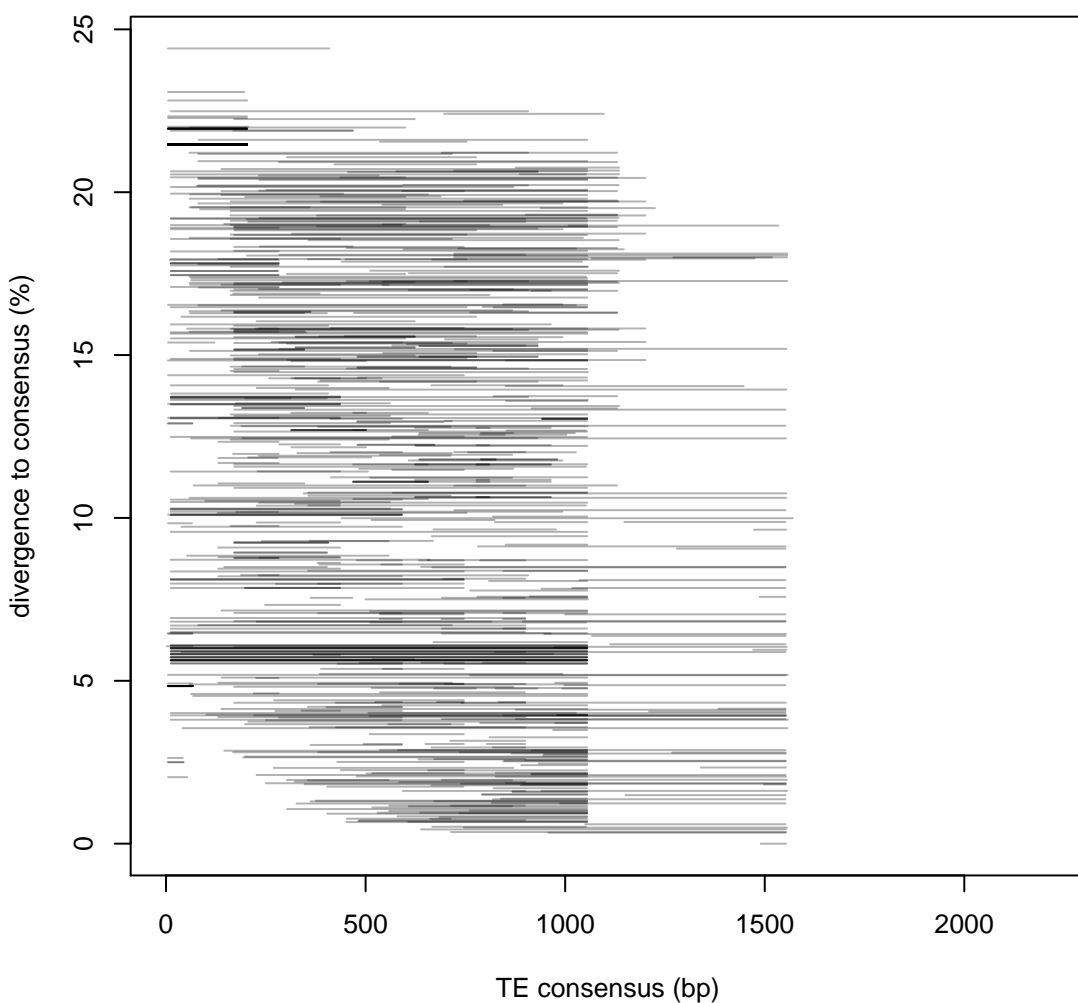
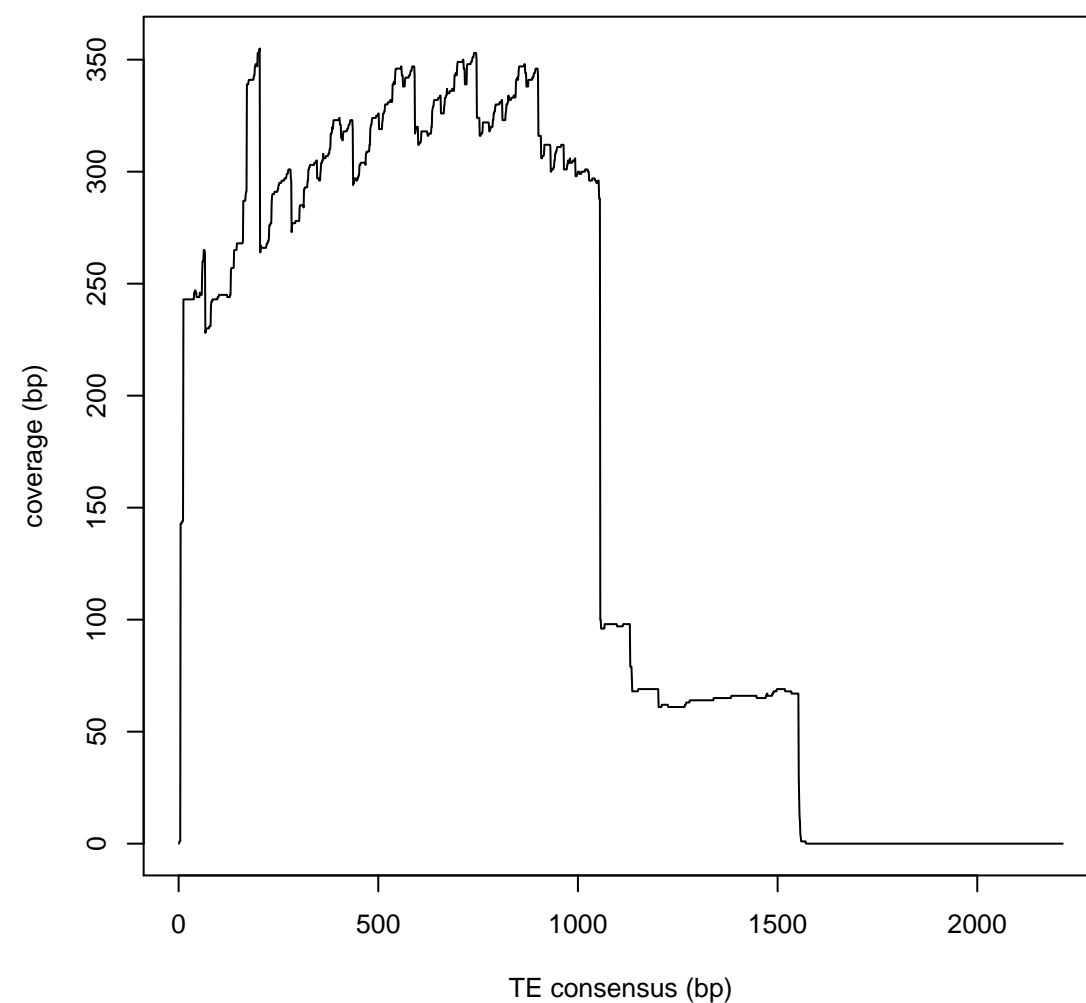


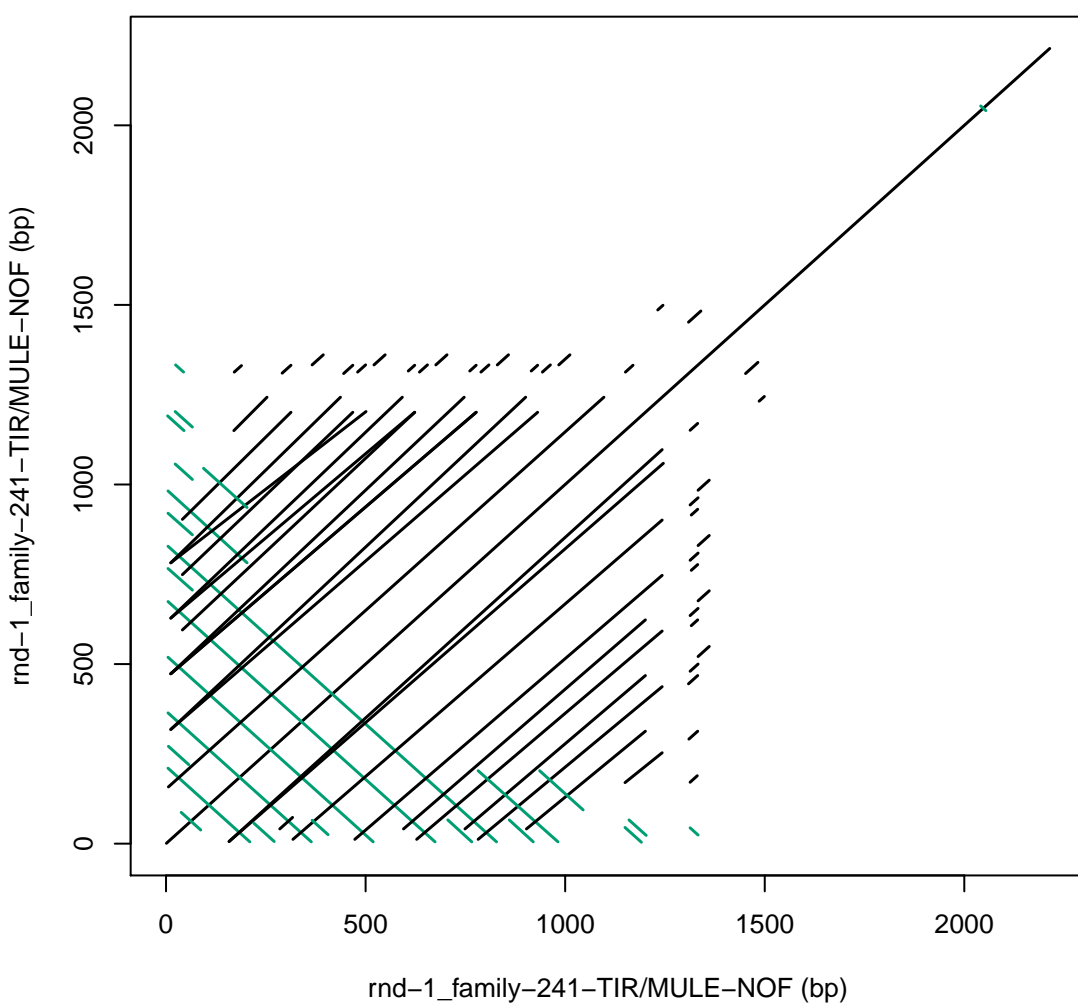
TE: rnd-1_family-241-TIR/MULE-NOF
consensus size: 2214bp; fragments: 803; full length: 0 (>=1992.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

