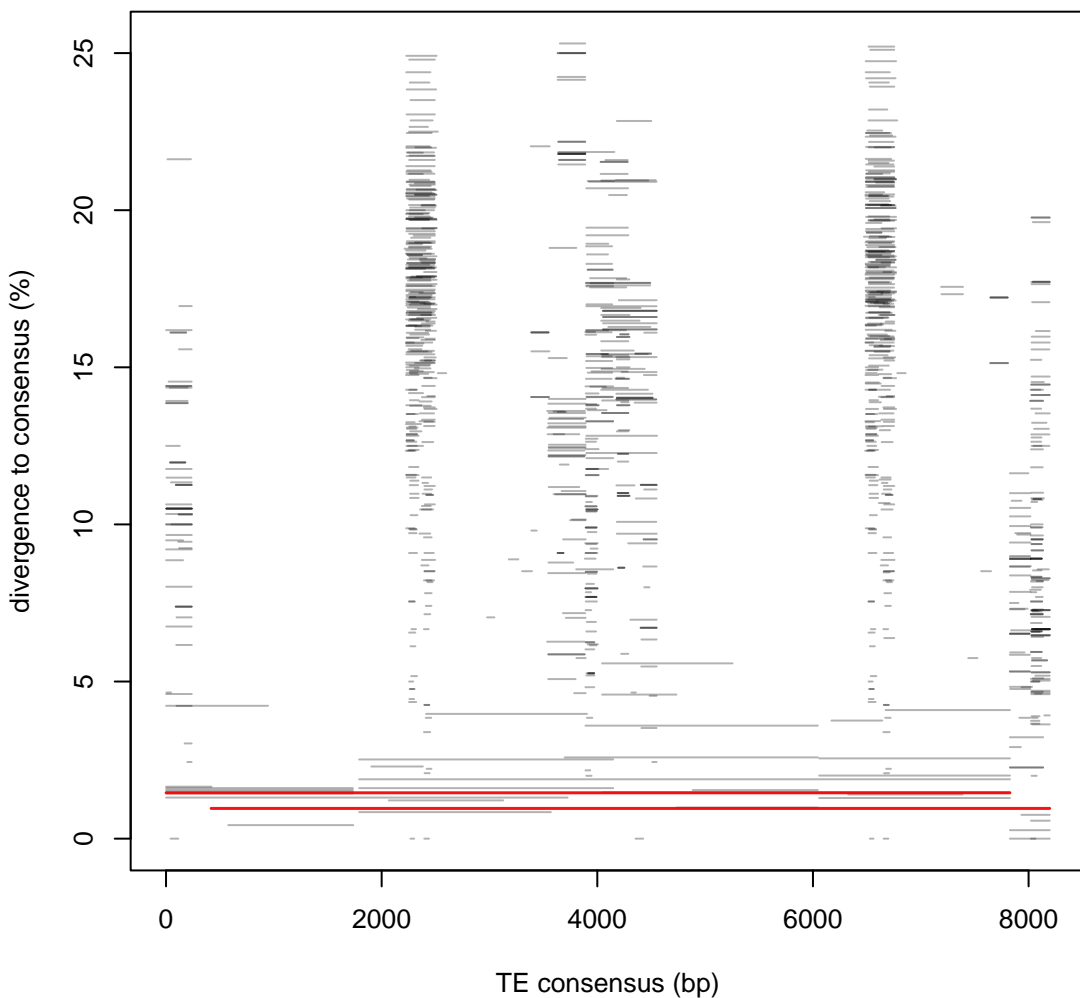
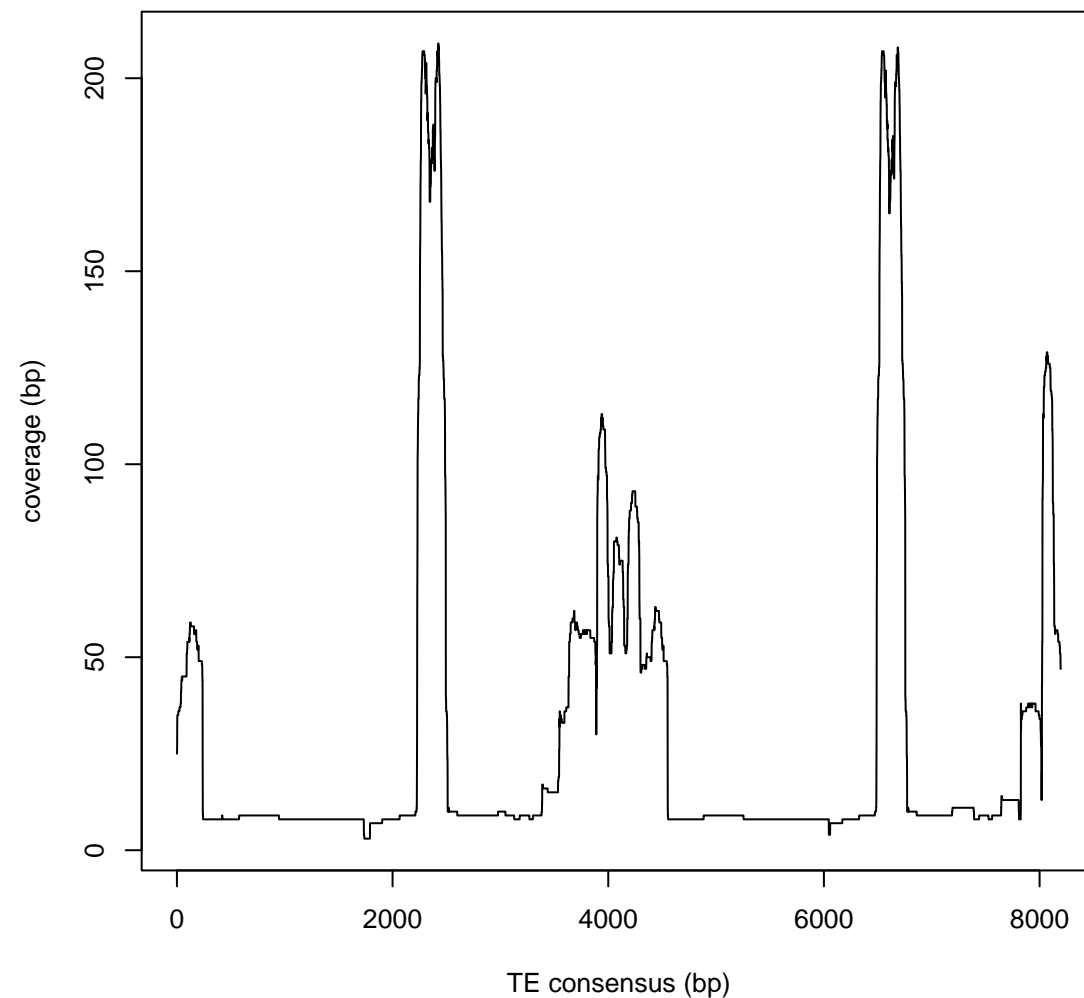


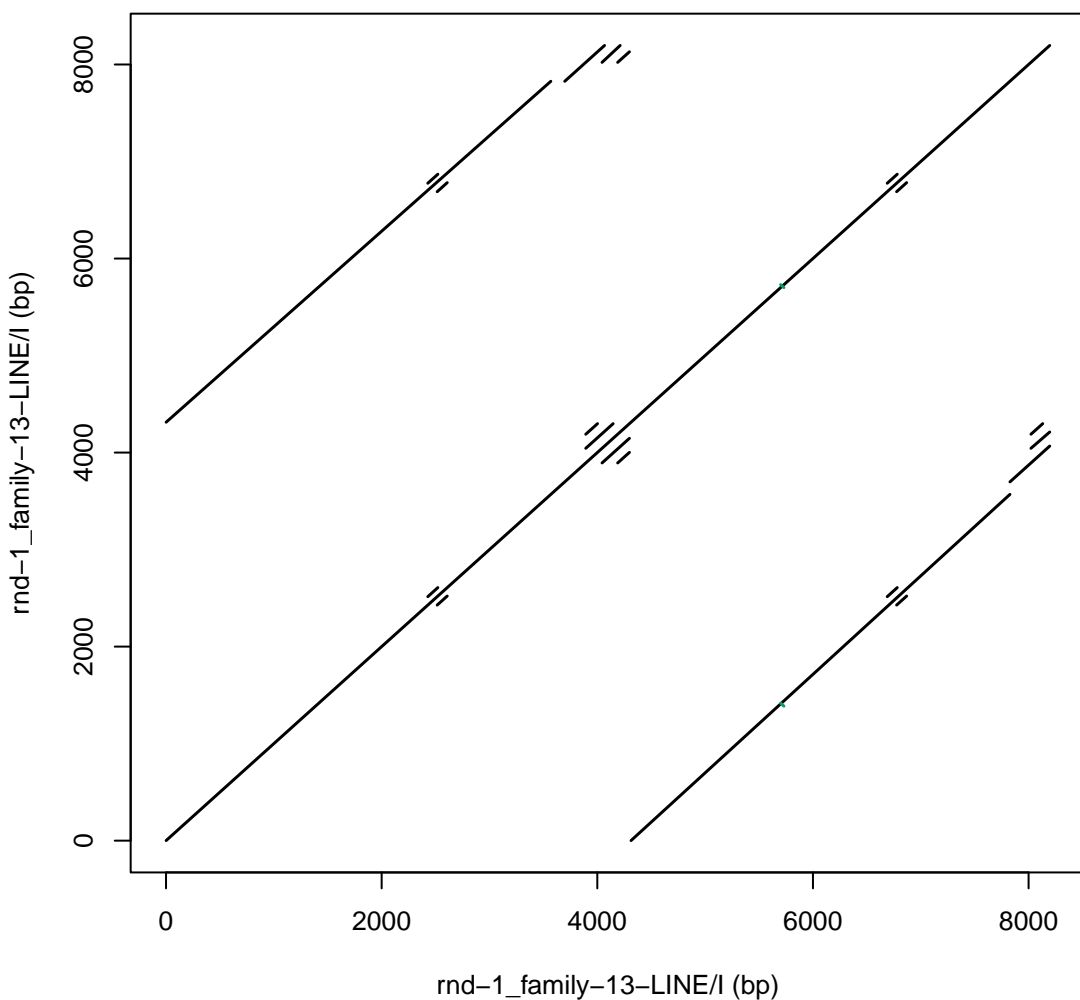
TE: rnd-1_family-13-LINE/I
consensus size: 8196bp; fragments: 1099; full length: 2 (>=7376.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

