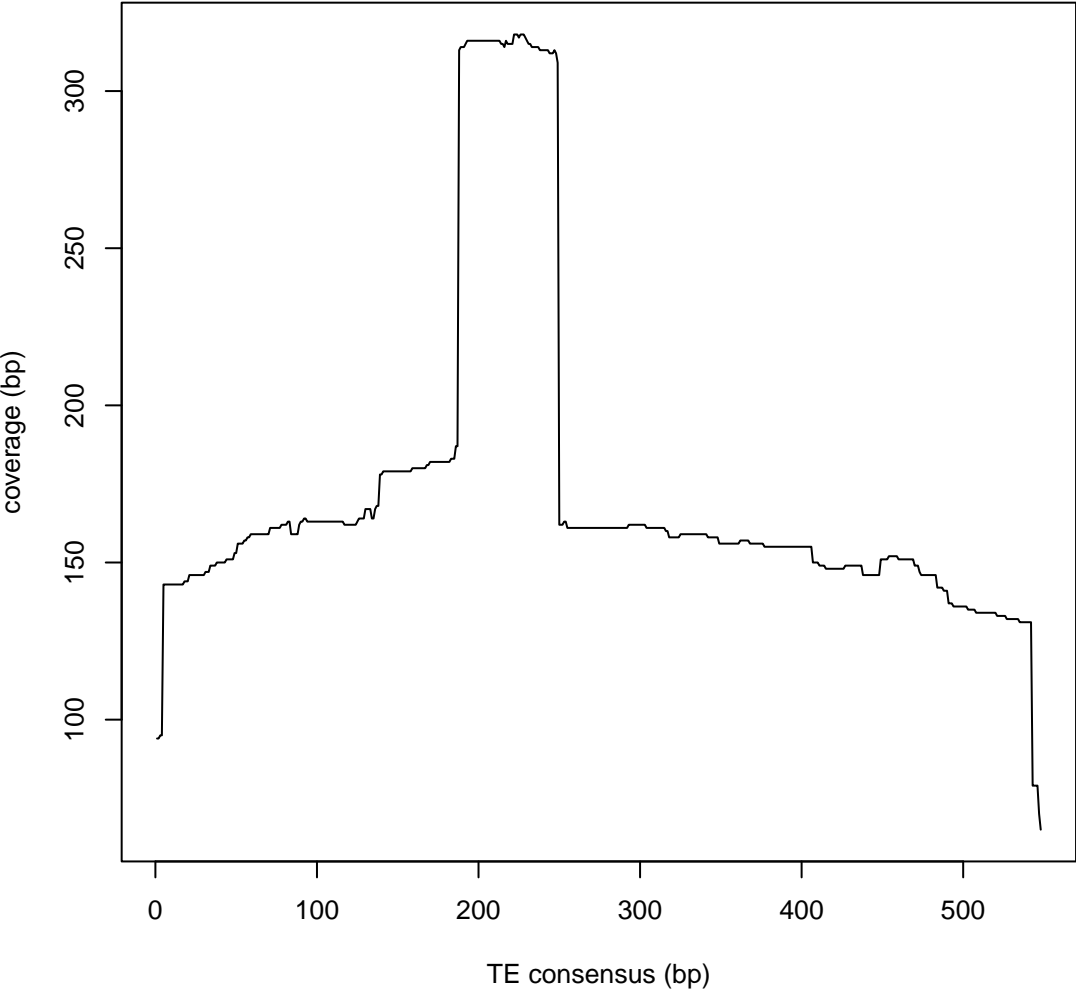


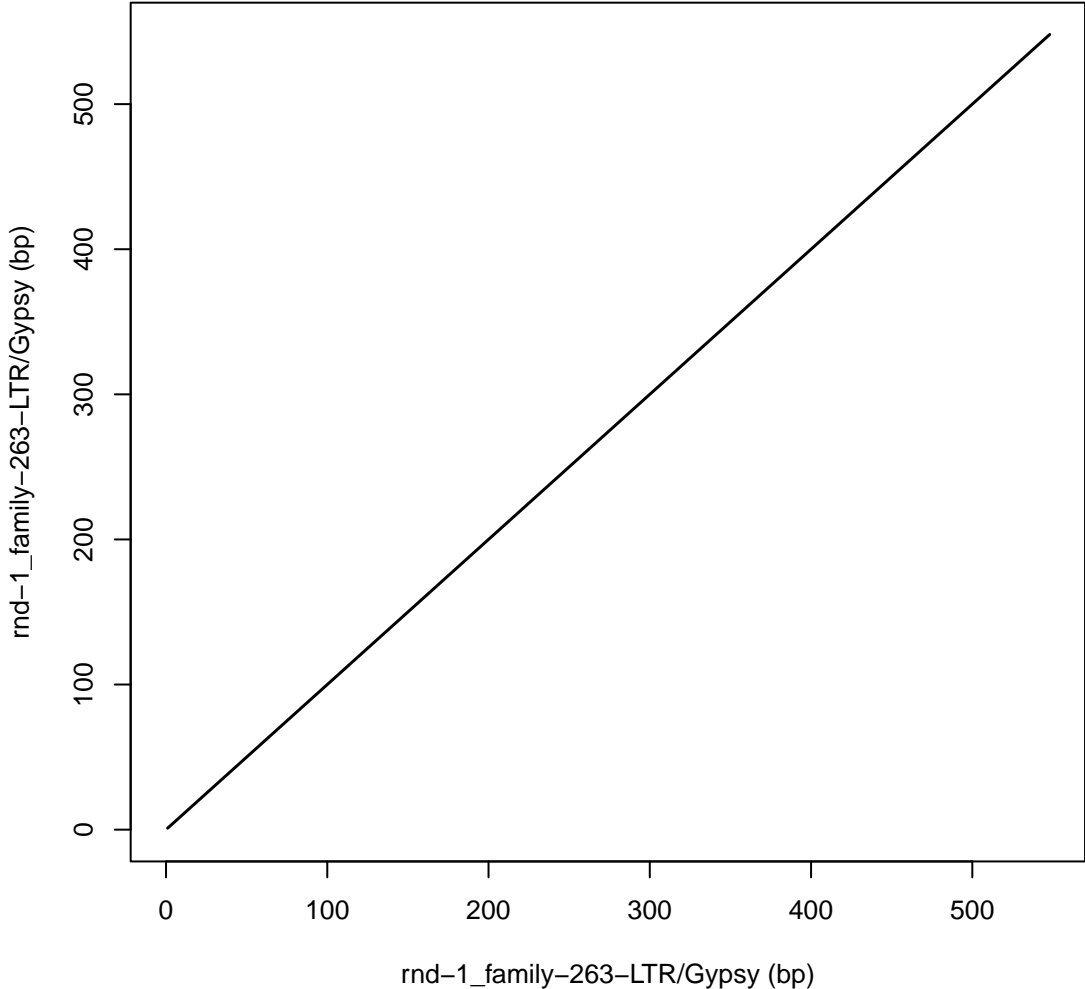
TE: rnd-1\_family-263-LTR/Gypsy  
consensus size: 548bp; fragments: 348; full length: 3 (>=493.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

