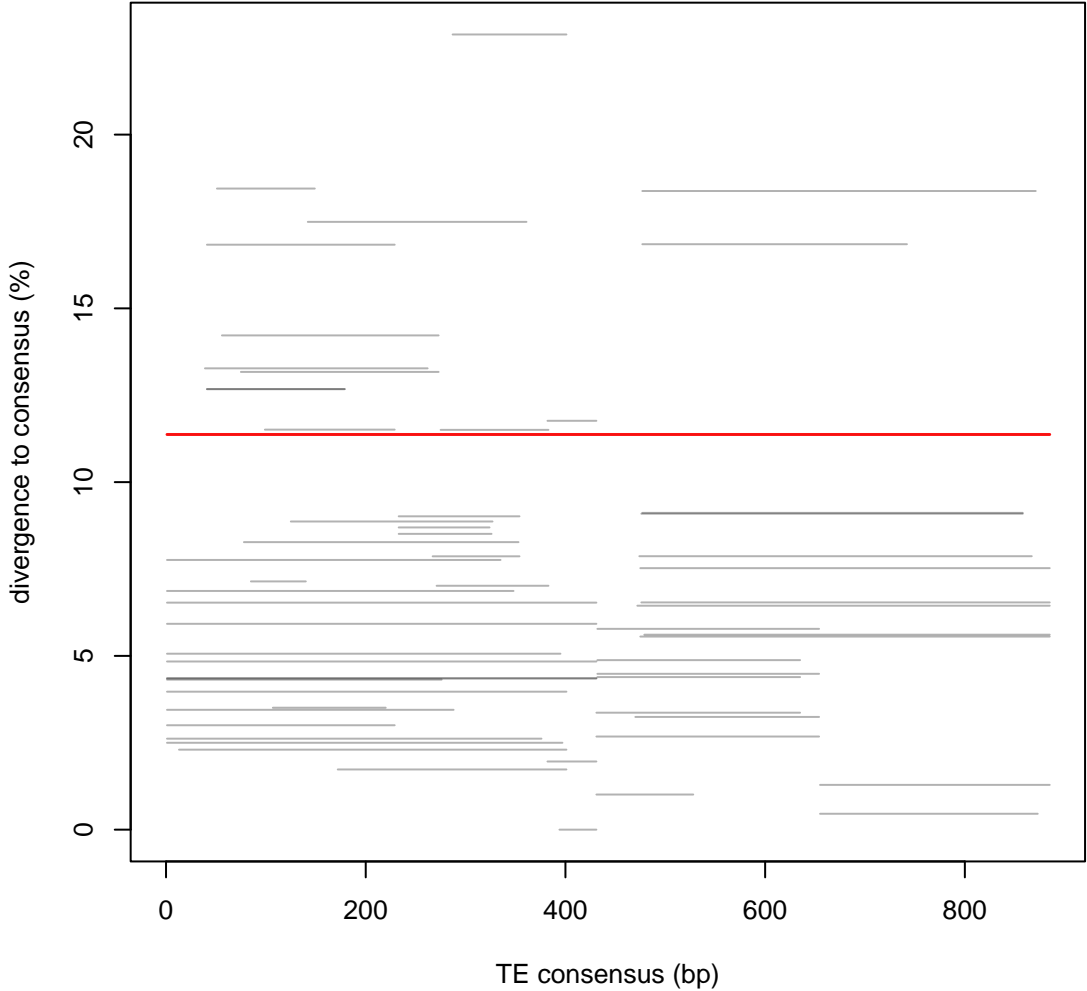
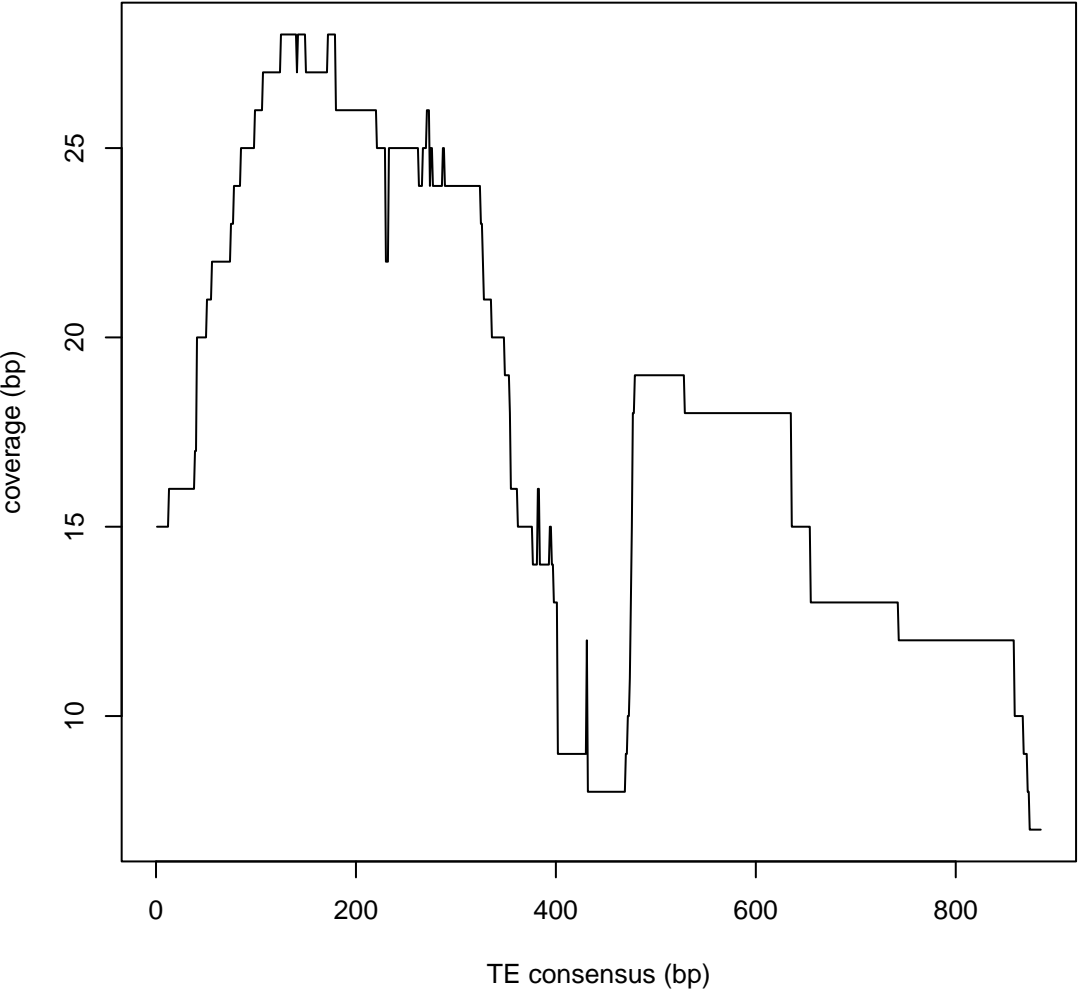


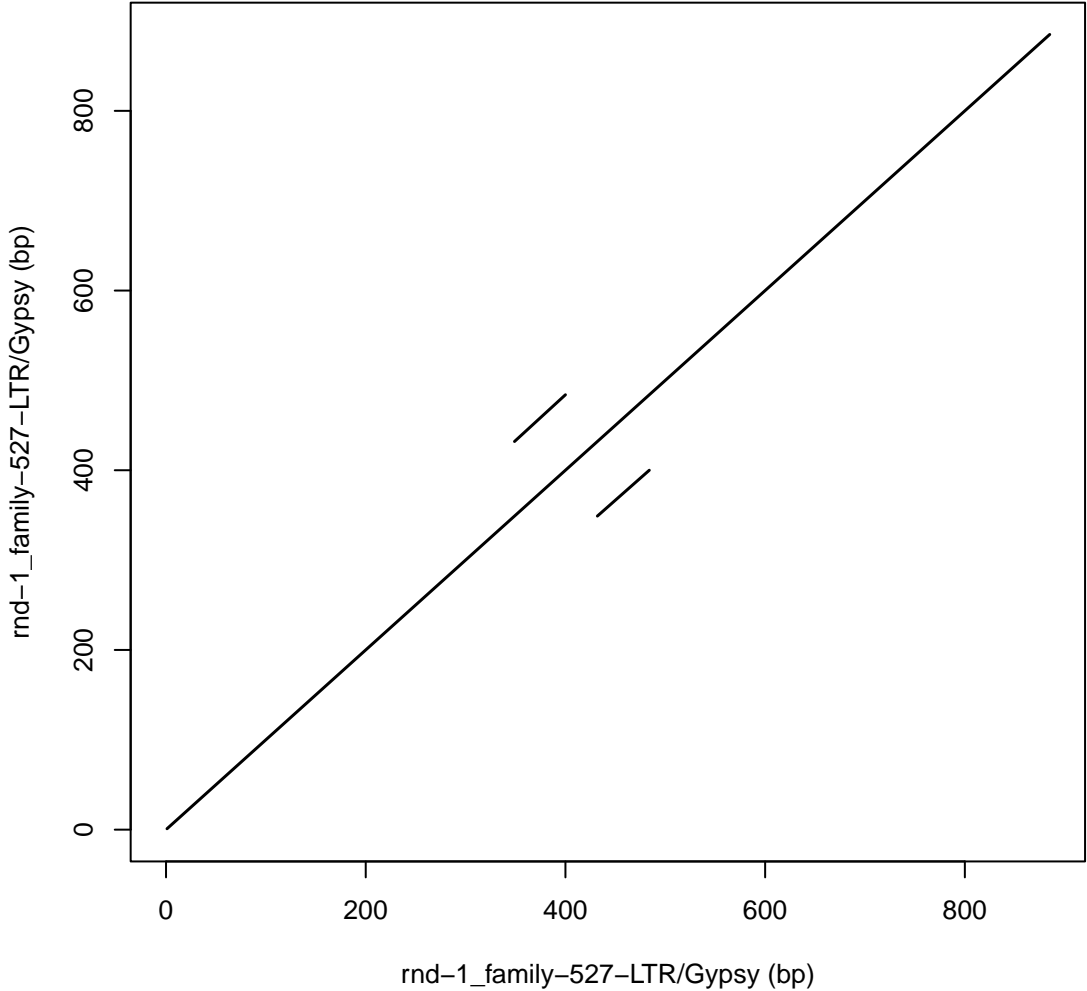
TE: rnd-1\_family-527-LTR/Gypsy  
consensus size: 885bp; fragments: 60; full length: 1 (>=796.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

