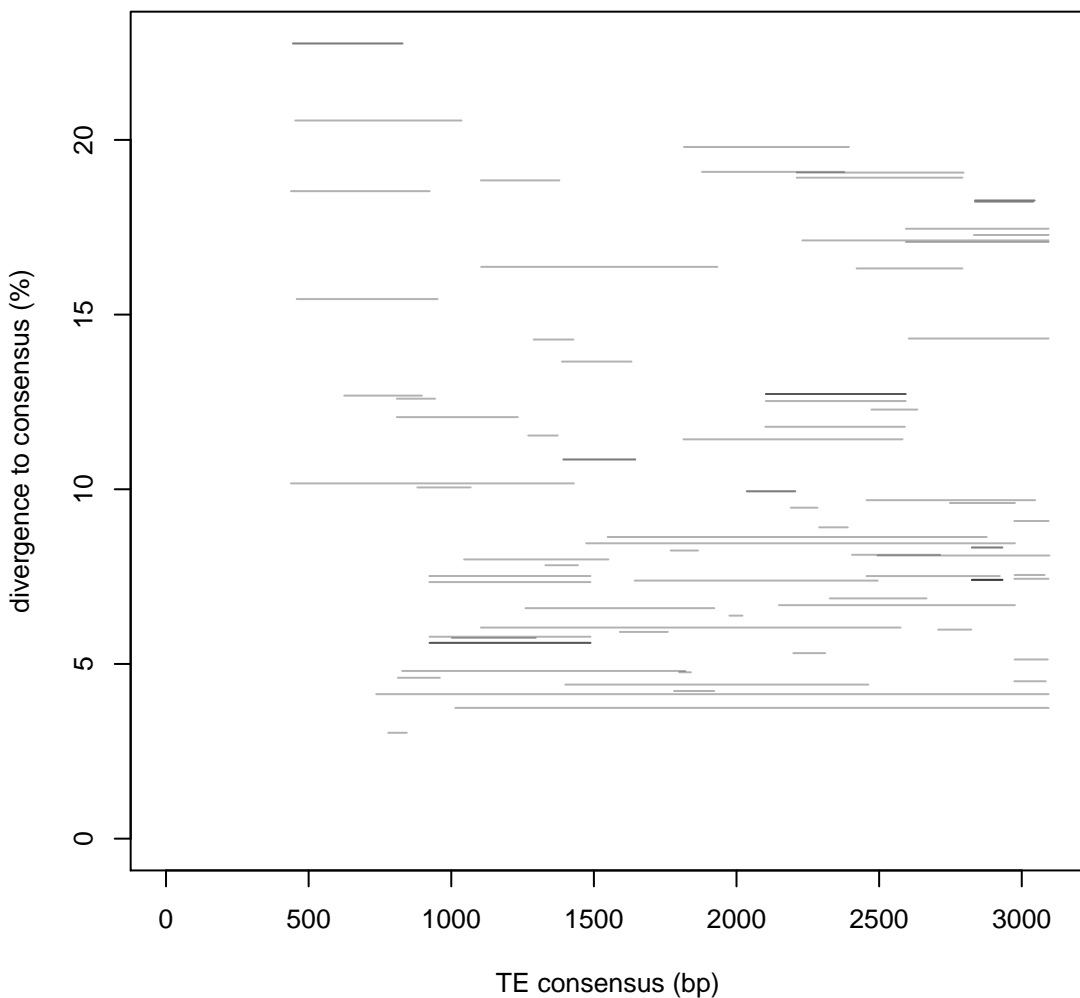
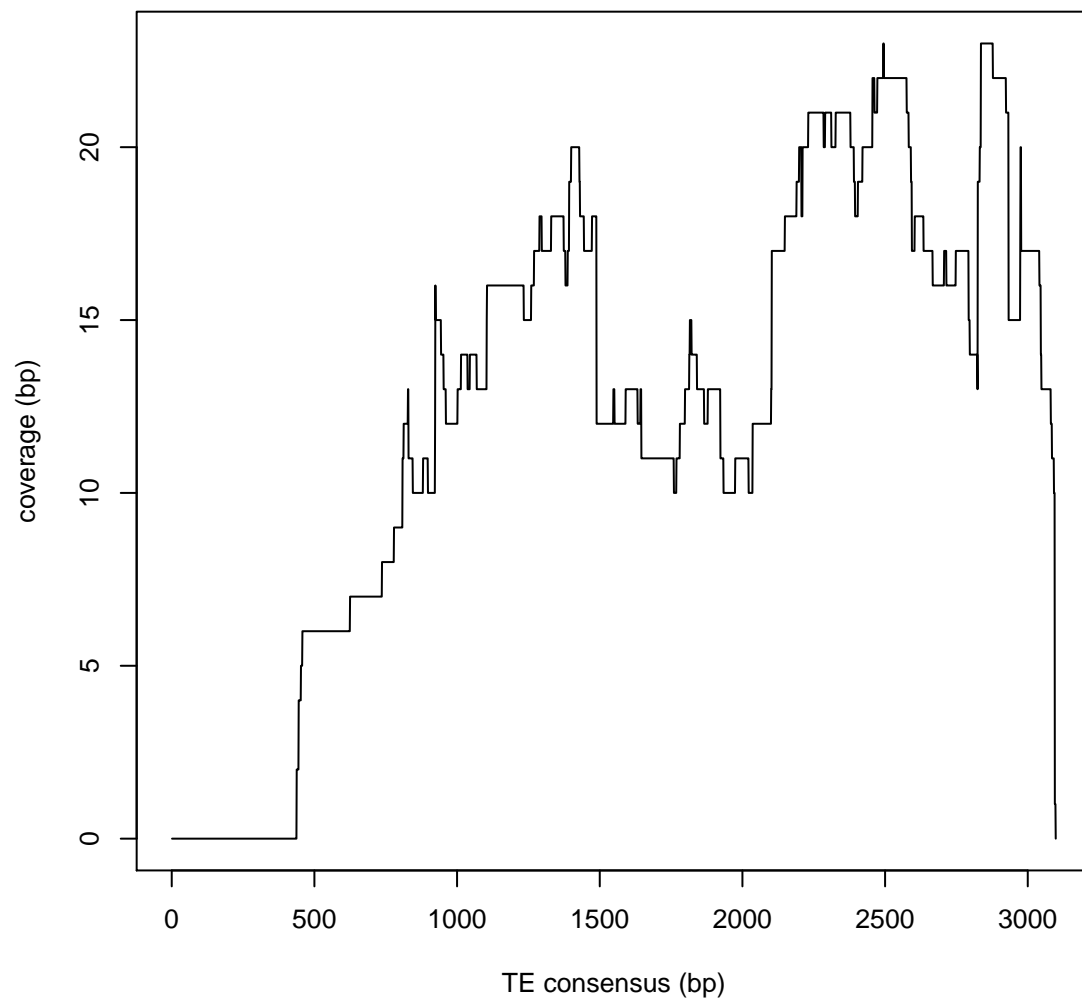


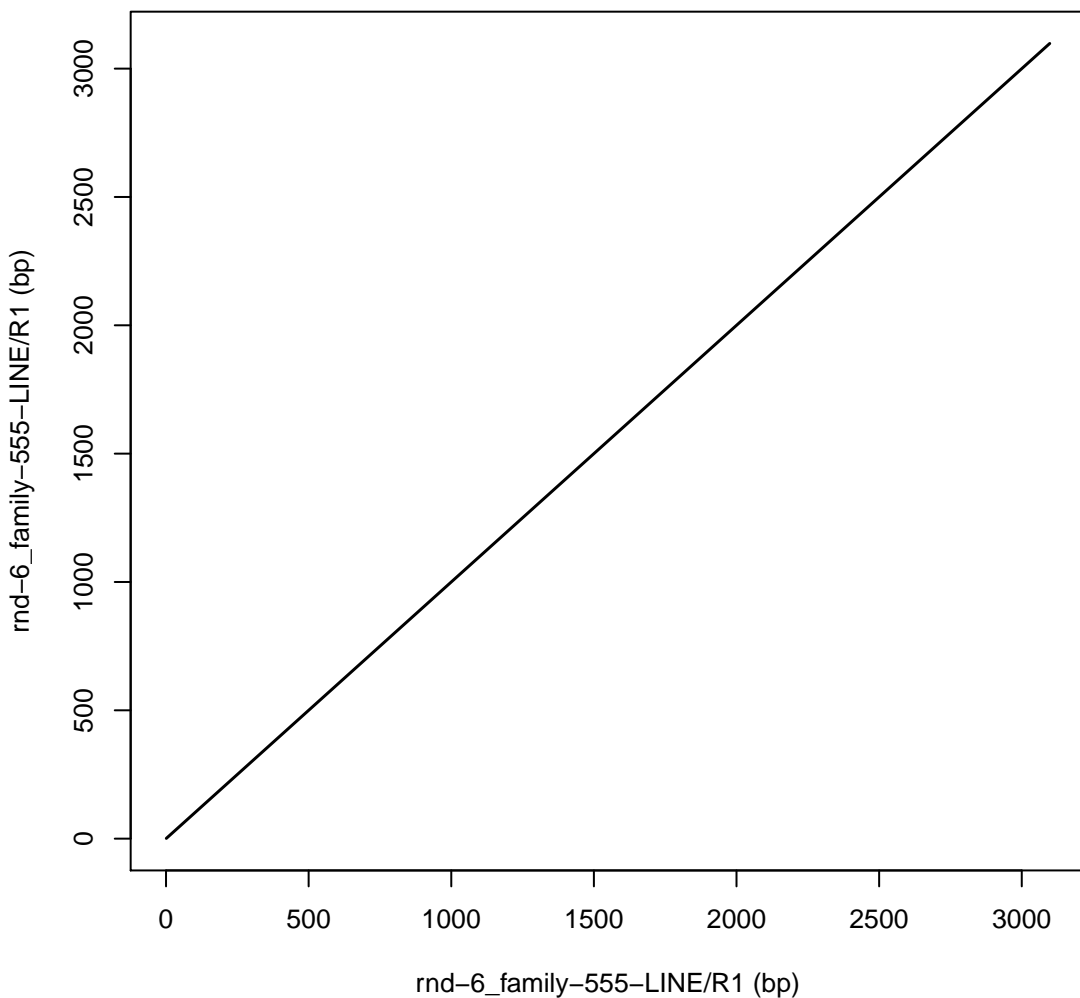
TE: rnd-6_family-555-LINE/R1
consensus size: 3098bp; fragments: 86; full length: 0 (>=2788.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

