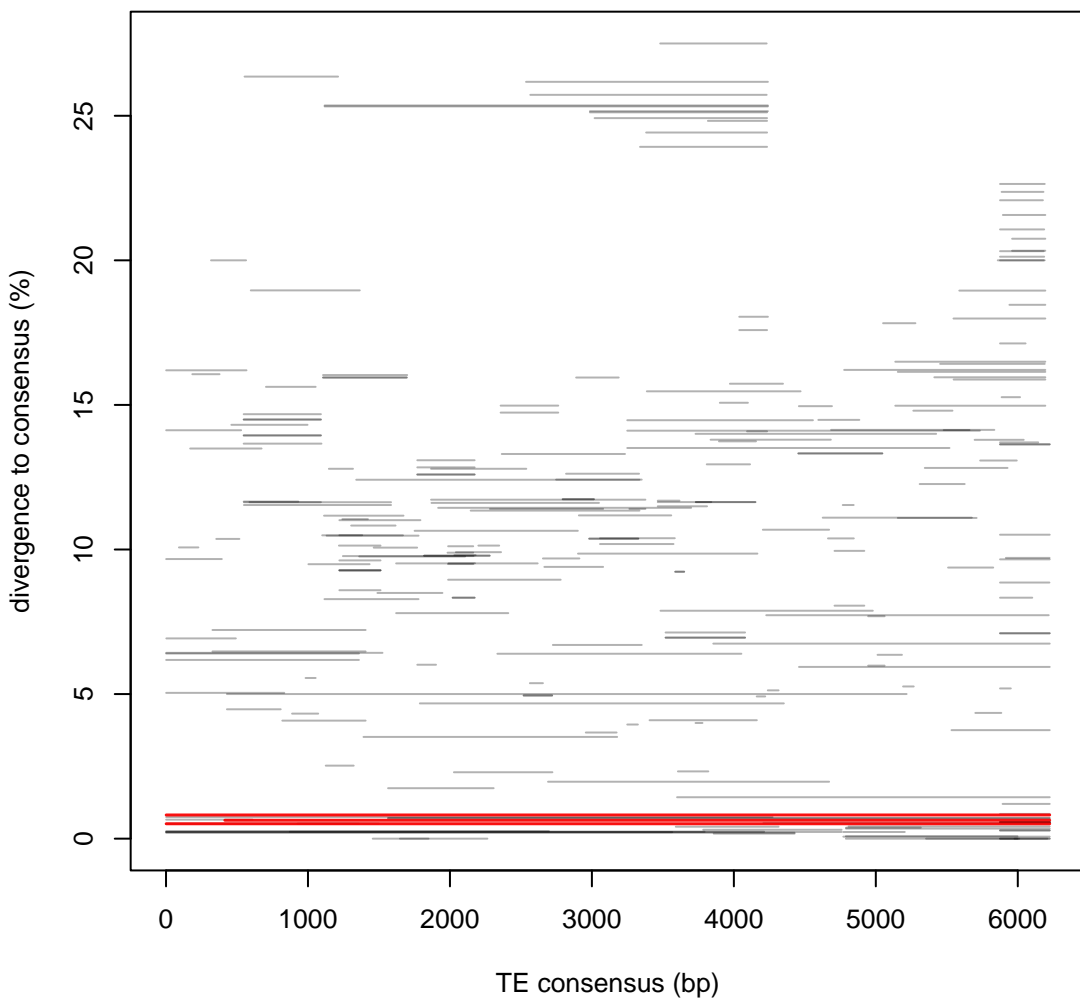
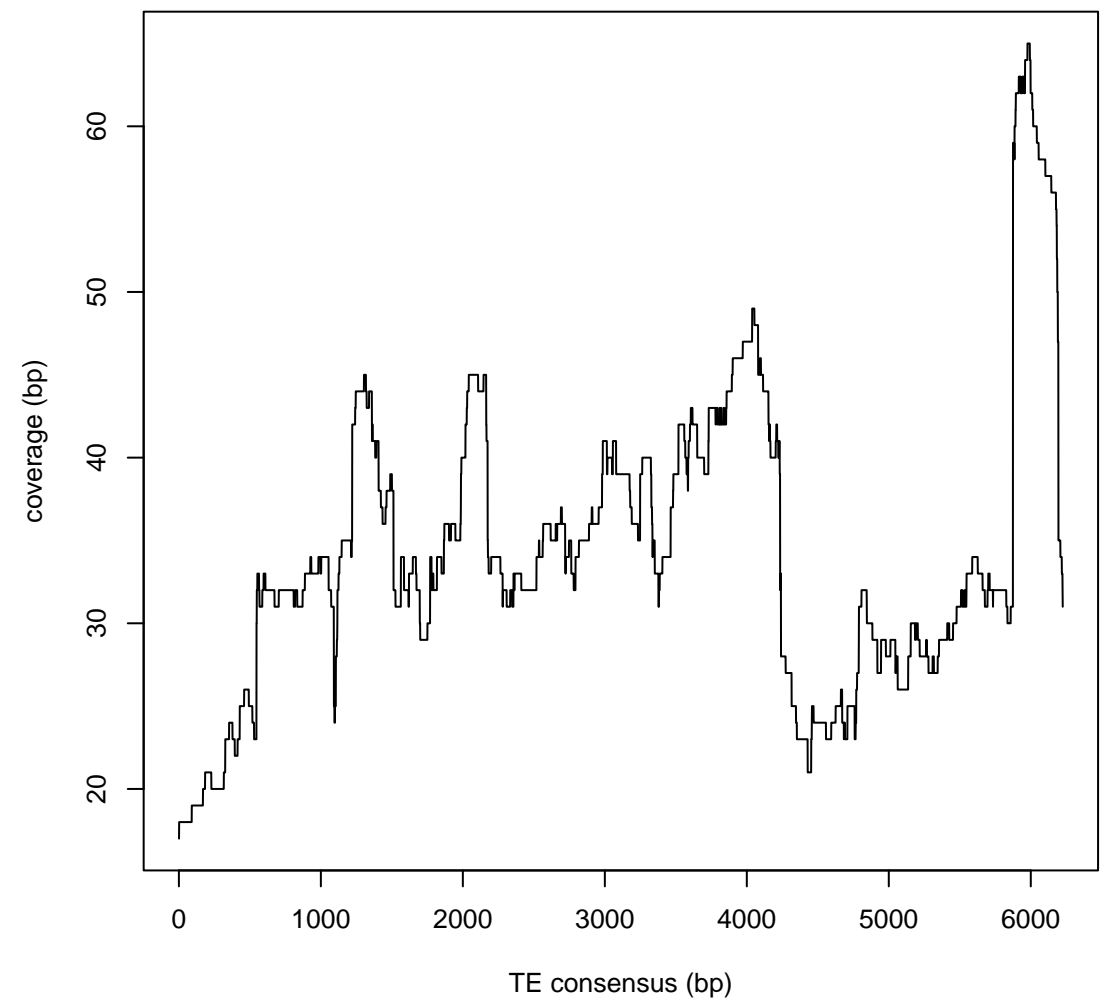


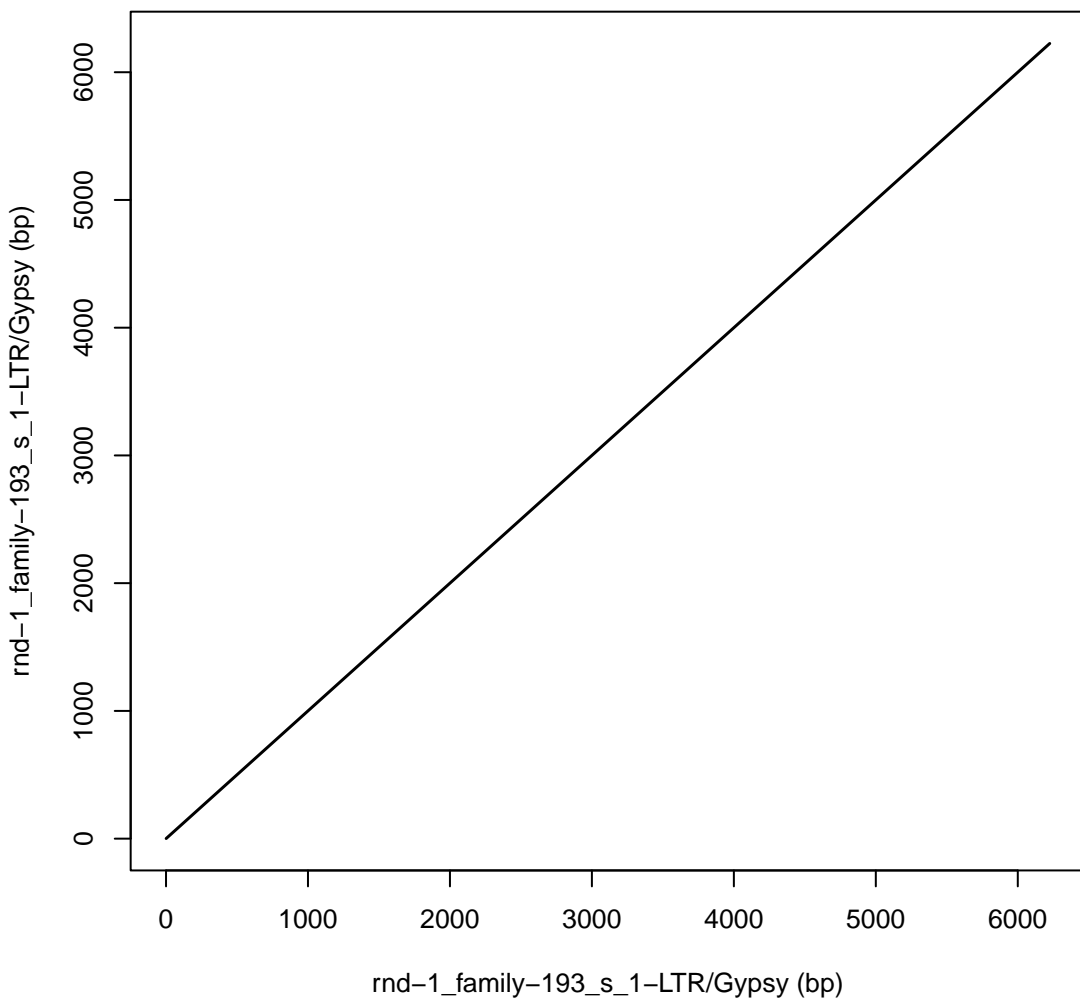
TE: rnd-1\_family-193\_s\_1-LTR/Gypsy  
consensus size: 6225bp; fragments: 260; full length: 3 (>=5602.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

