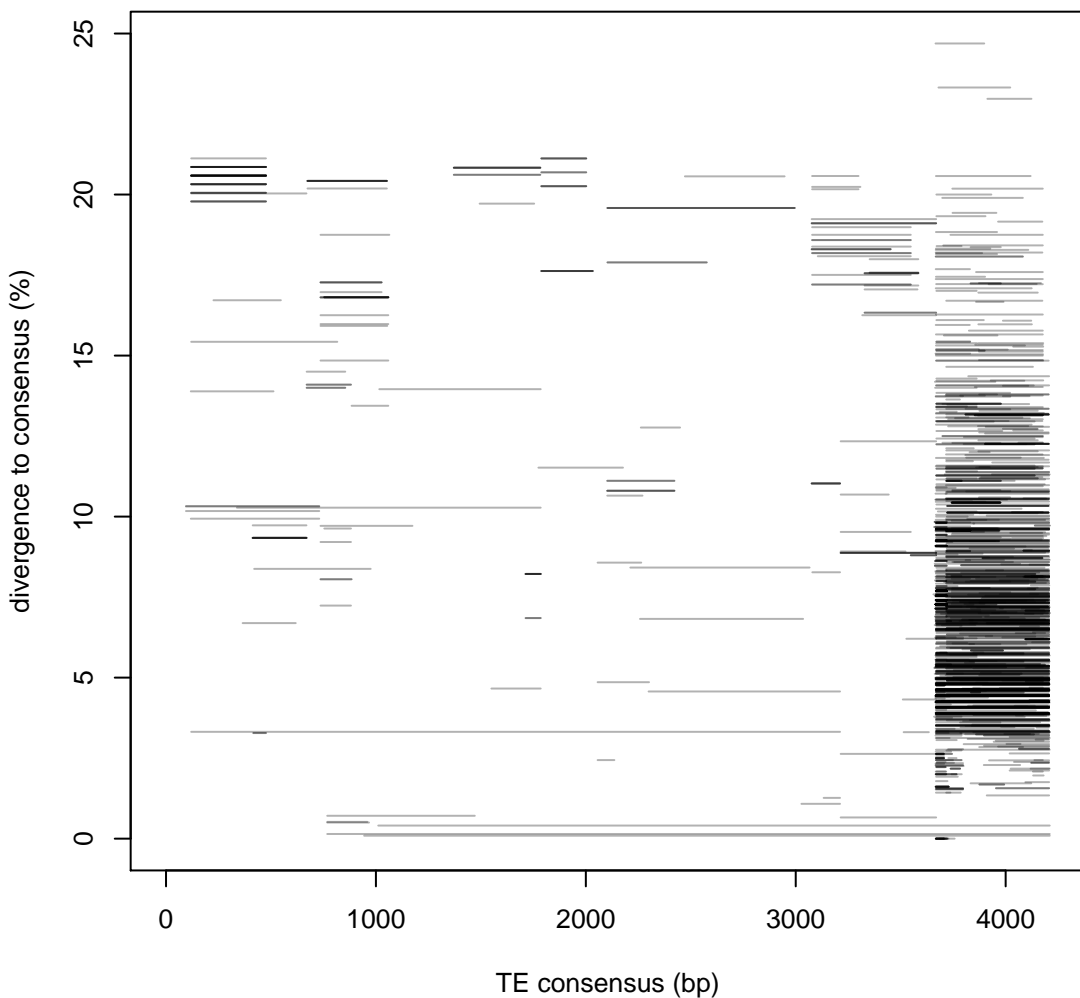
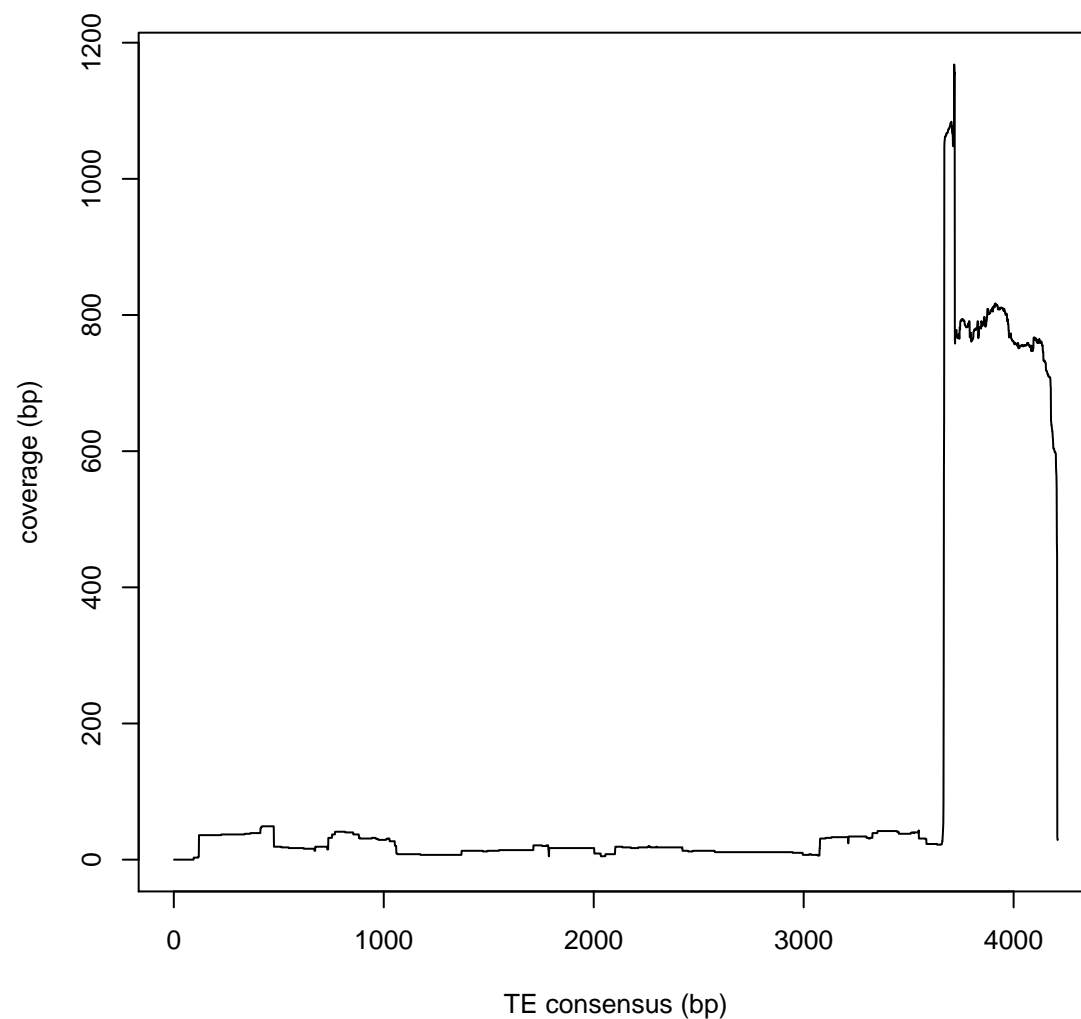


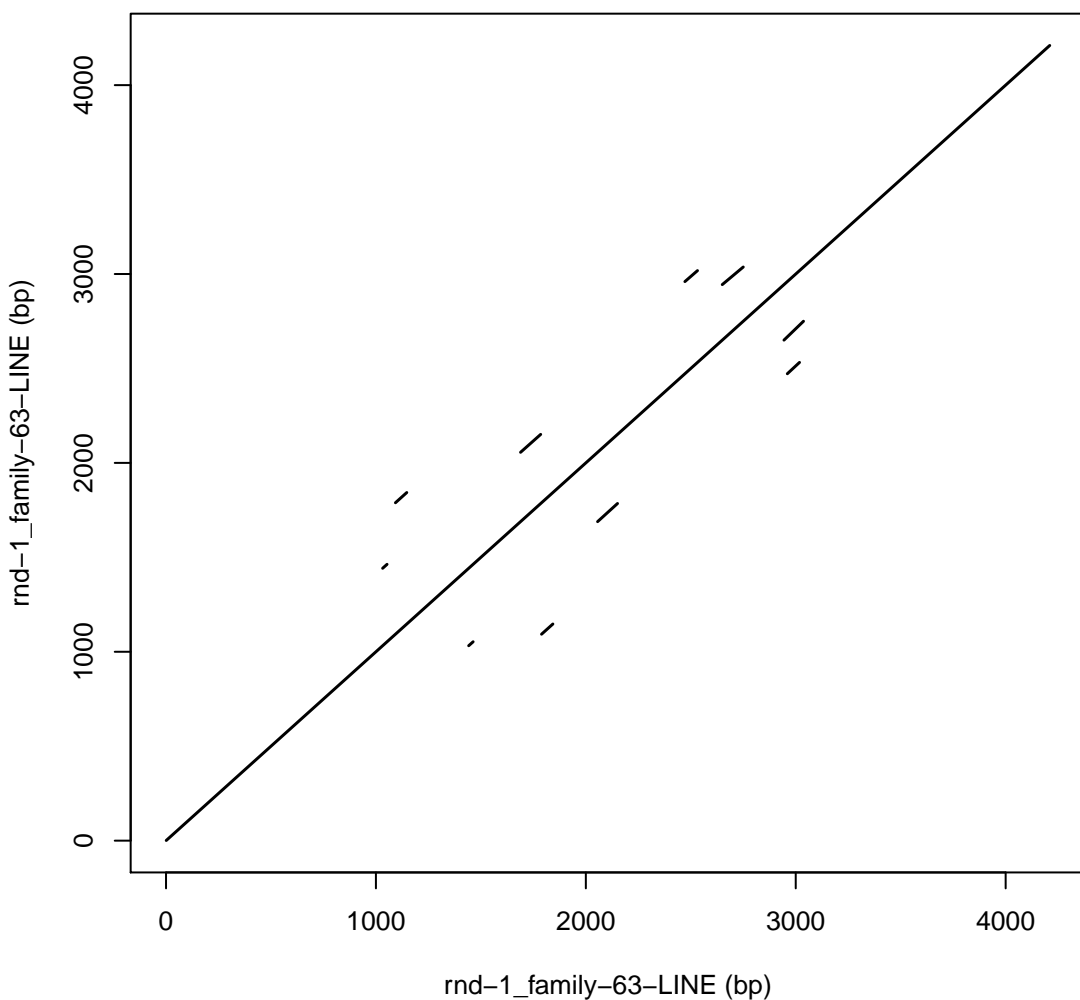
TE: rnd-1_family-63-LINE
consensus size: 4210bp; fragments: 1810; full length: 0 (>=3789bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

