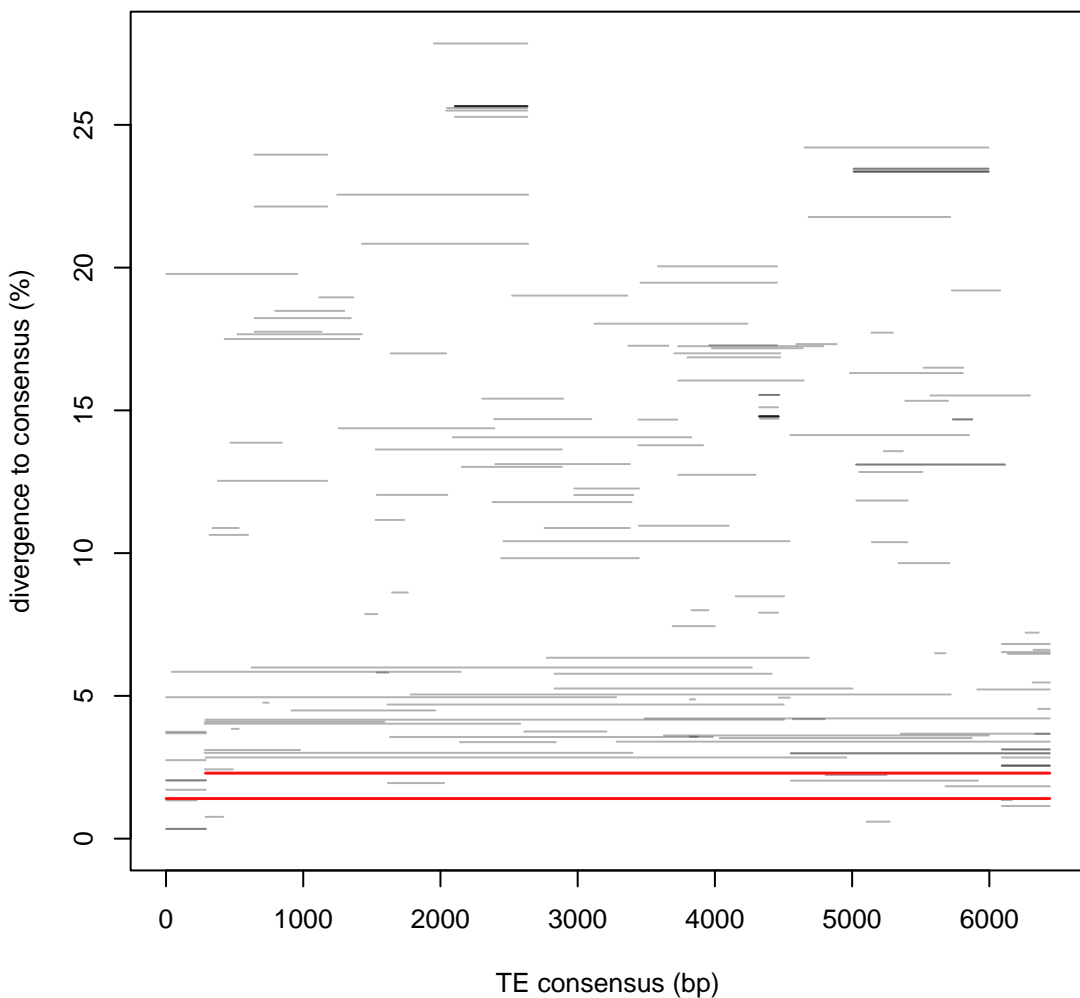
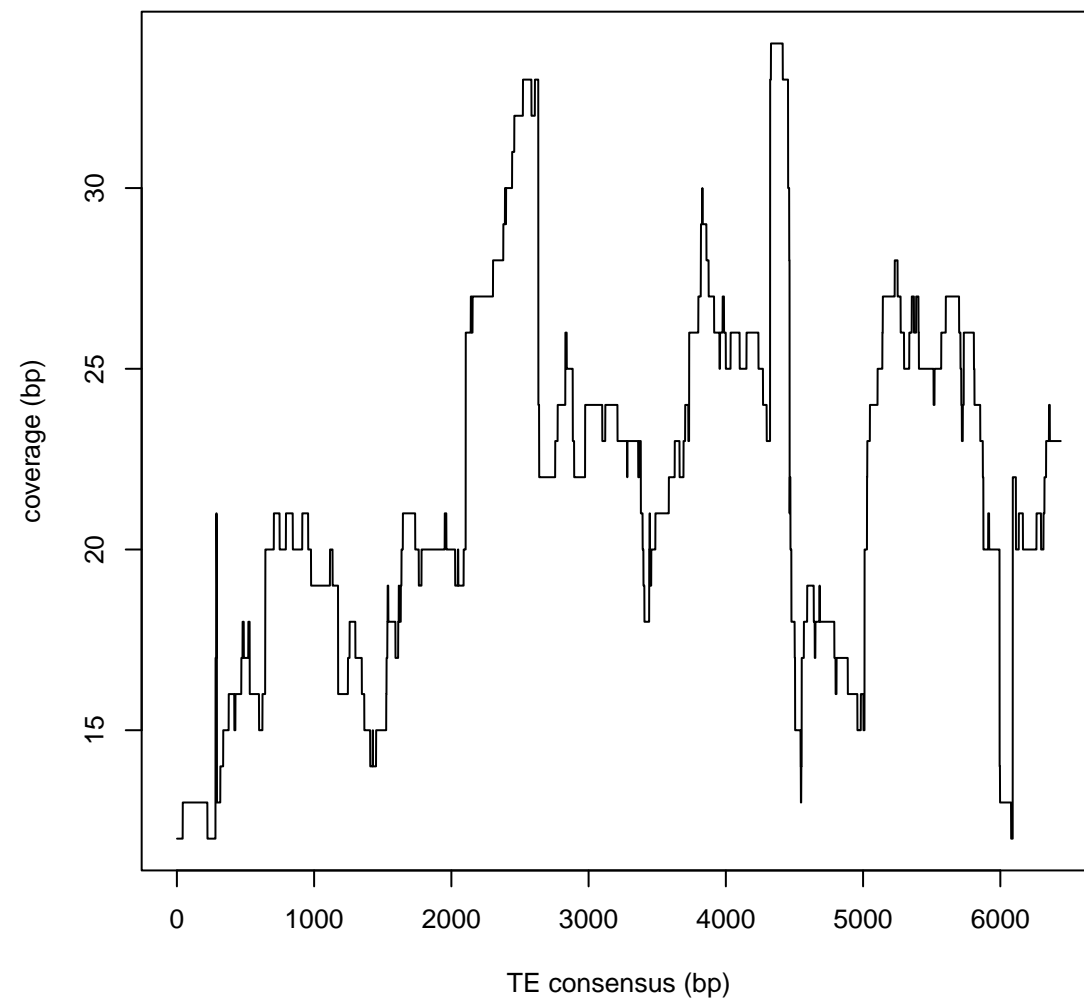


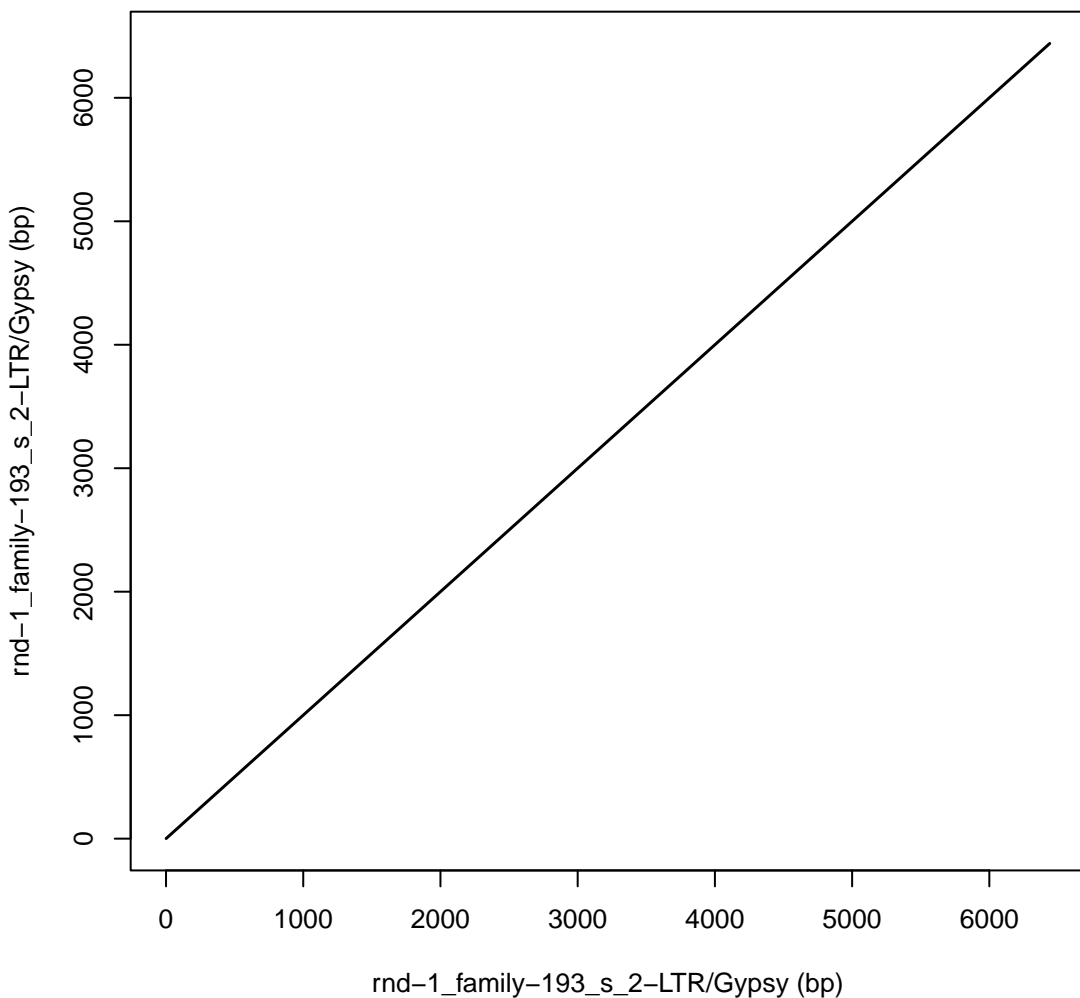
TE: rnd-1\_family-193\_s\_2-LTR/Gypsy  
consensus size: 6440bp; fragments: 163; full length: 2 (>=5796bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

