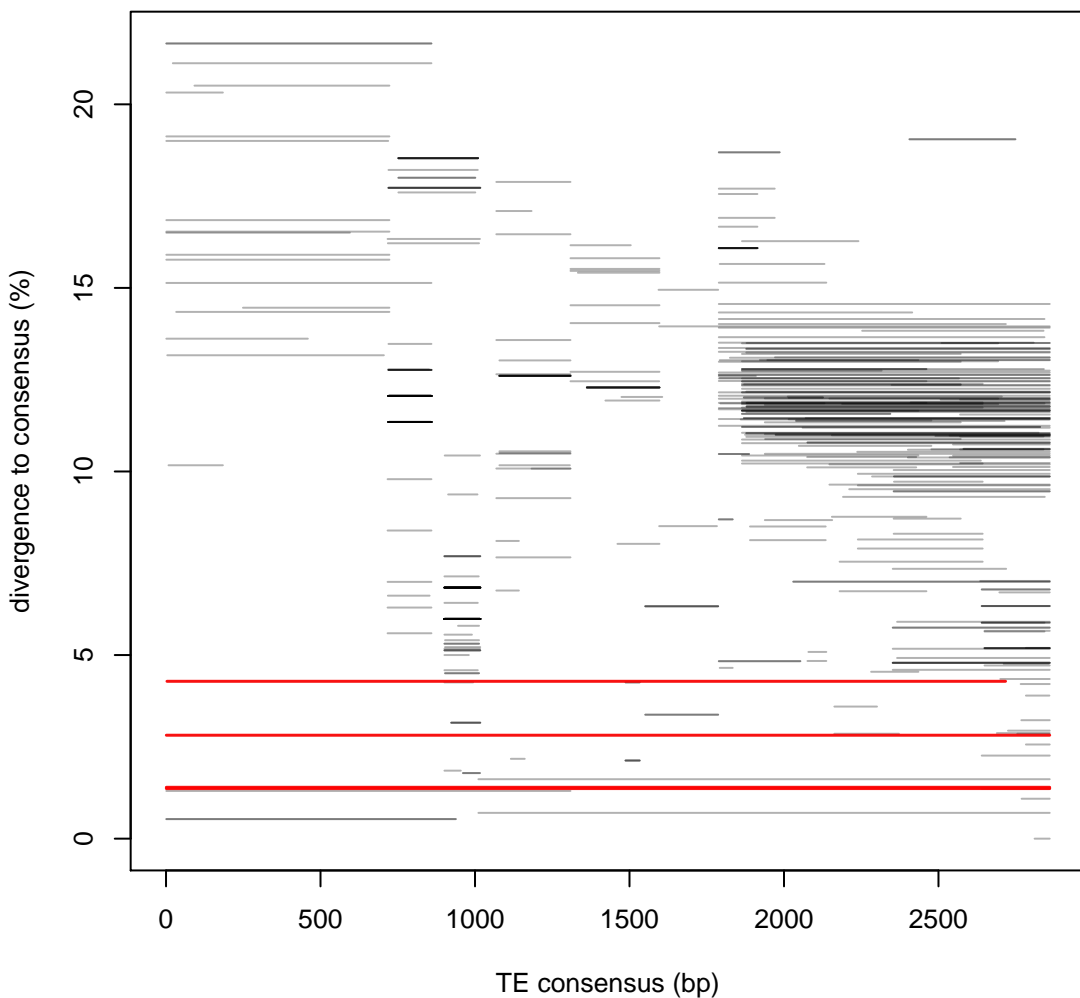
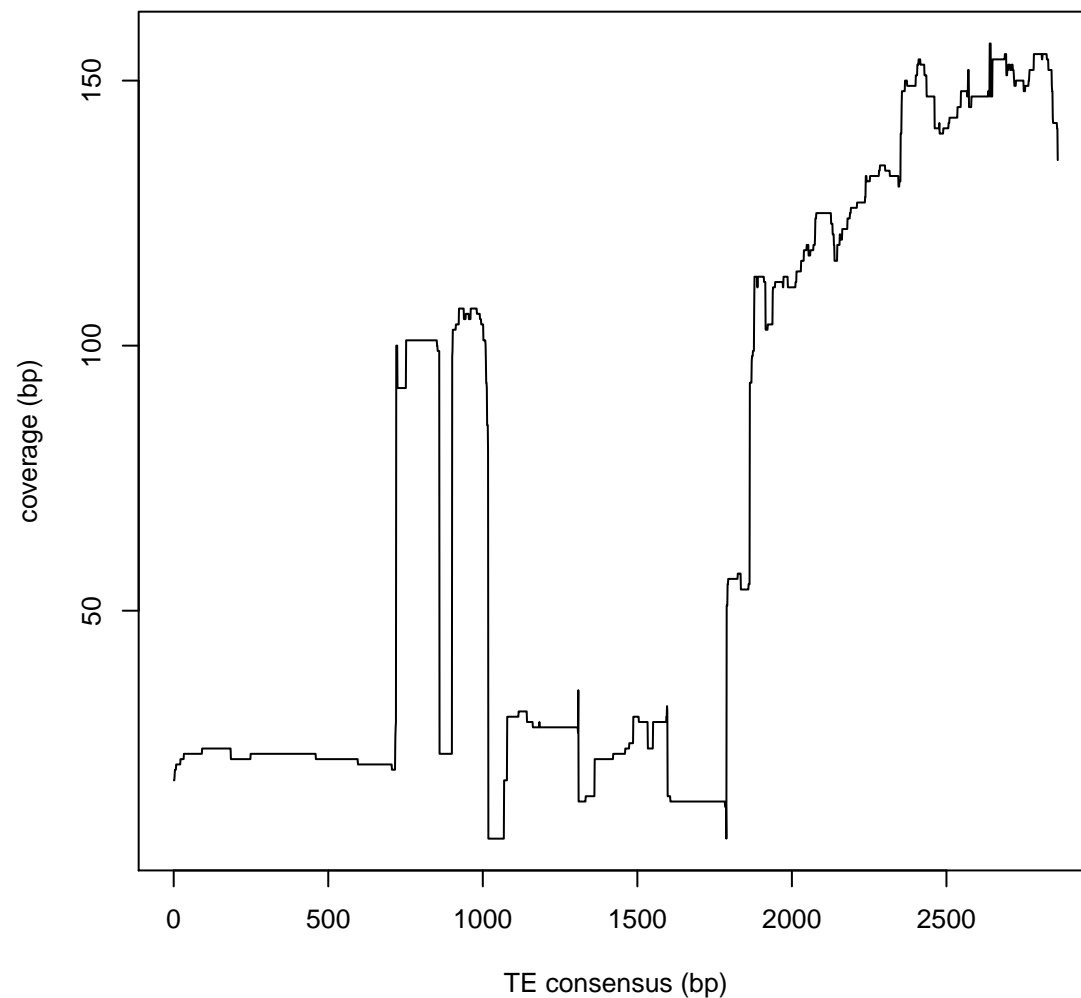


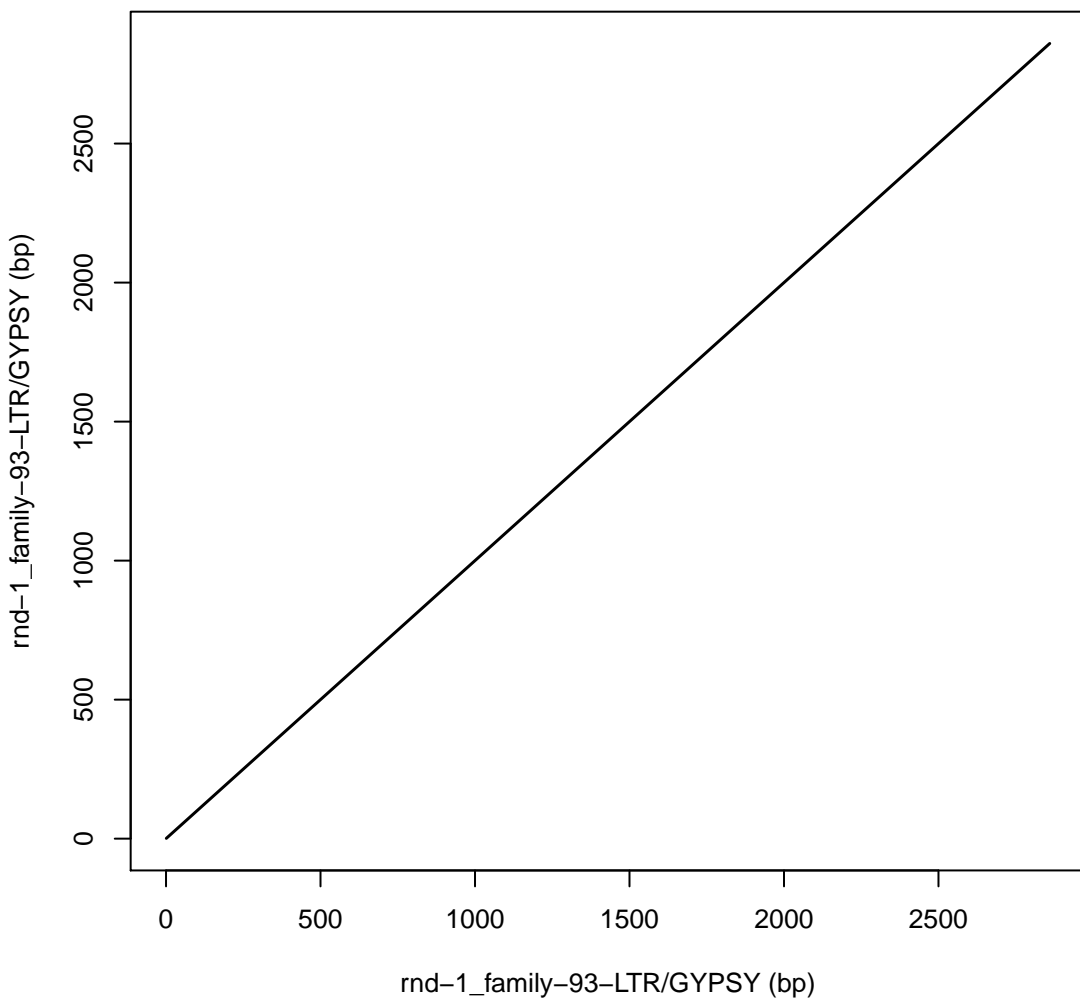
TE: rnd-1_family-93-LTR/GYPHY
consensus size: 2860bp; fragments: 492; full length: 4 (>=2574bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

