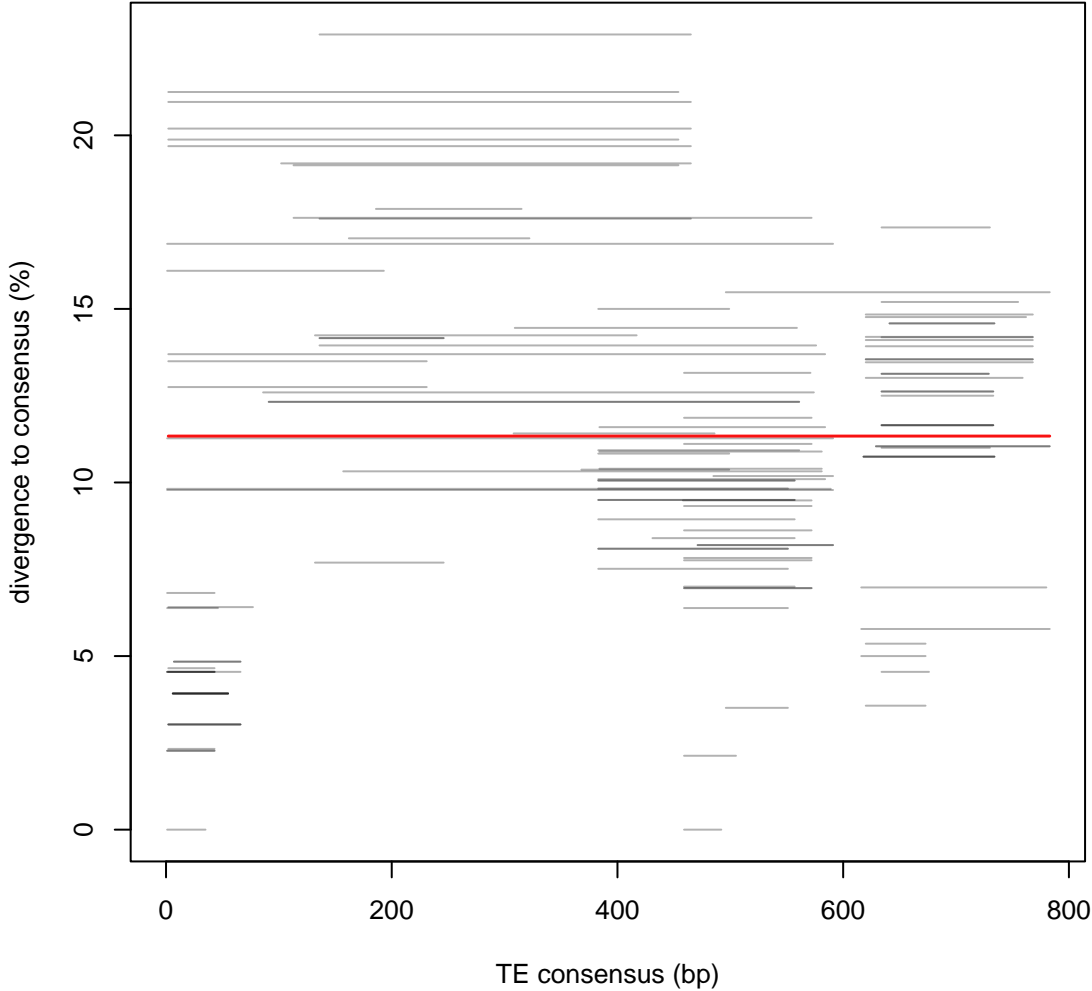
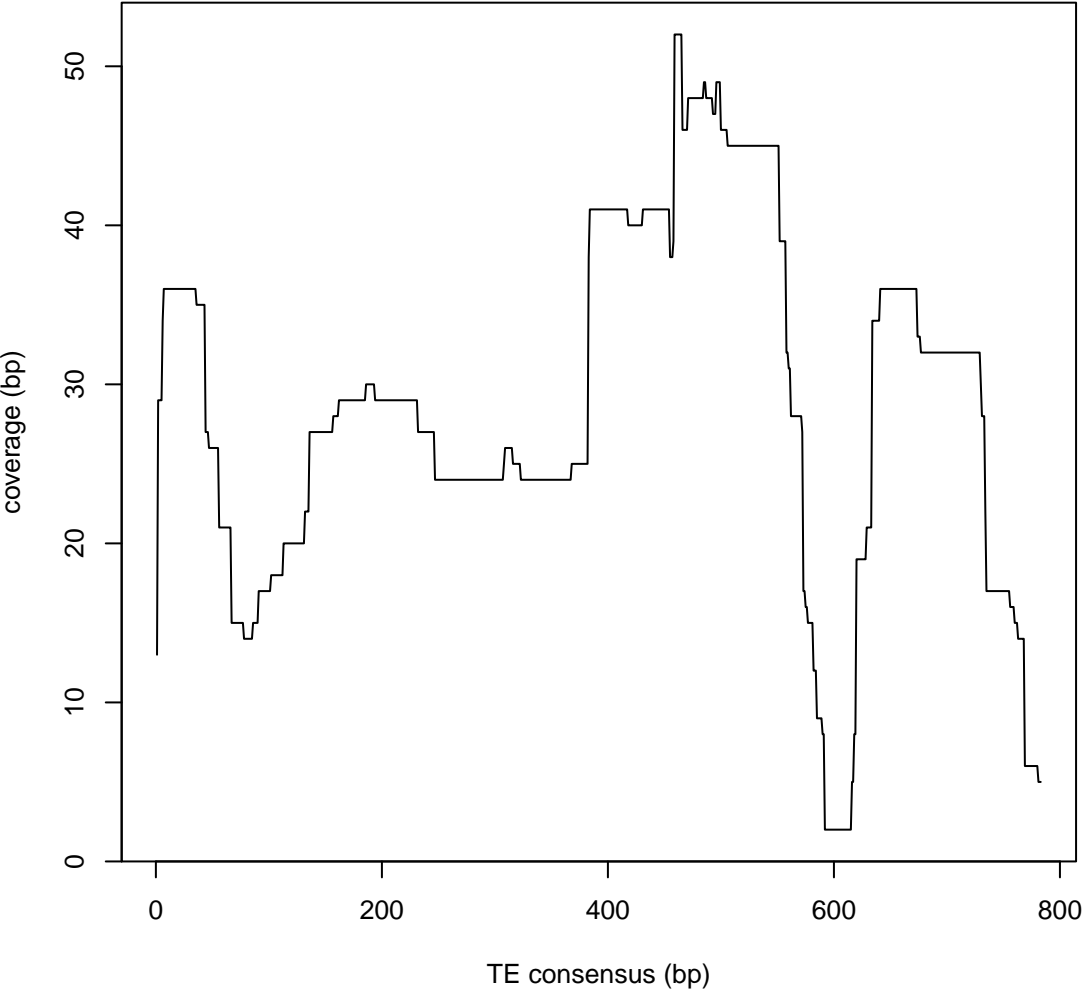


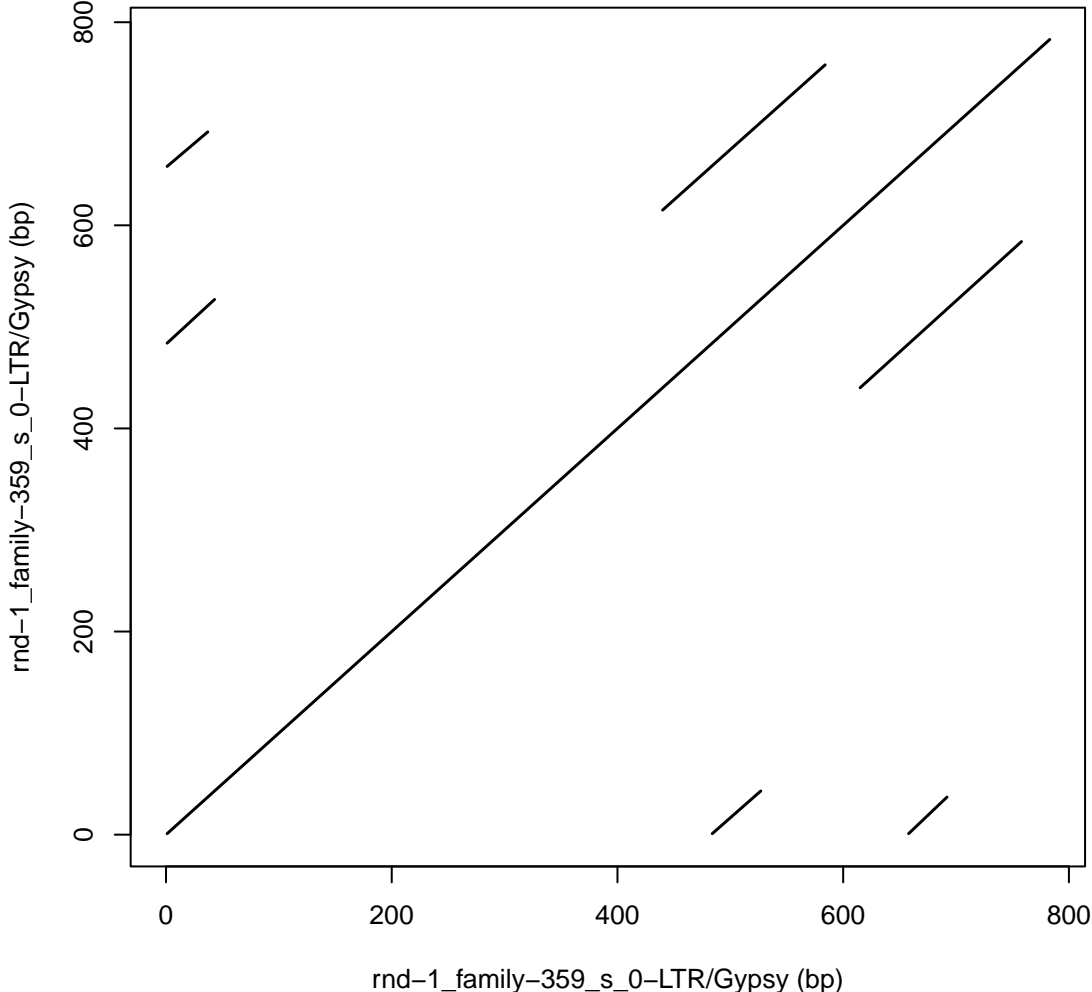
TE: rnd-1\_family-359\_s\_0-LTR/Gypsy  
consensus size: 783bp; fragments: 125; full length: 1 (>=704.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

