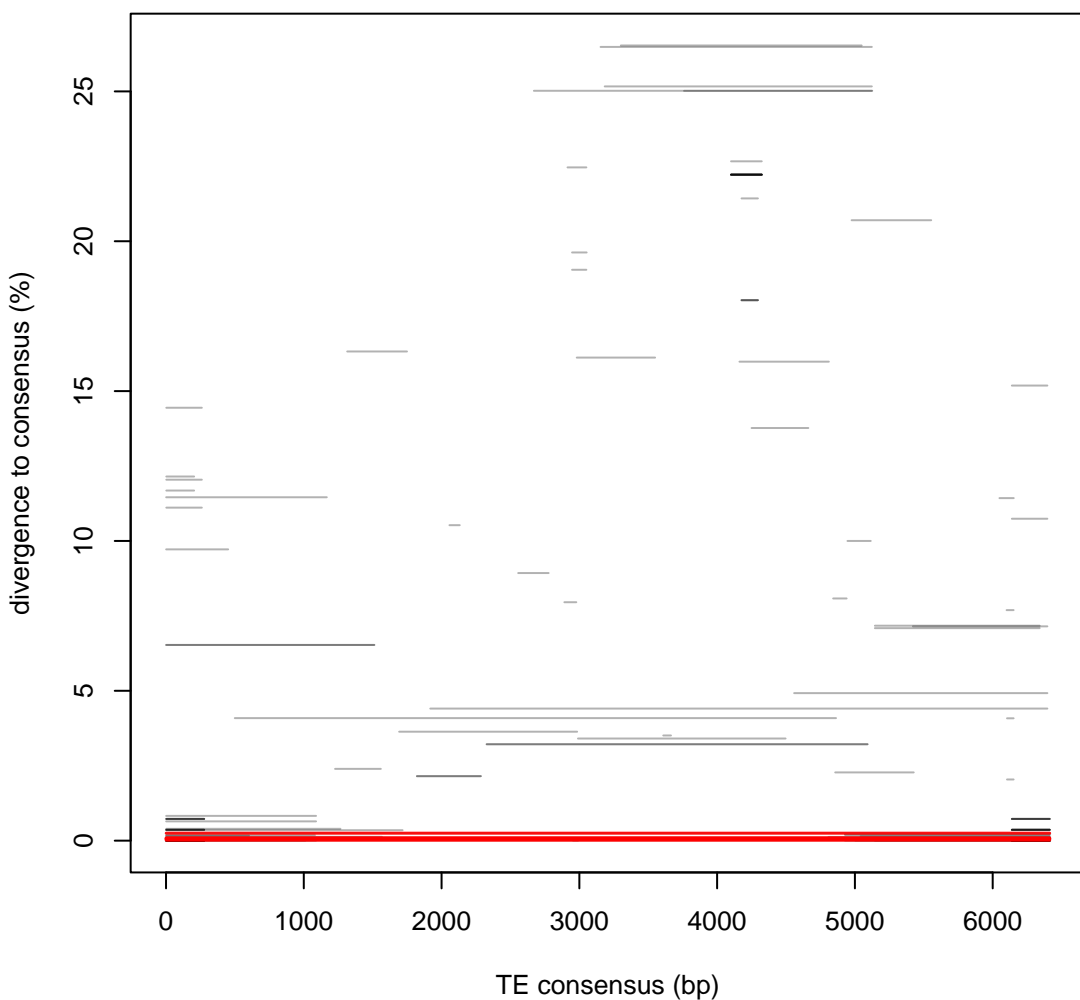
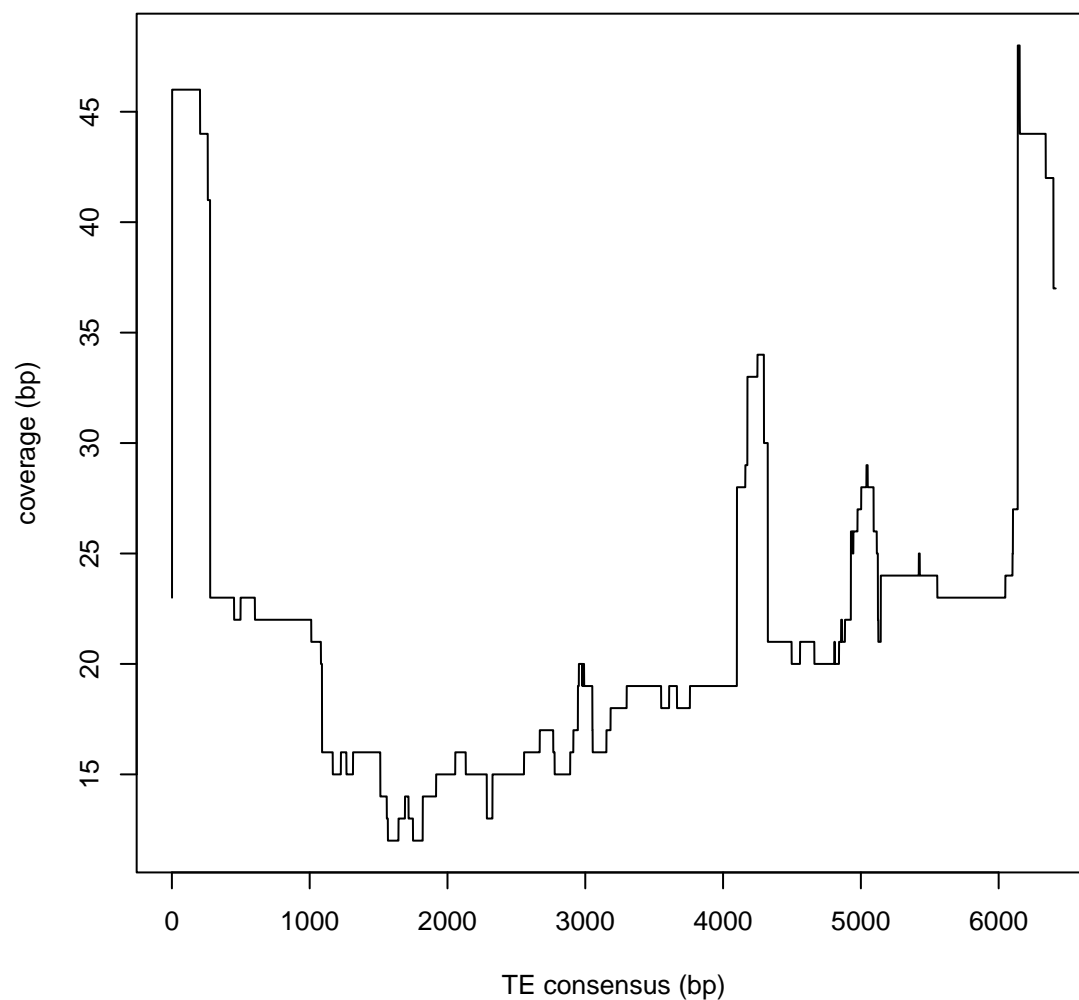


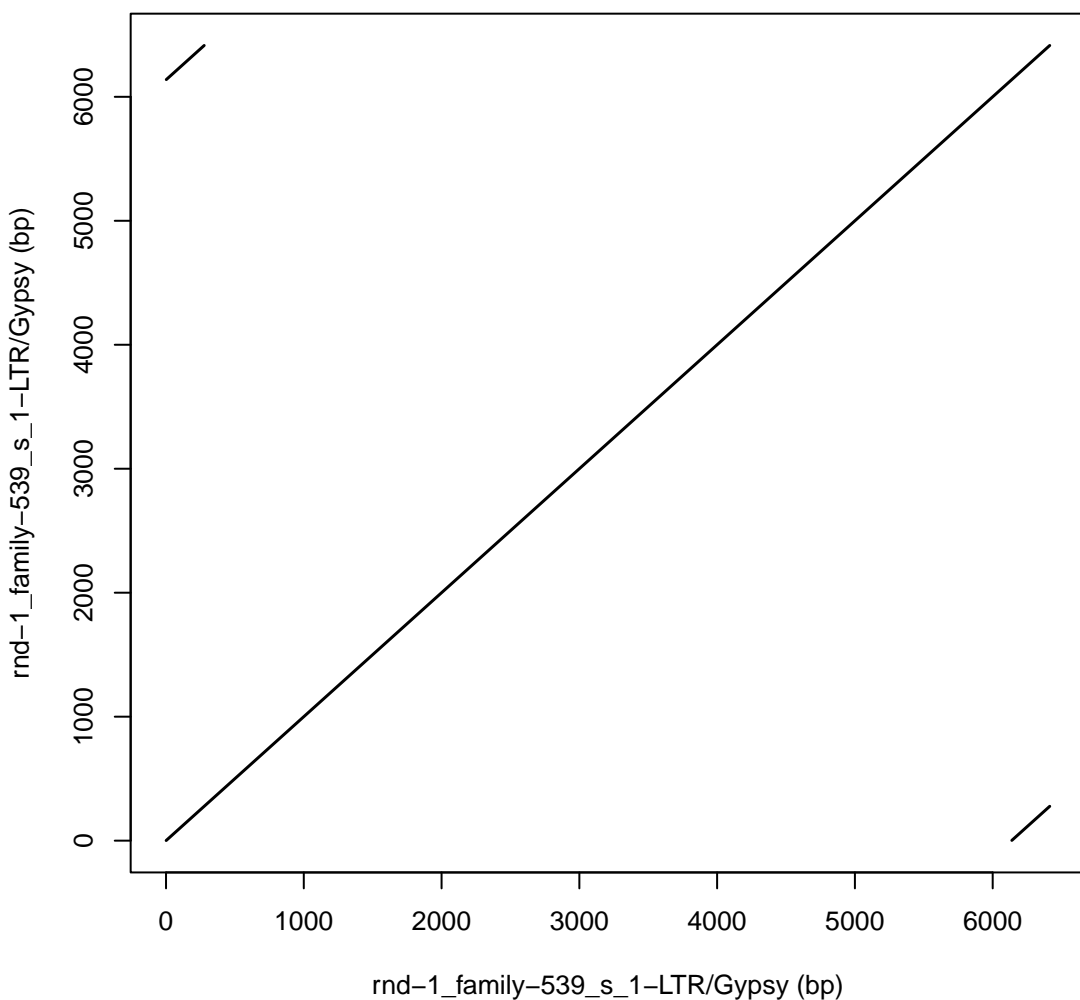
TE: rnd-1\_family-539\_s\_1-LTR/Gypsy  
consensus size: 6414bp; fragments: 128; full length: 8 (>=5772.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

