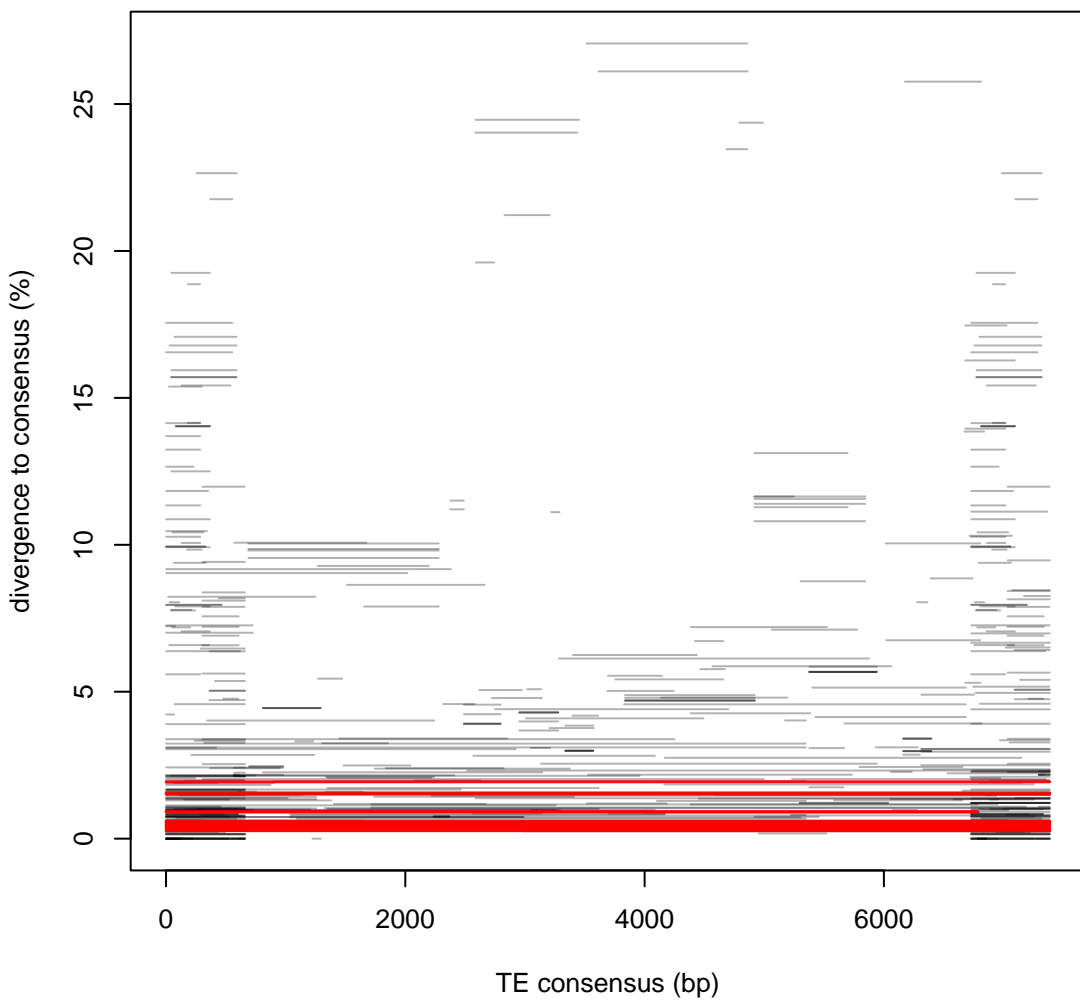
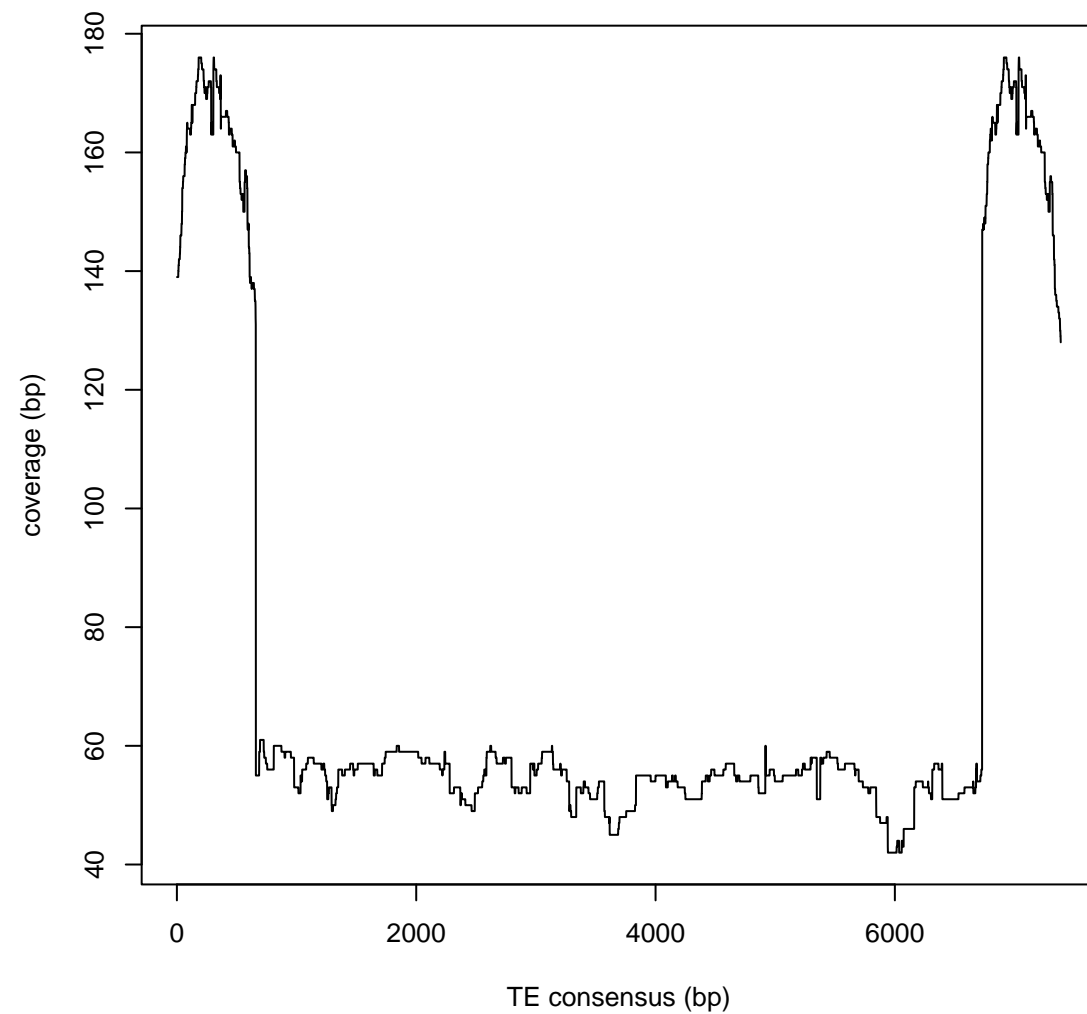


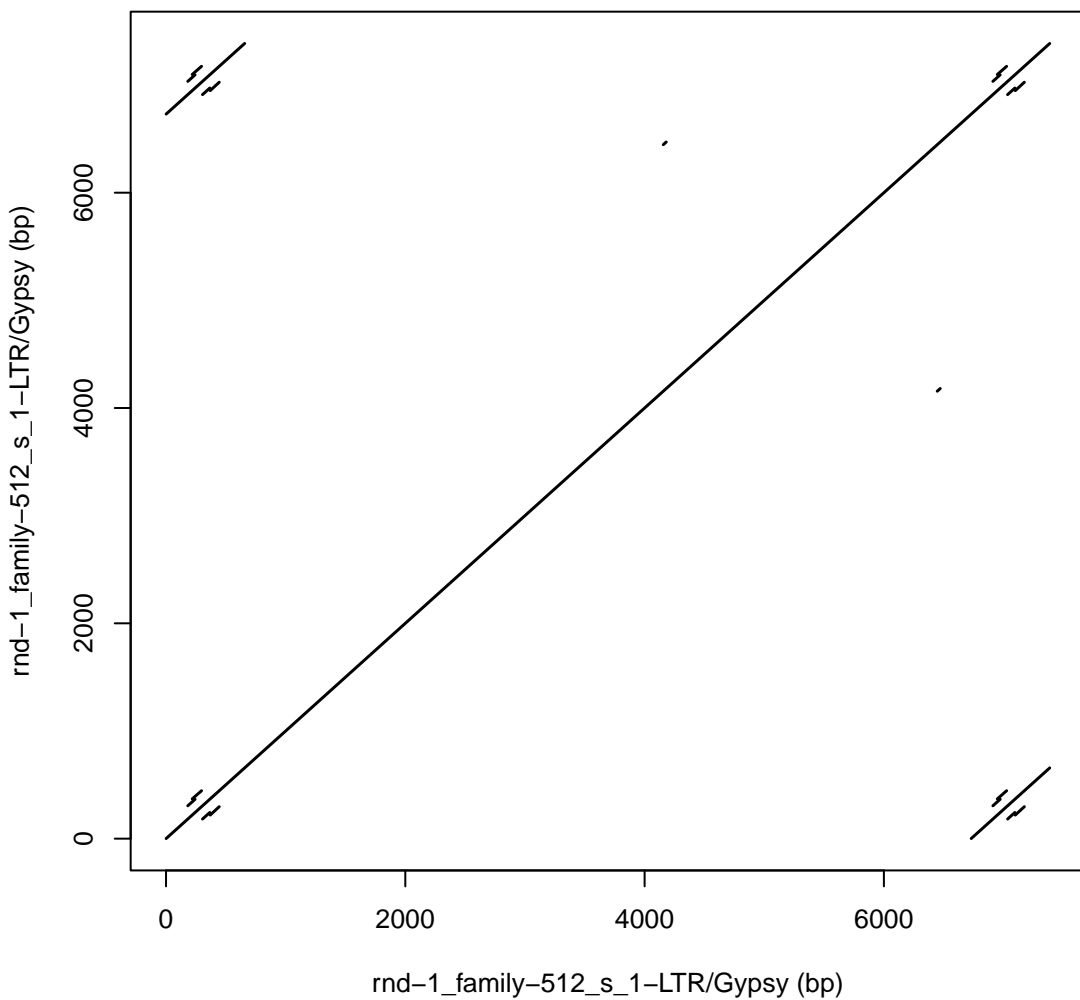
TE: rnd-1\_family-512\_s\_1-LTR/Gypsy  
consensus size: 7386bp; fragments: 578; full length: 16 (>=6647.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

