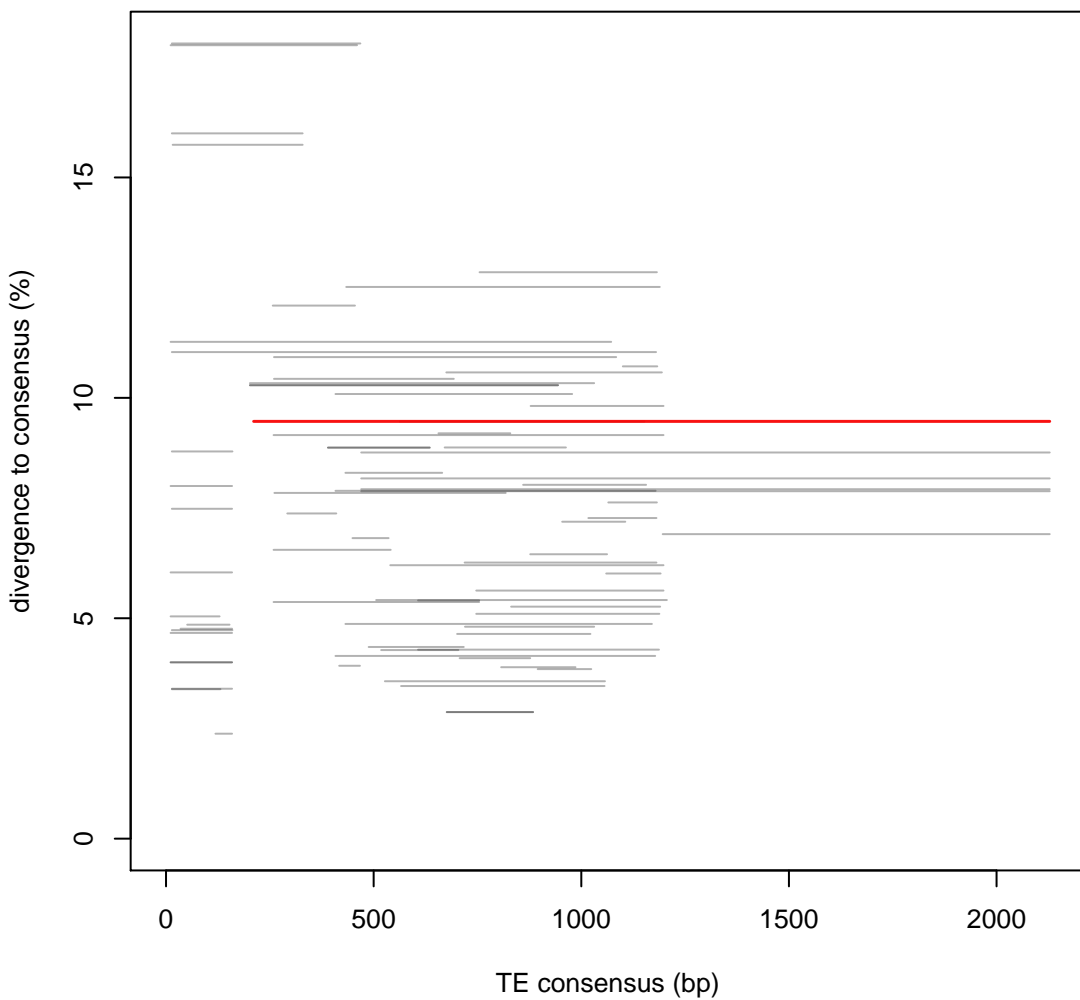
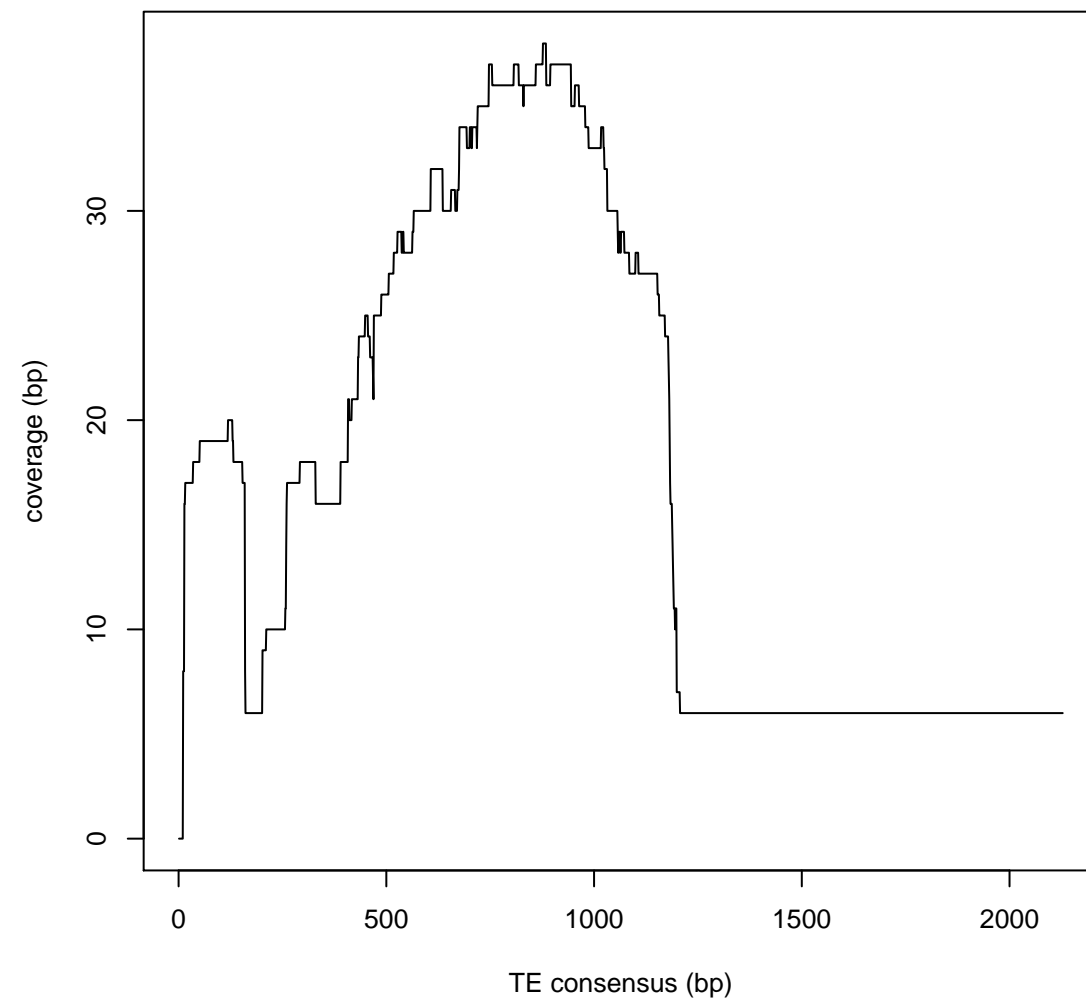


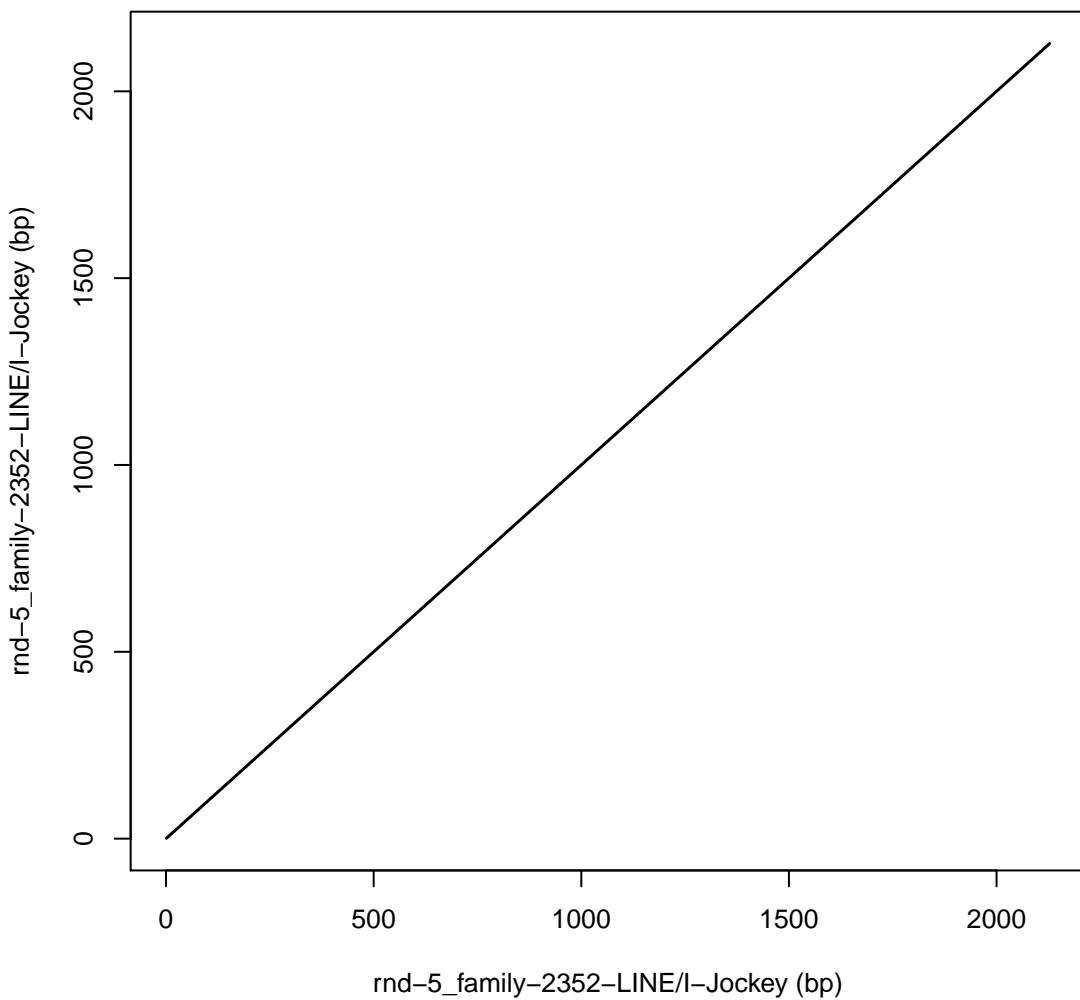
TE: rnd-5_family-2352-LINE/l-Jockey
consensus size: 2128bp; fragments: 79; full length: 1 (≥ 1915.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

