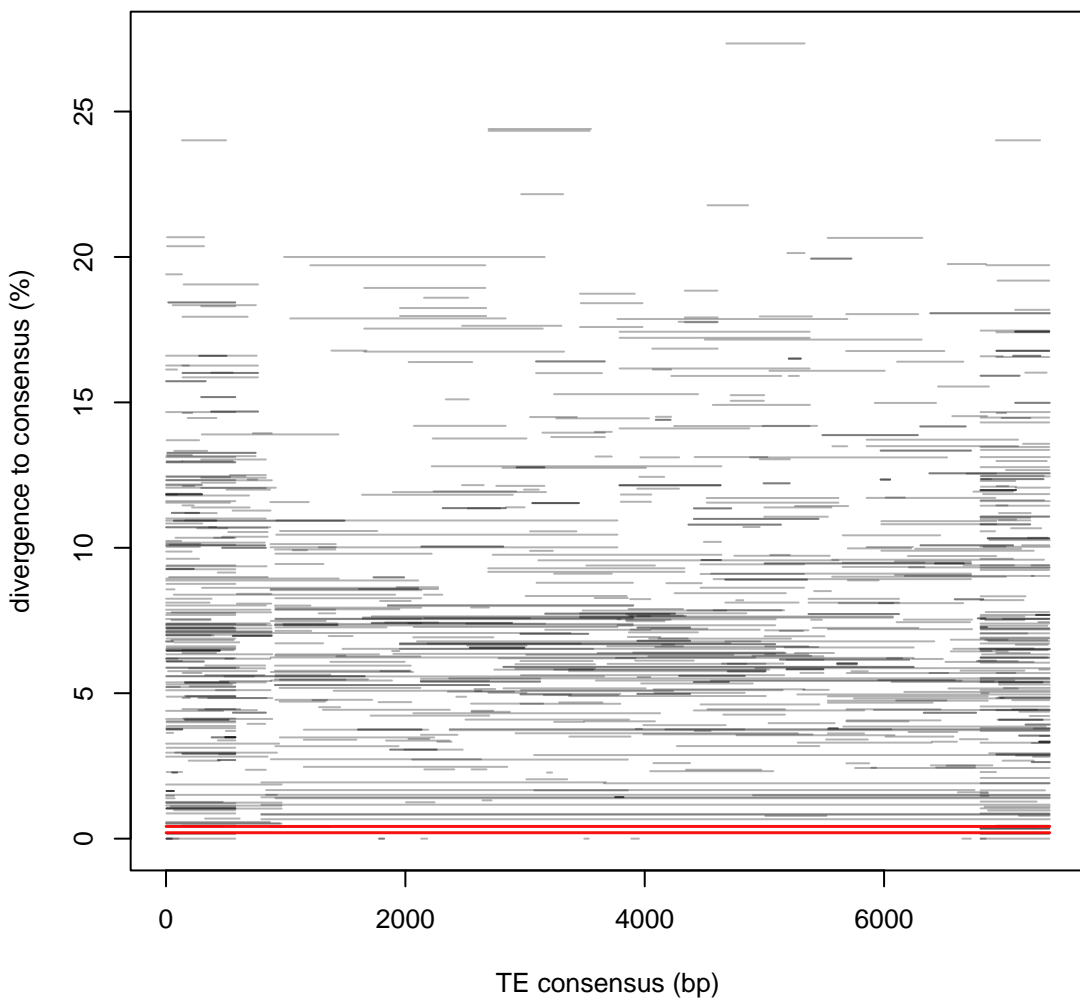
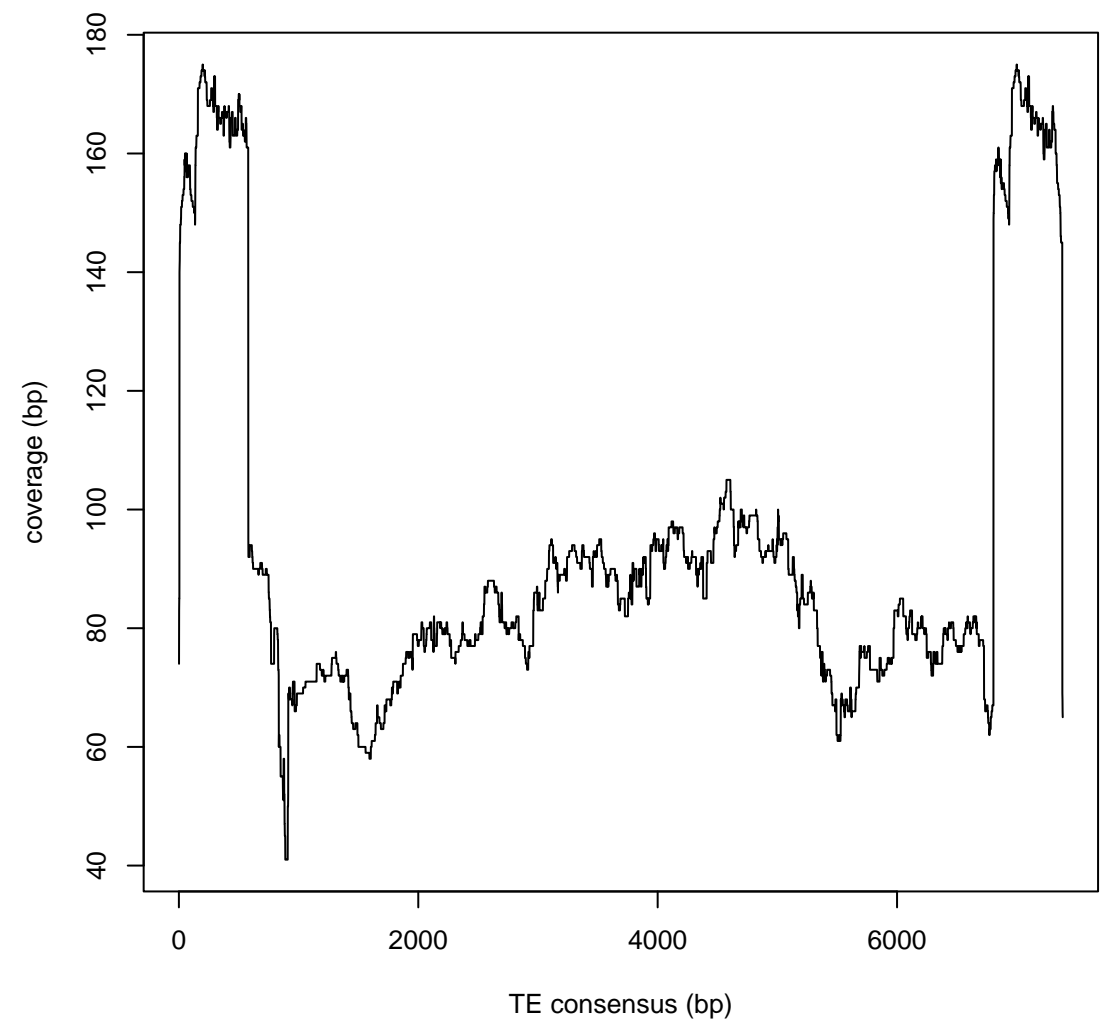


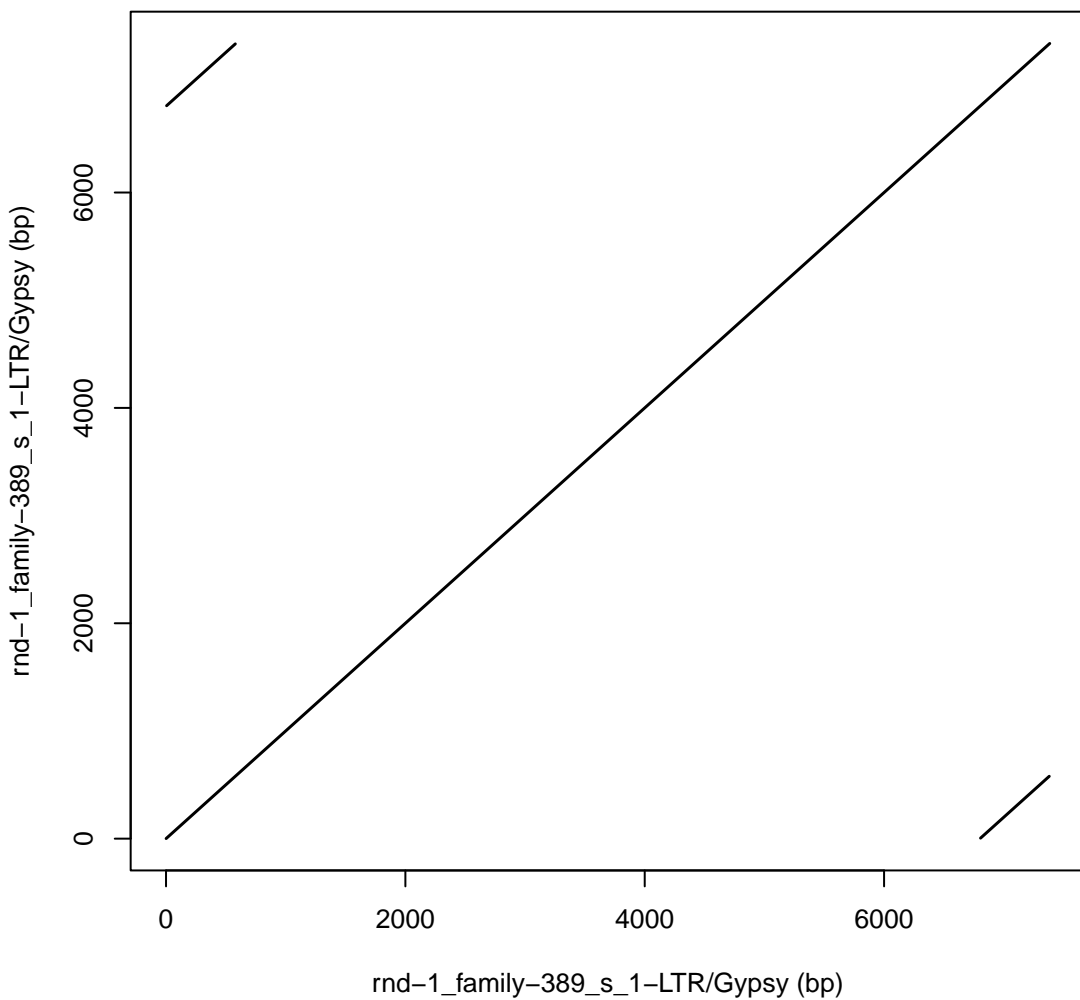
TE: rnd-1\_family-389\_s\_1-LTR/Gypsy  
consensus size: 7384bp; fragments: 962; full length: 2 (>=6645.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

