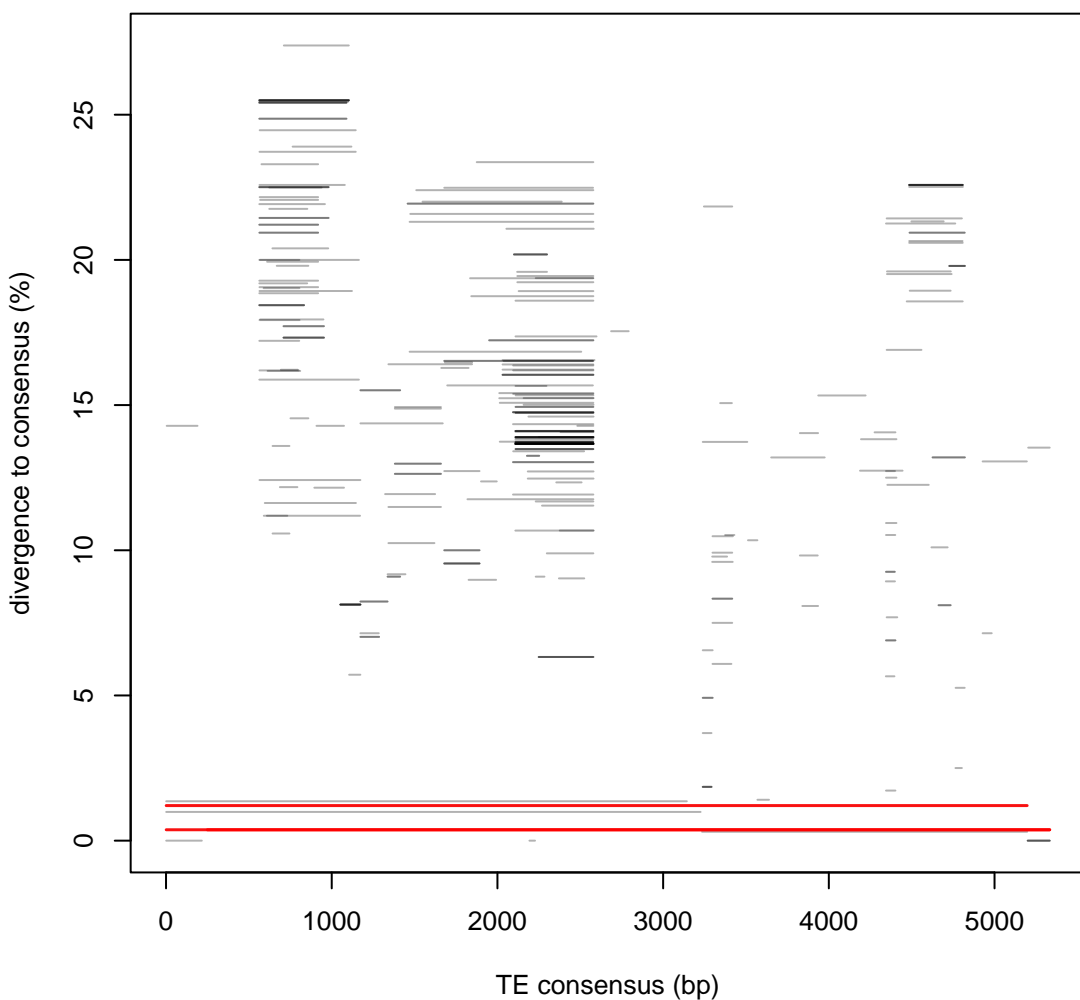
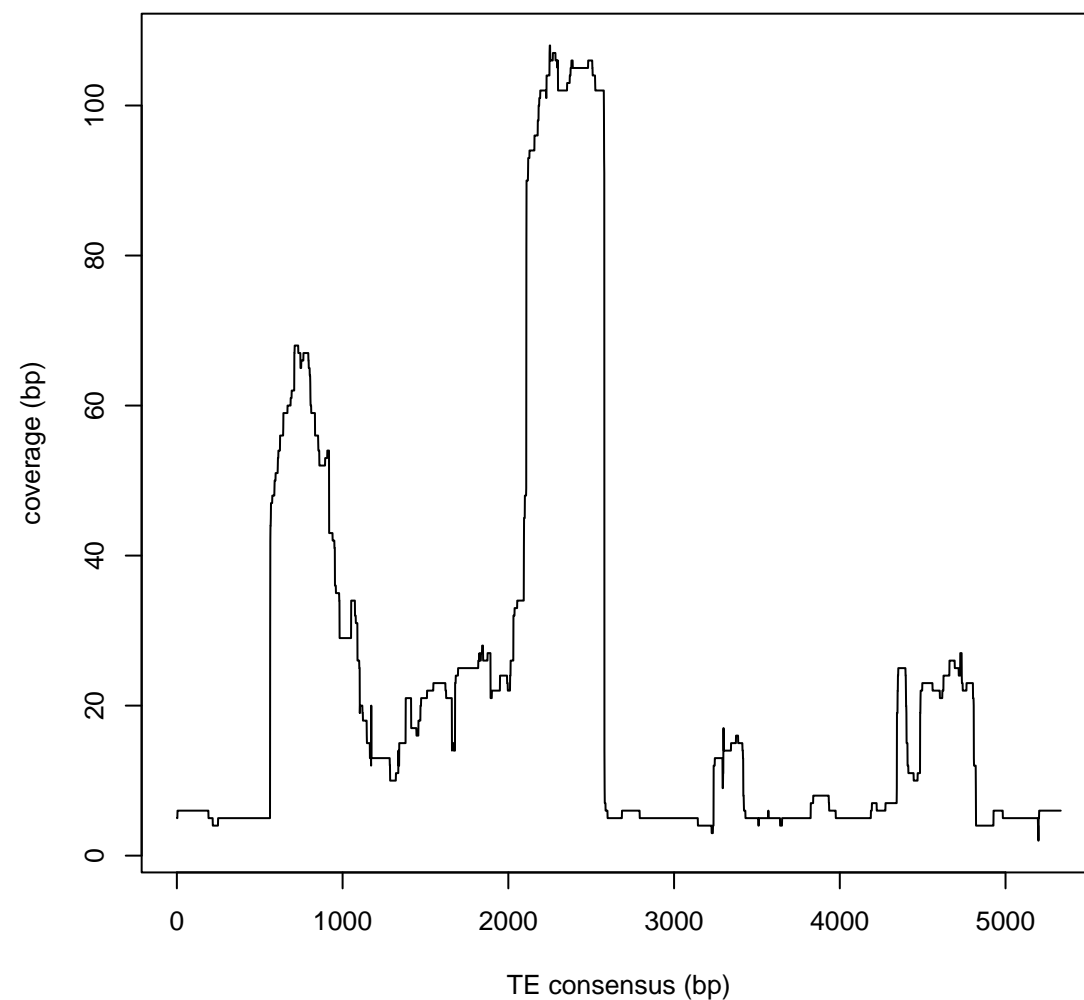


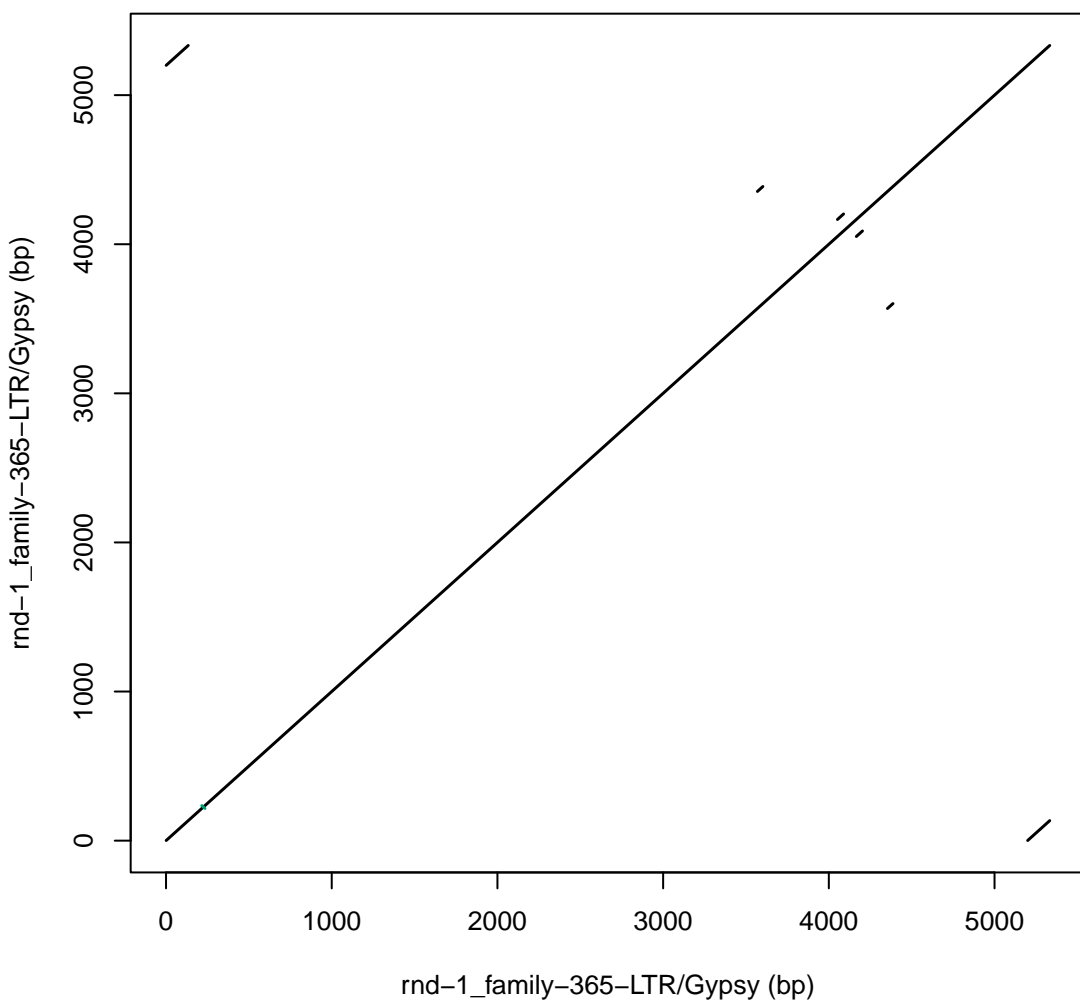
TE: rnd-1\_family-365-LTR/Gypsy  
consensus size: 5333bp; fragments: 300; full length: 3 (>=4799.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

