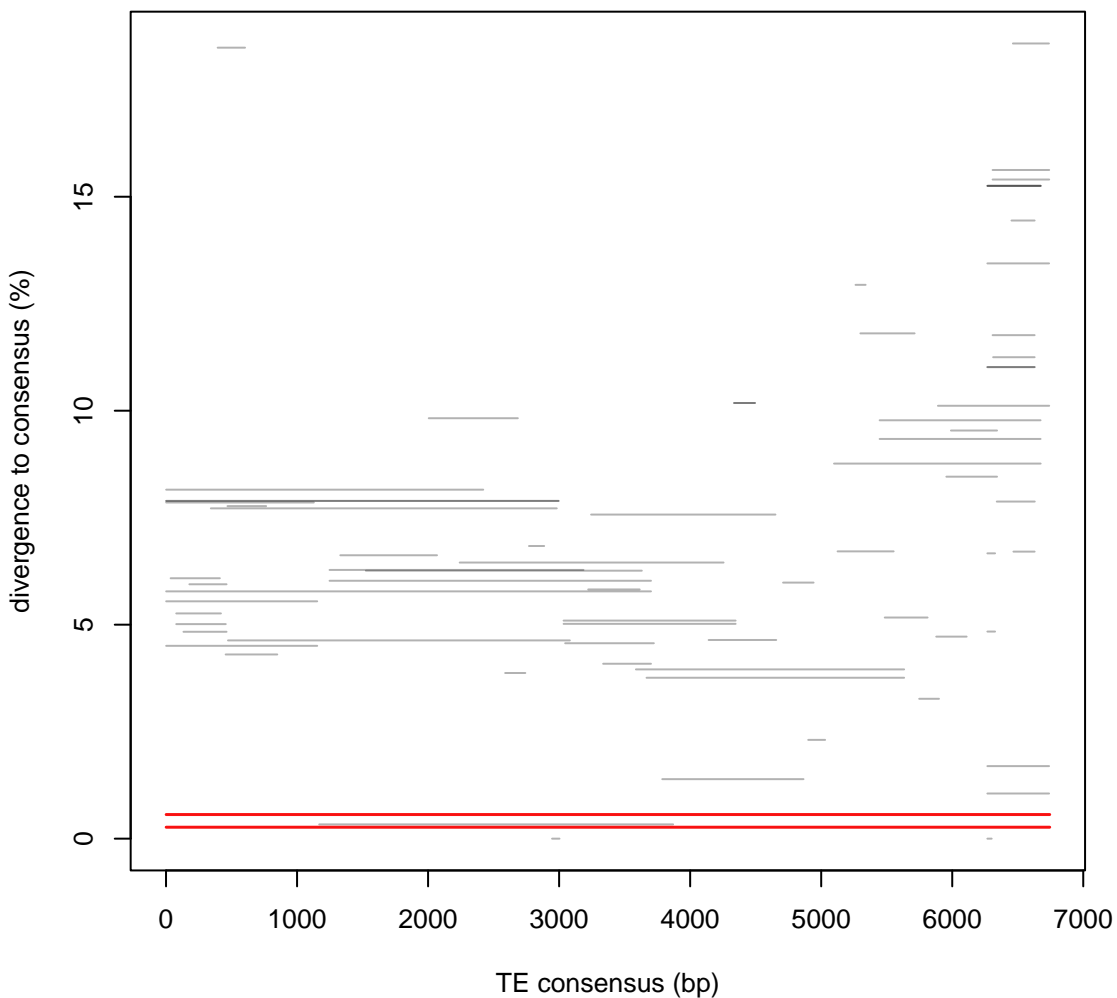
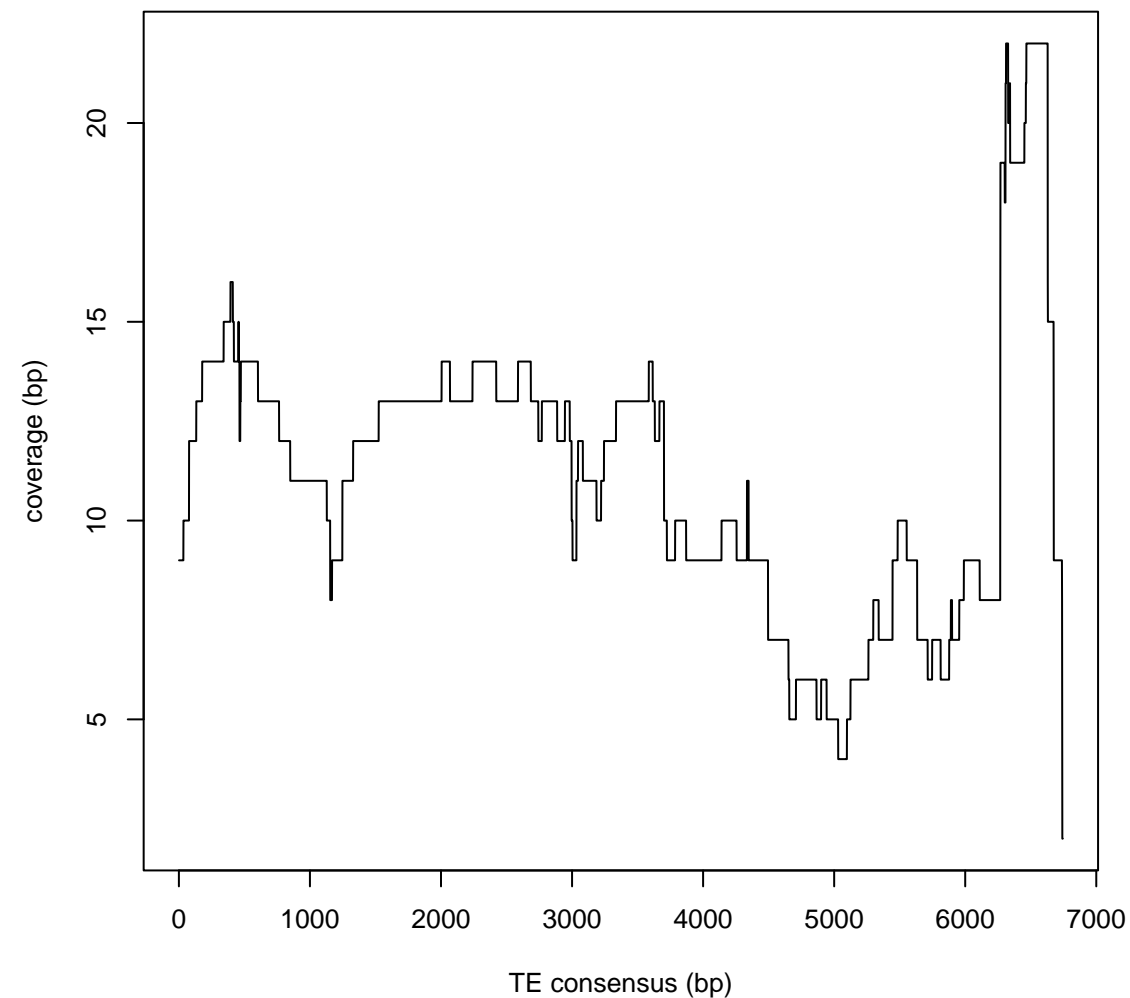


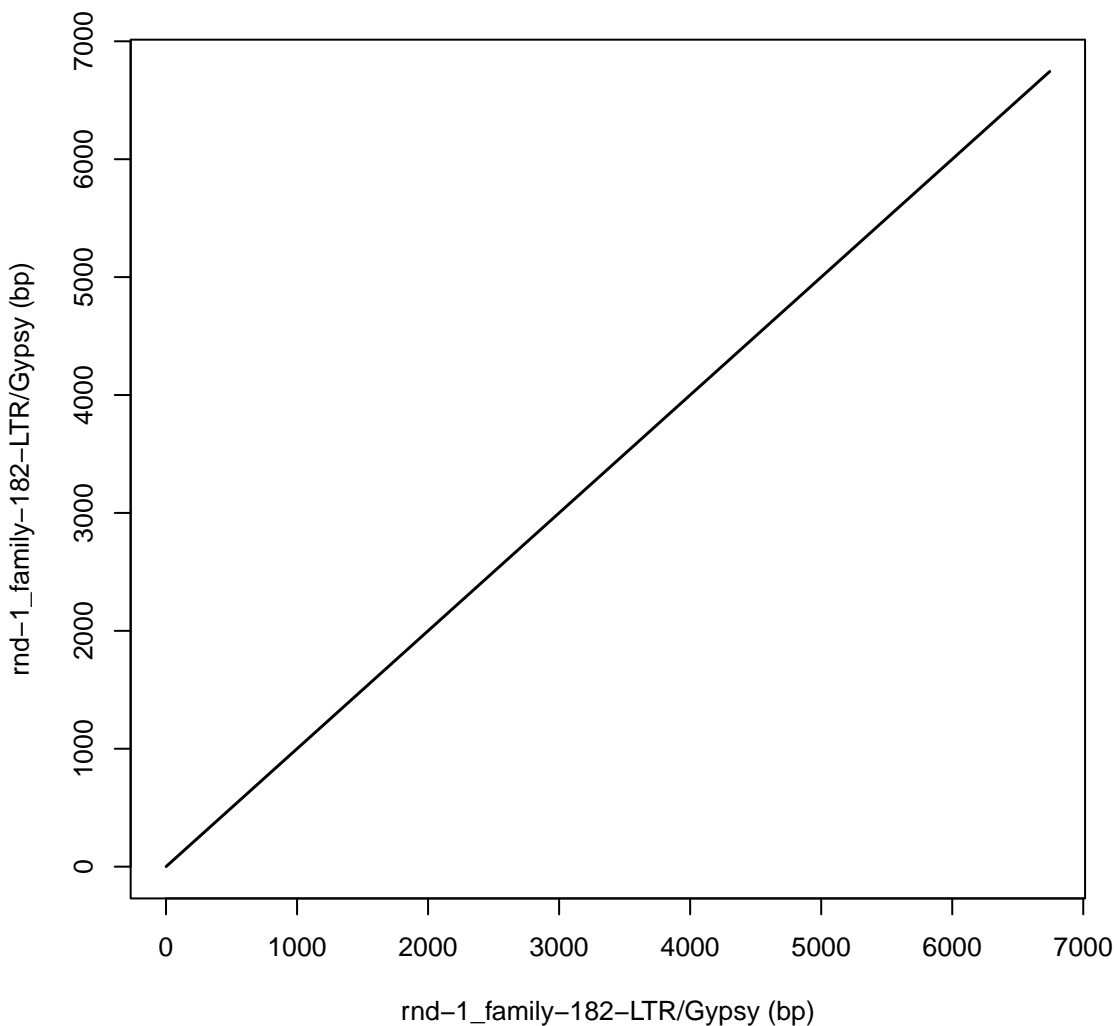
TE: rnd-1\_family-182-LTR/Gypsy  
consensus size: 6744bp; fragments: 74; full length: 2 (>=6069.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

