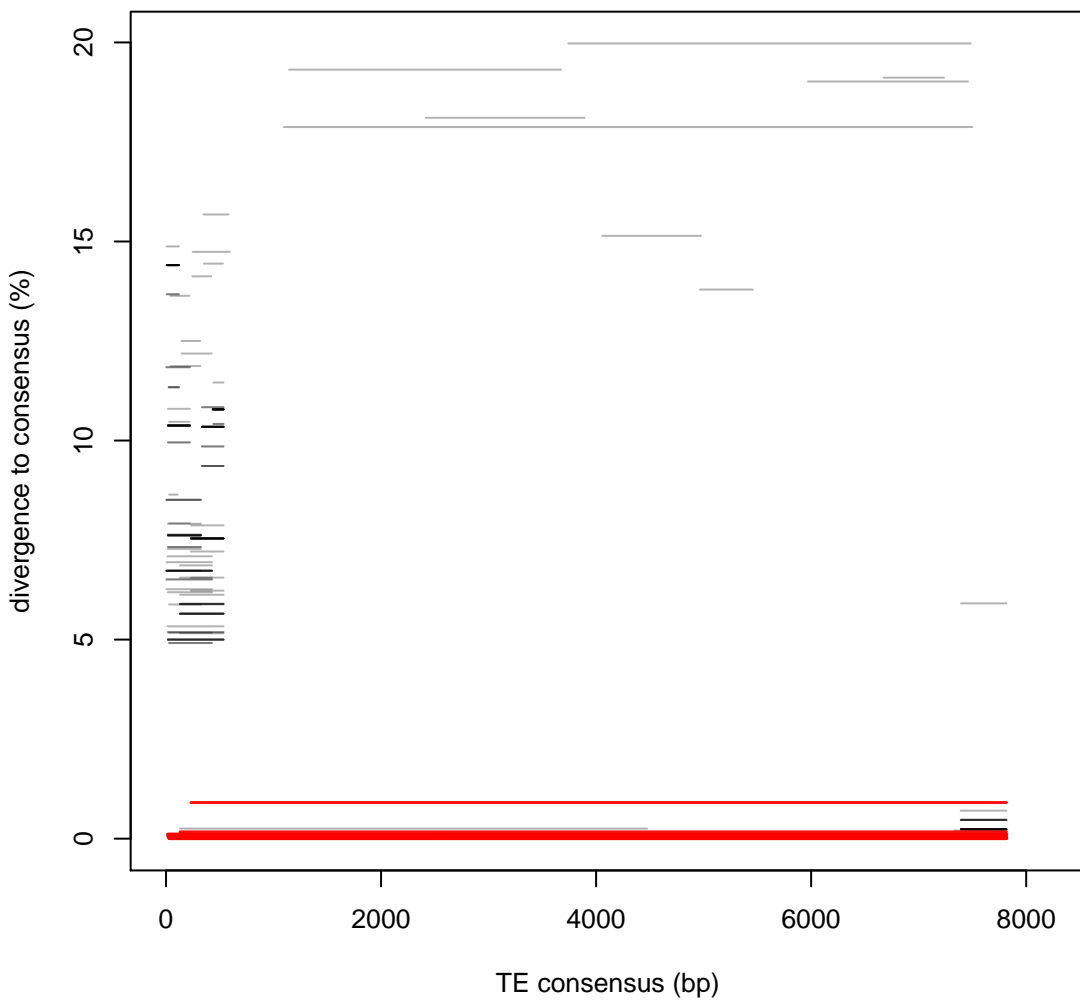
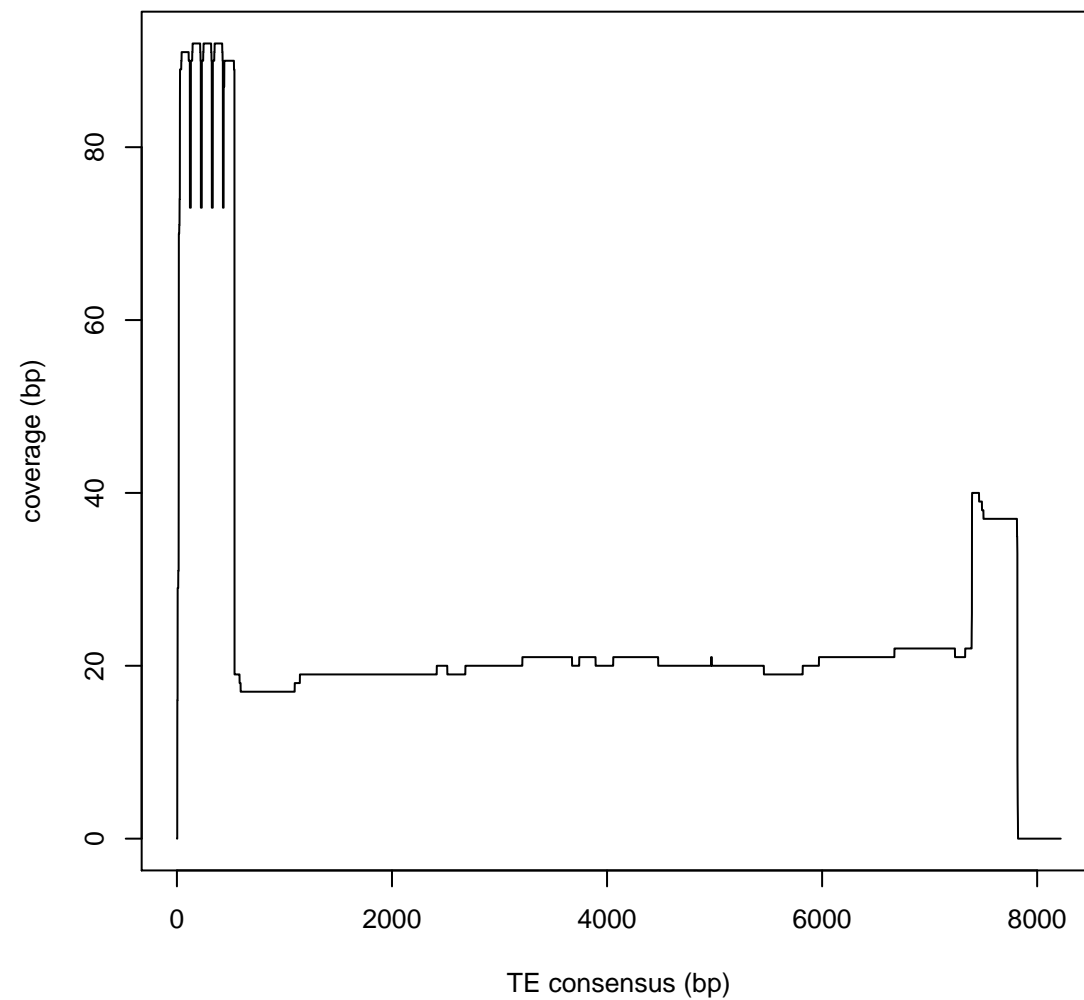


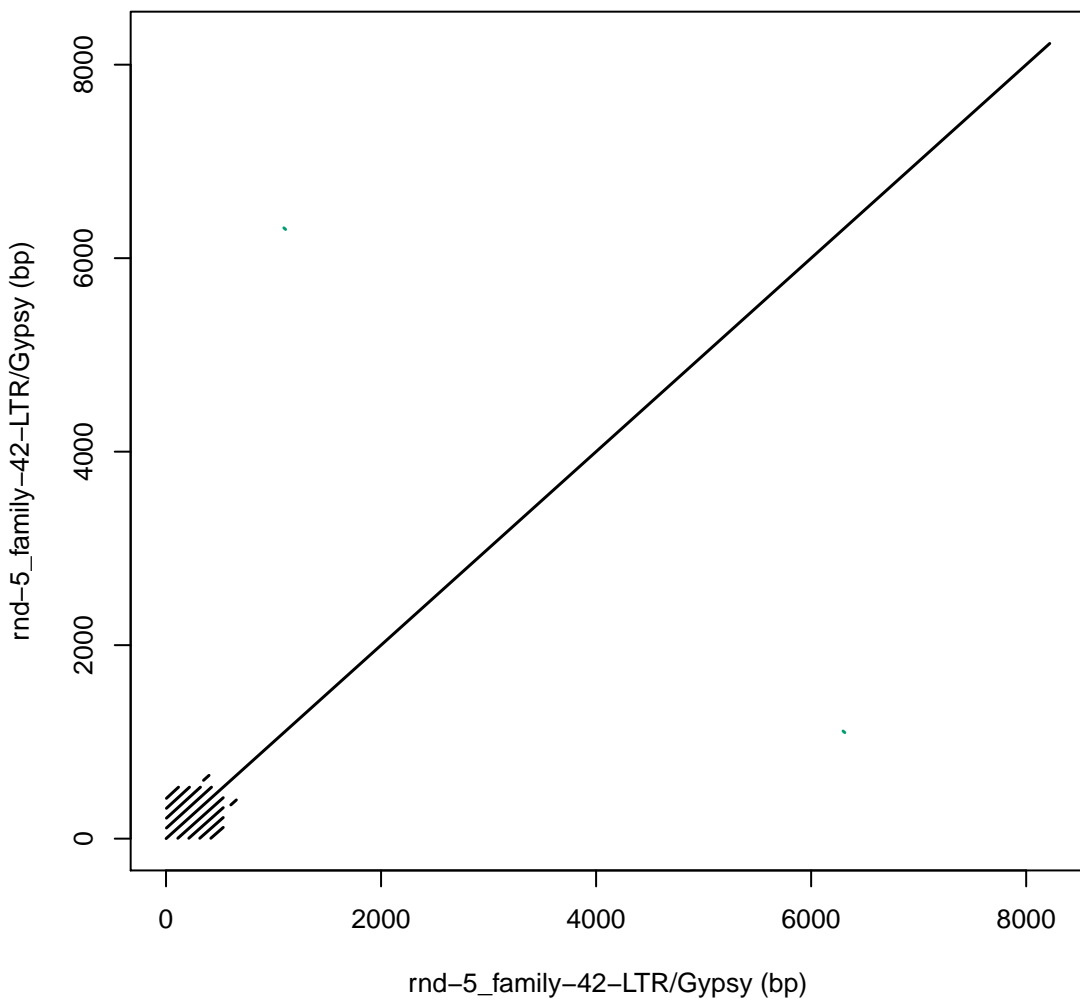
TE: rnd-5\_family-42-LTR/Gypsy  
consensus size: 8219bp; fragments: 195; full length: 15 ( $\geq 7397.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

