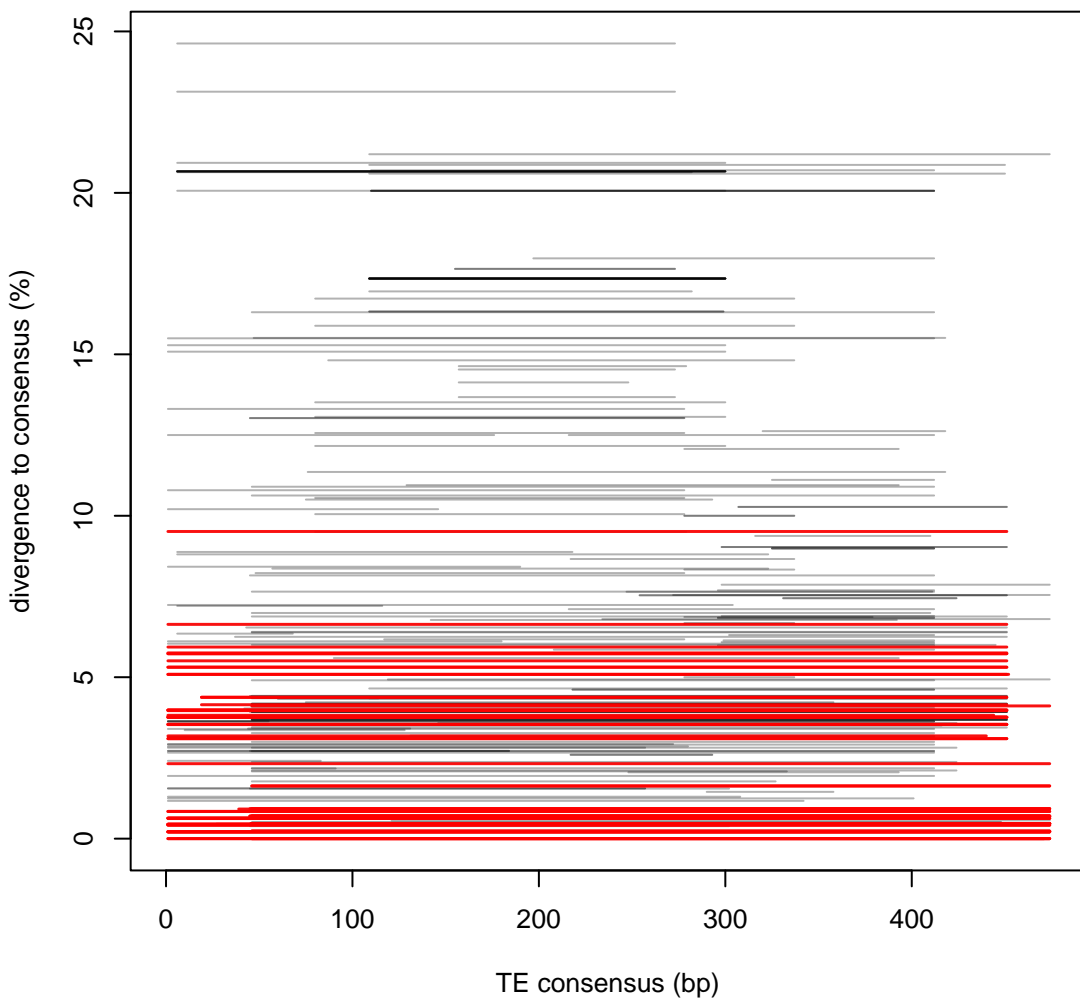
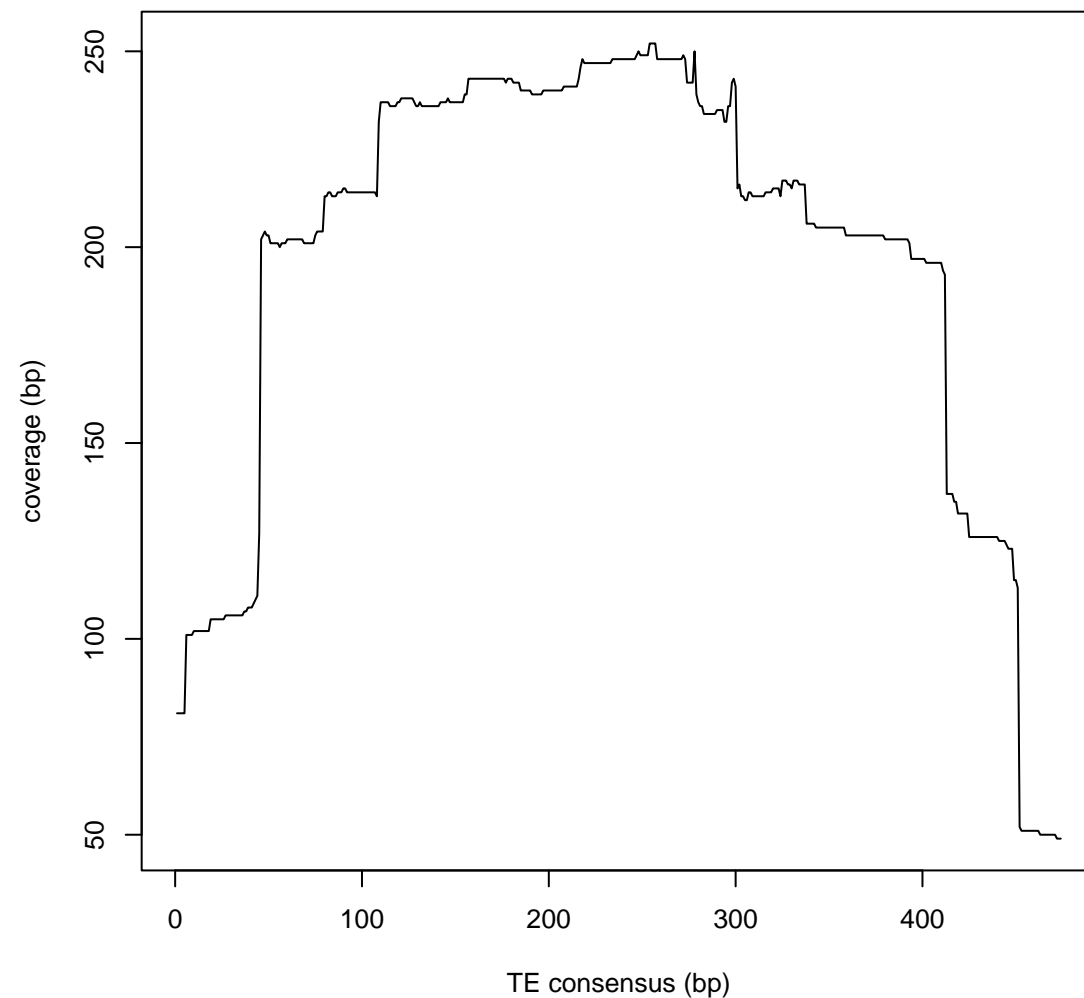


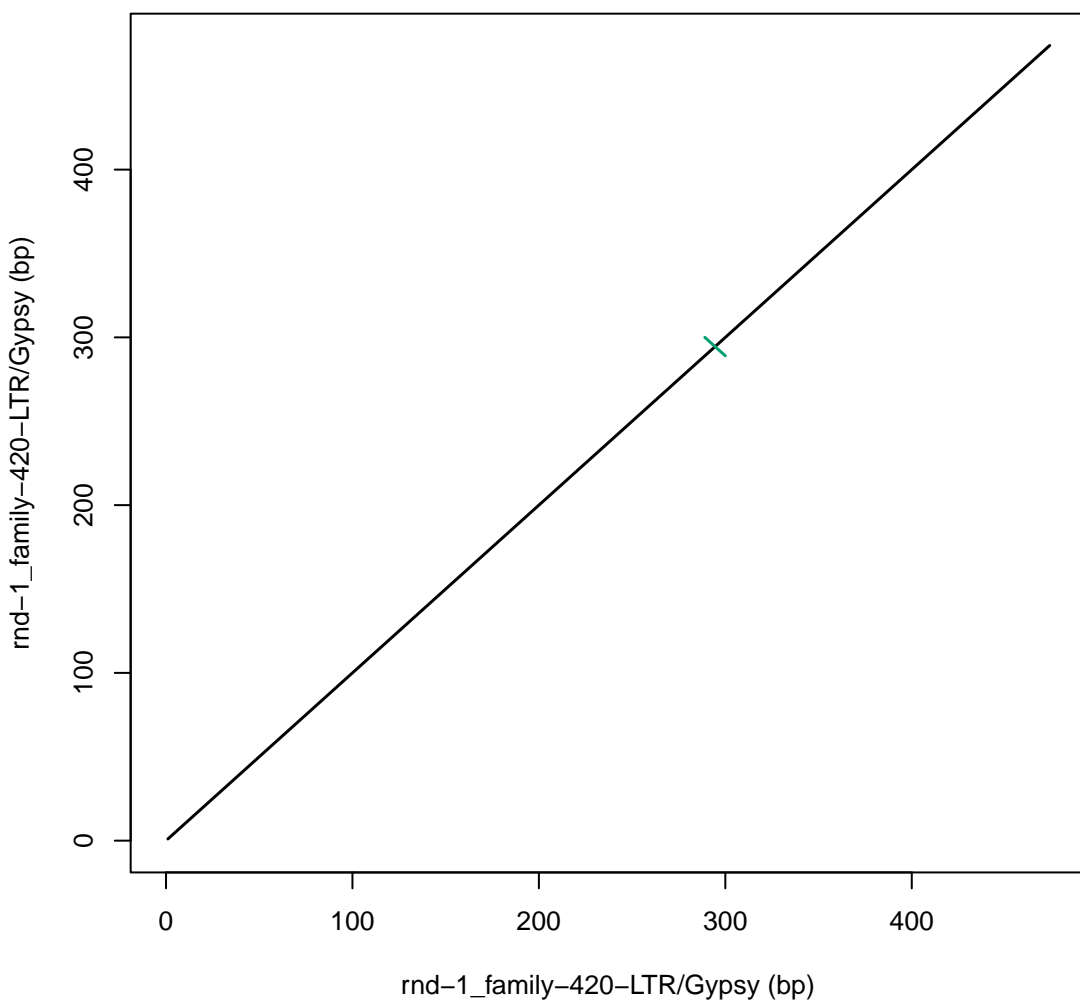
TE: rnd-1\_family-420-LTR/Gypsy  
consensus size: 474bp; fragments: 306; full length: 76 ( $\geq 426.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

