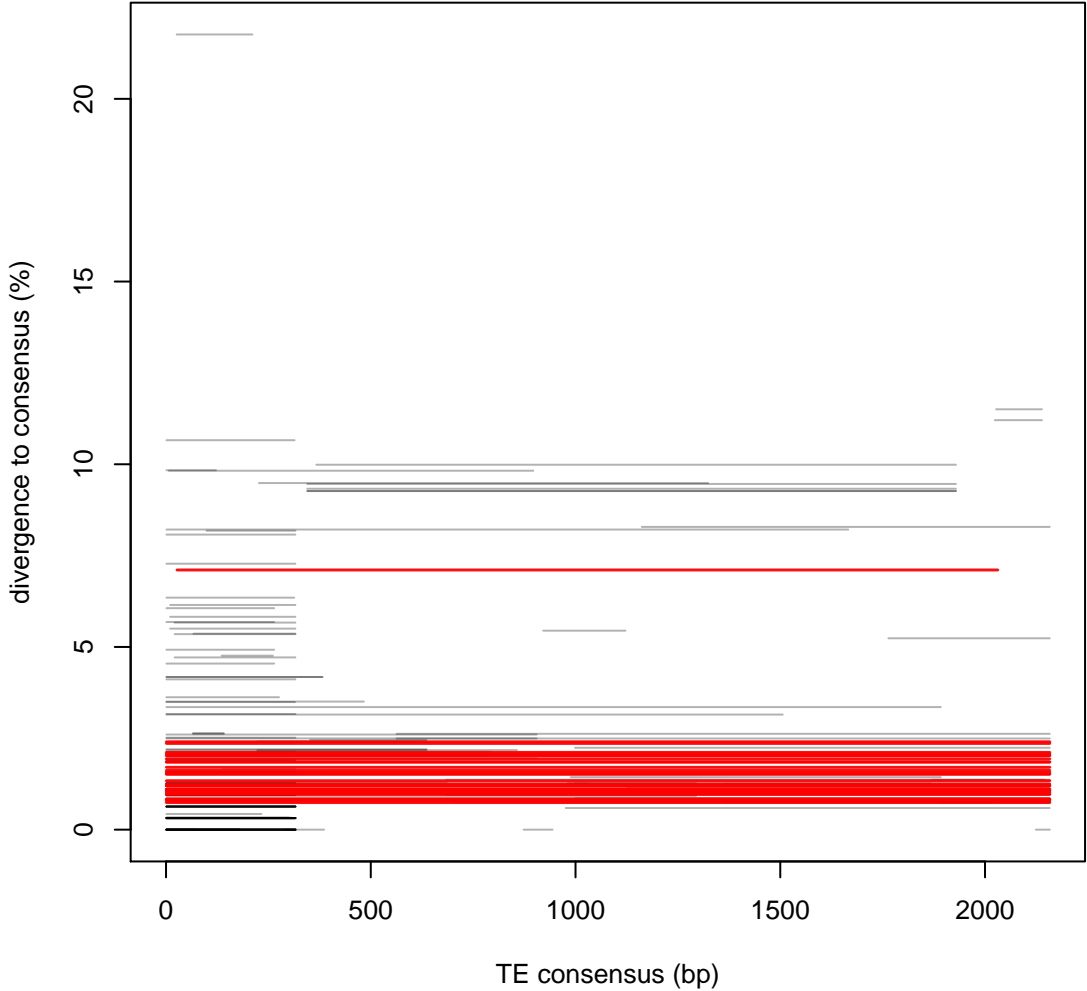
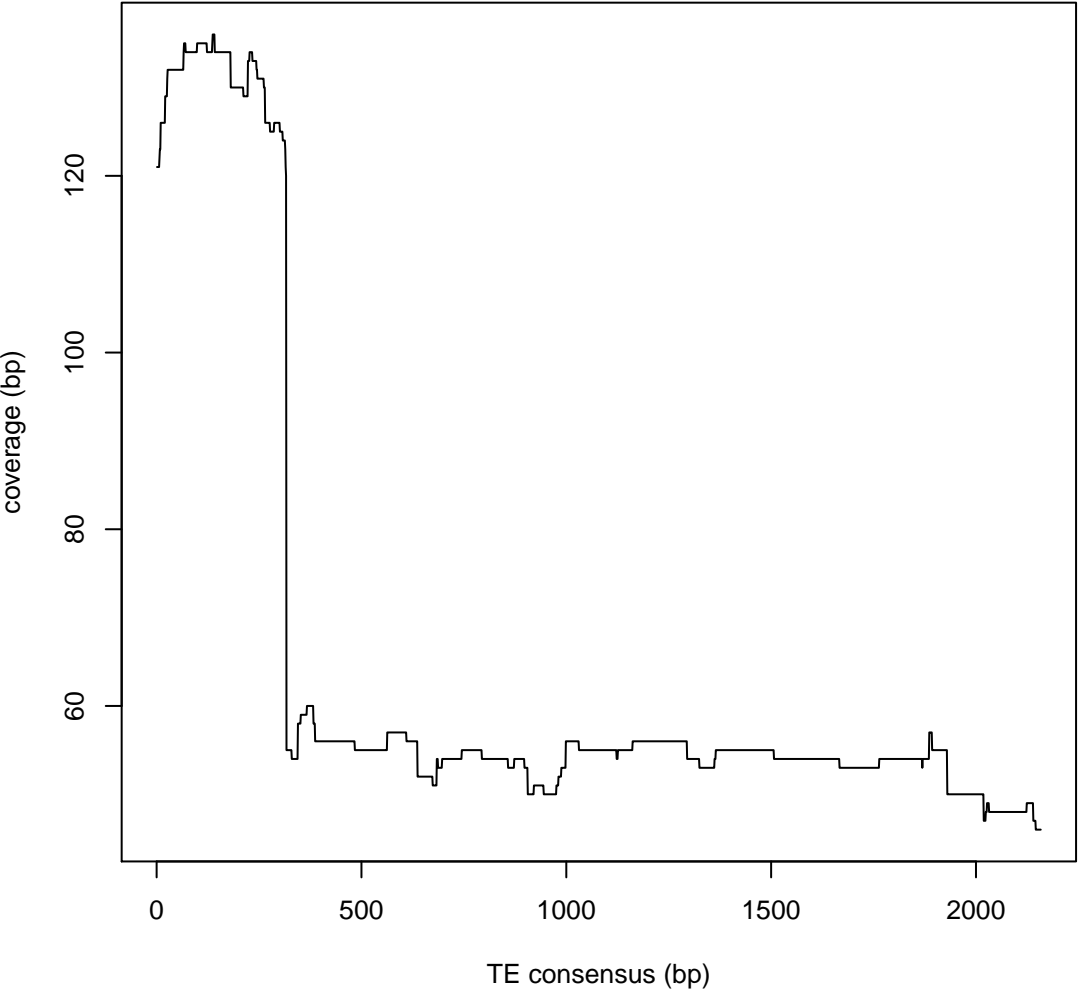


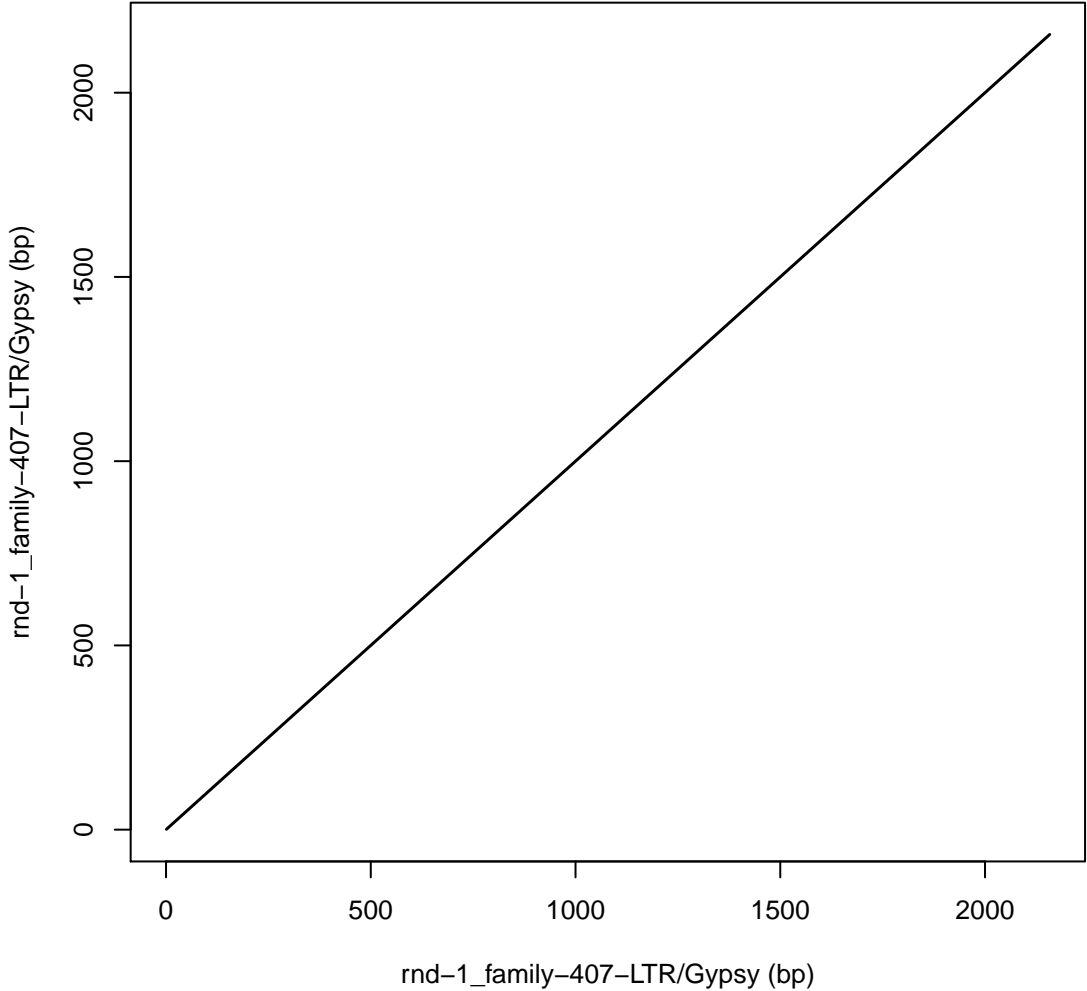
TE: rnd-1\_family-407-LTR/Gypsy  
consensus size: 2158bp; fragments: 177; full length: 31 (>=1942.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

