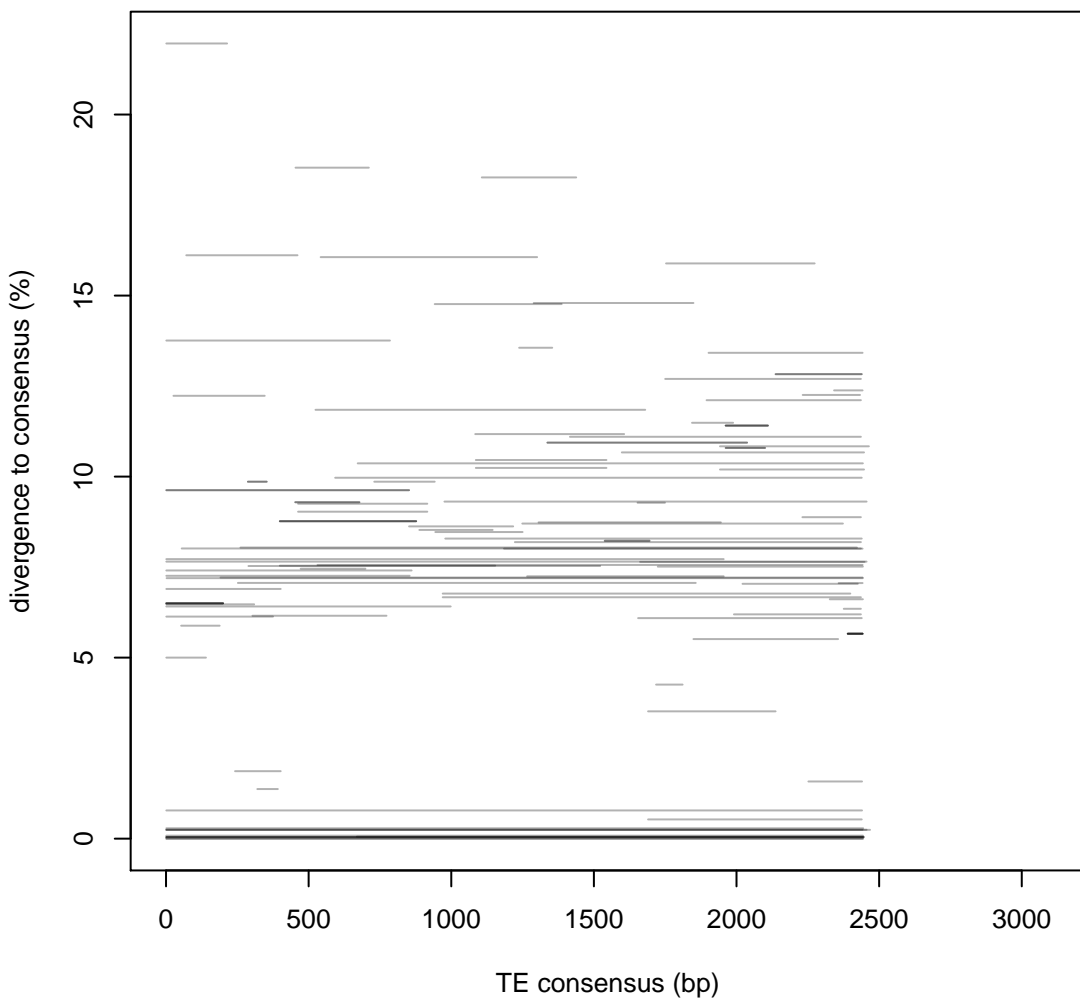
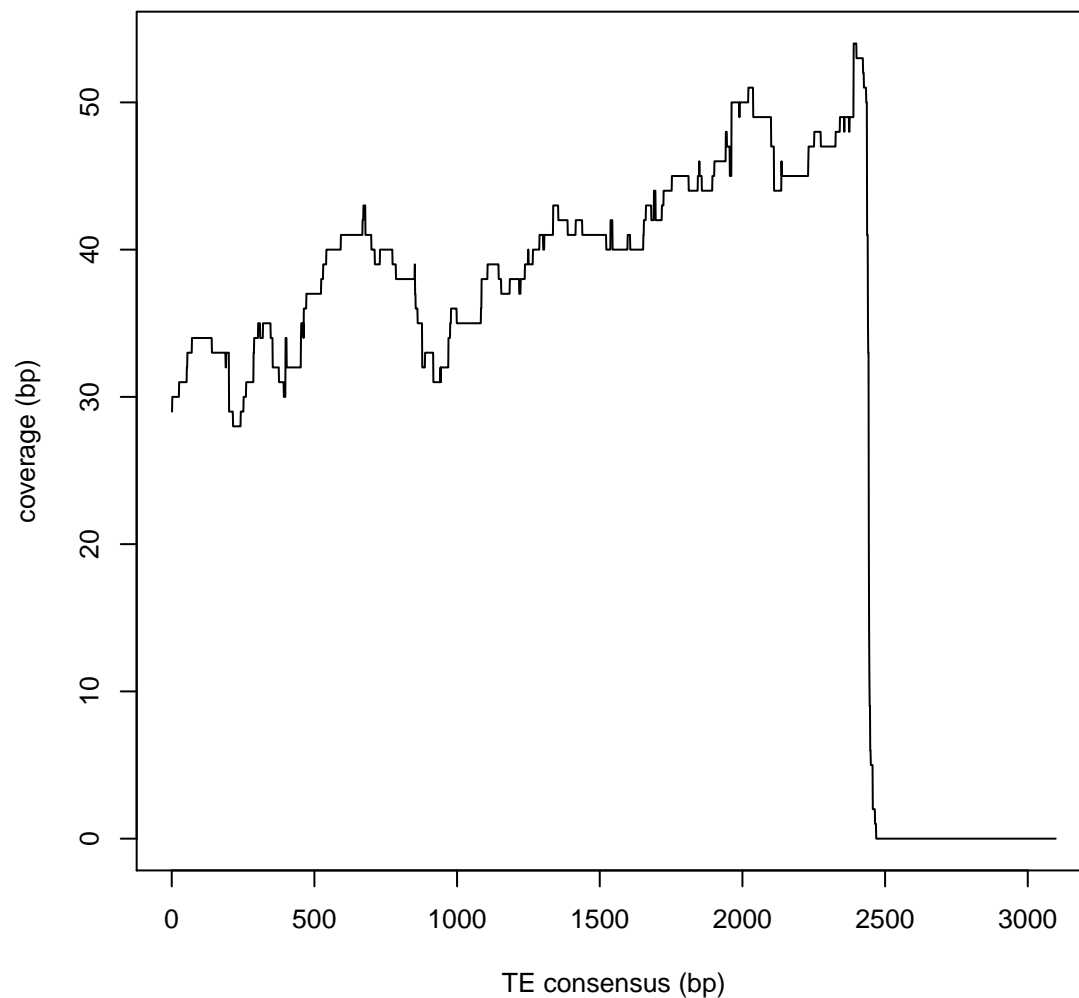


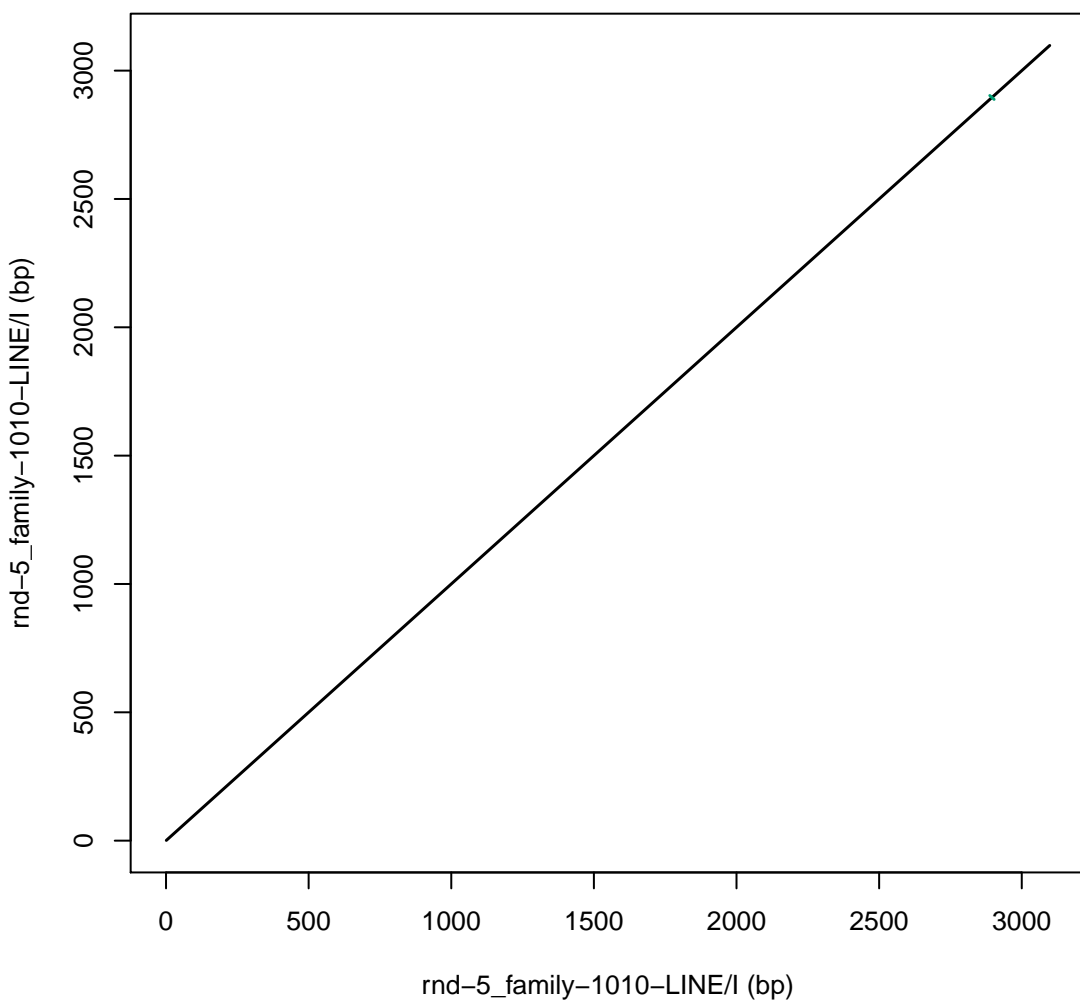
TE: rnd-5_family-1010-LINE/I
consensus size: 3098bp; fragments: 121; full length: 0 (>=2788.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

