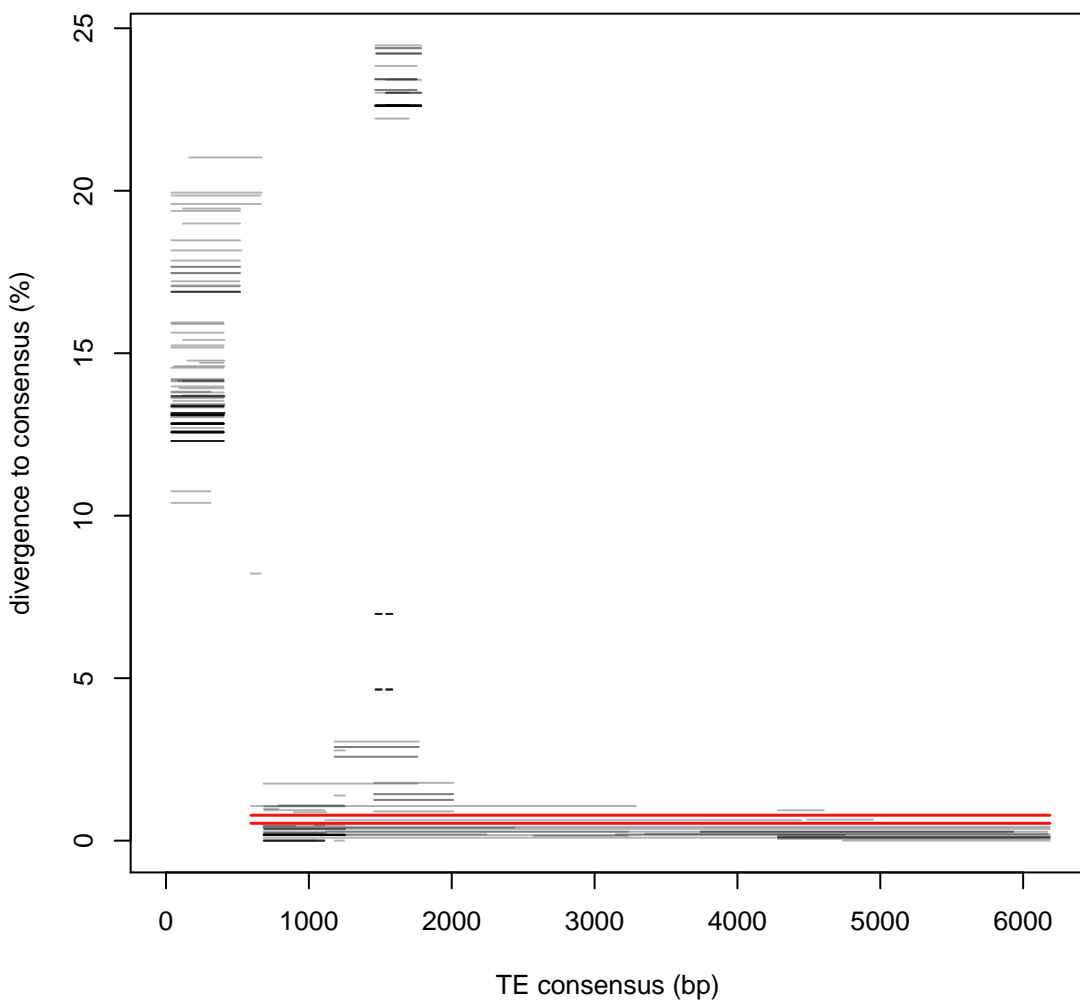
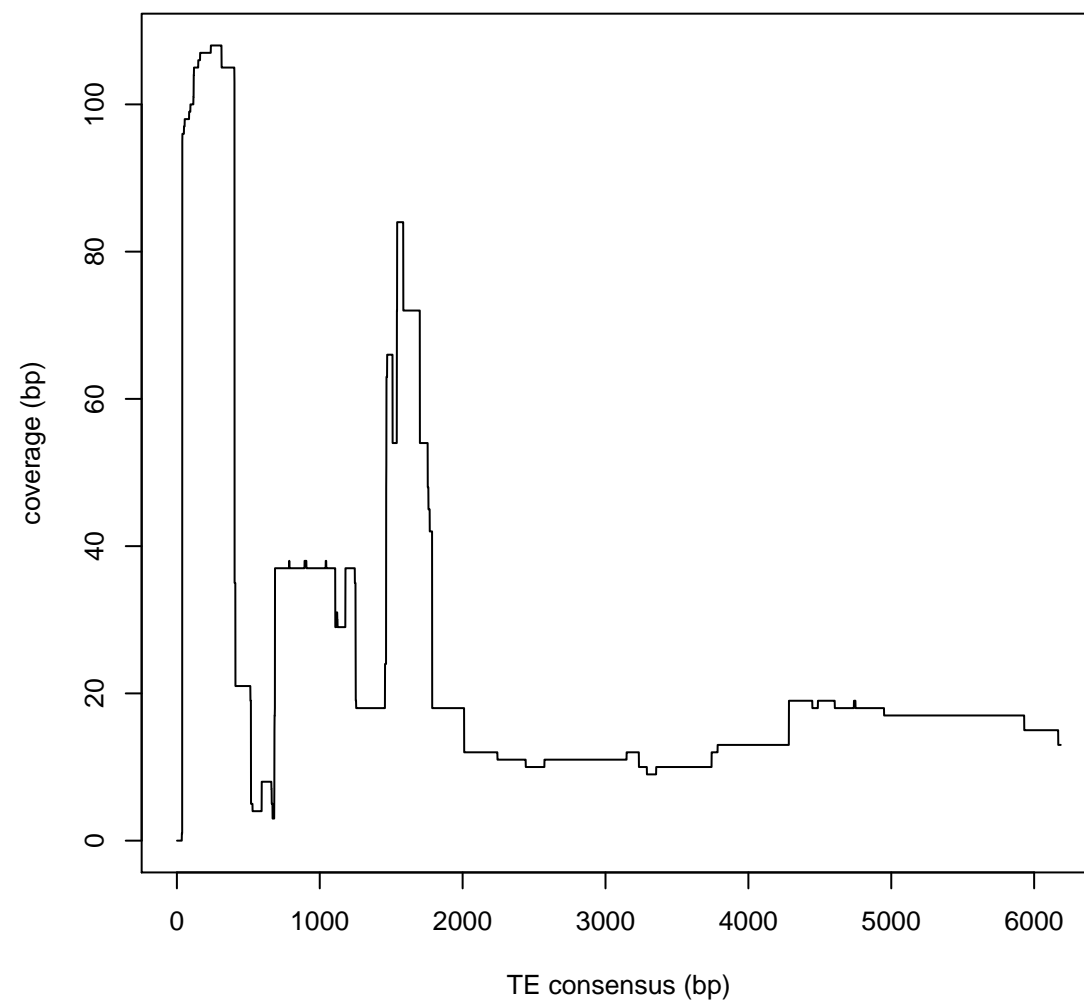


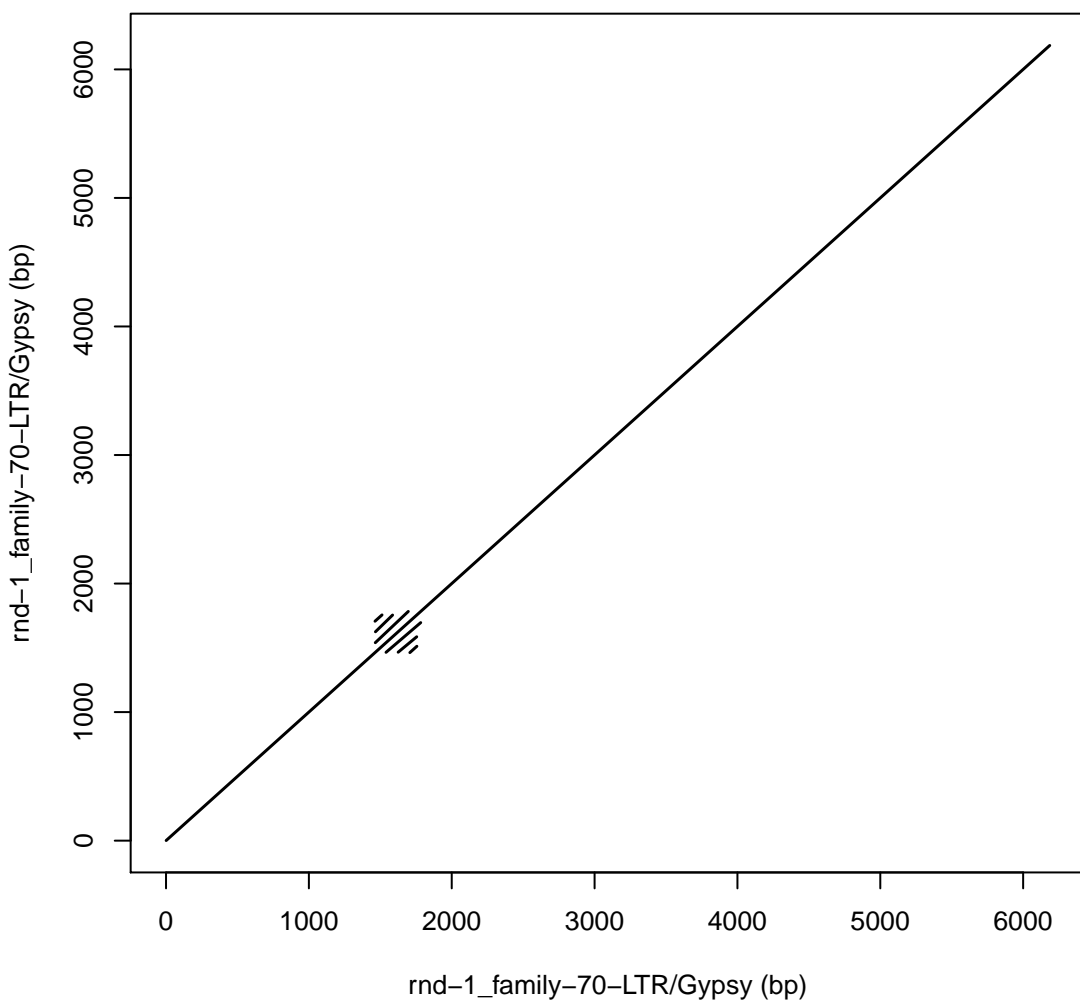
TE: rnd-1\_family-70-LTR/Gypsy  
consensus size: 6186bp; fragments: 251; full length: 2 (>=5567.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

