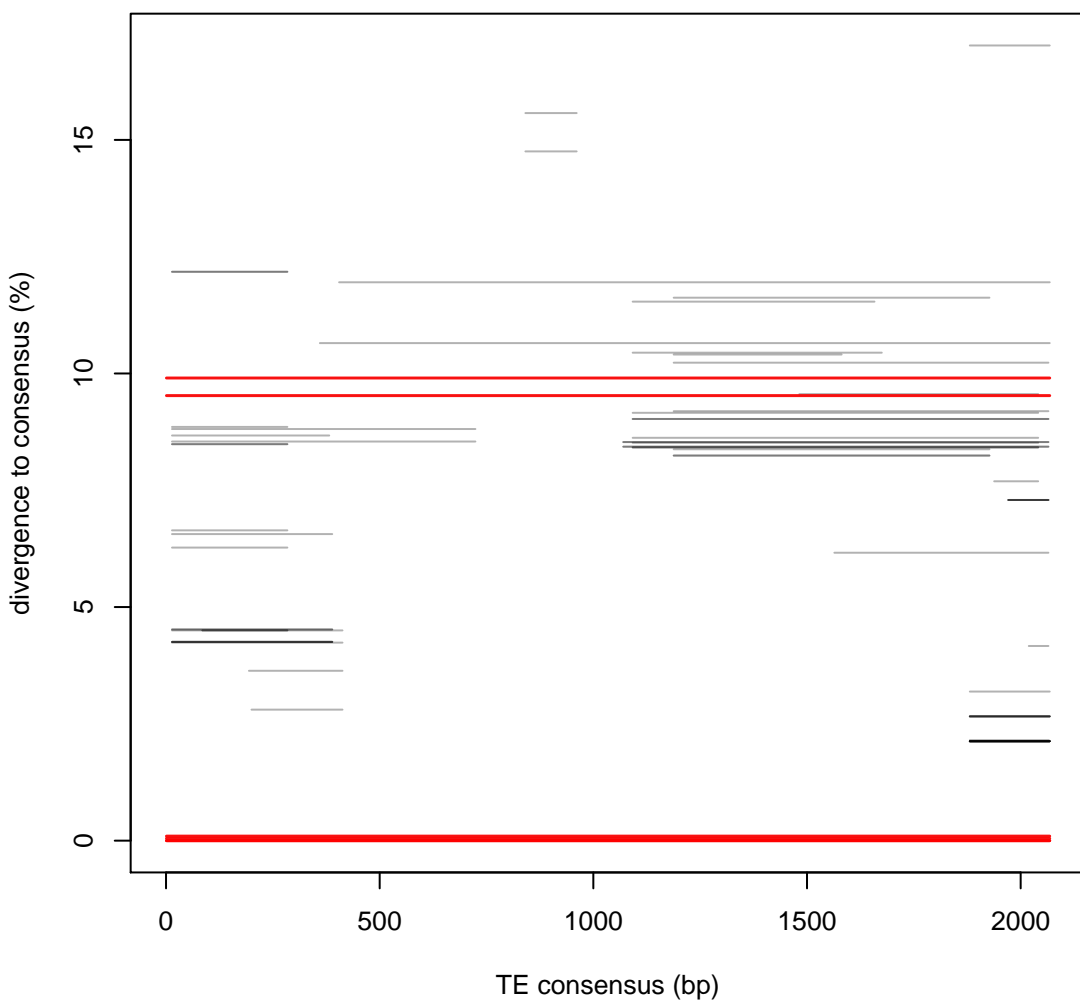
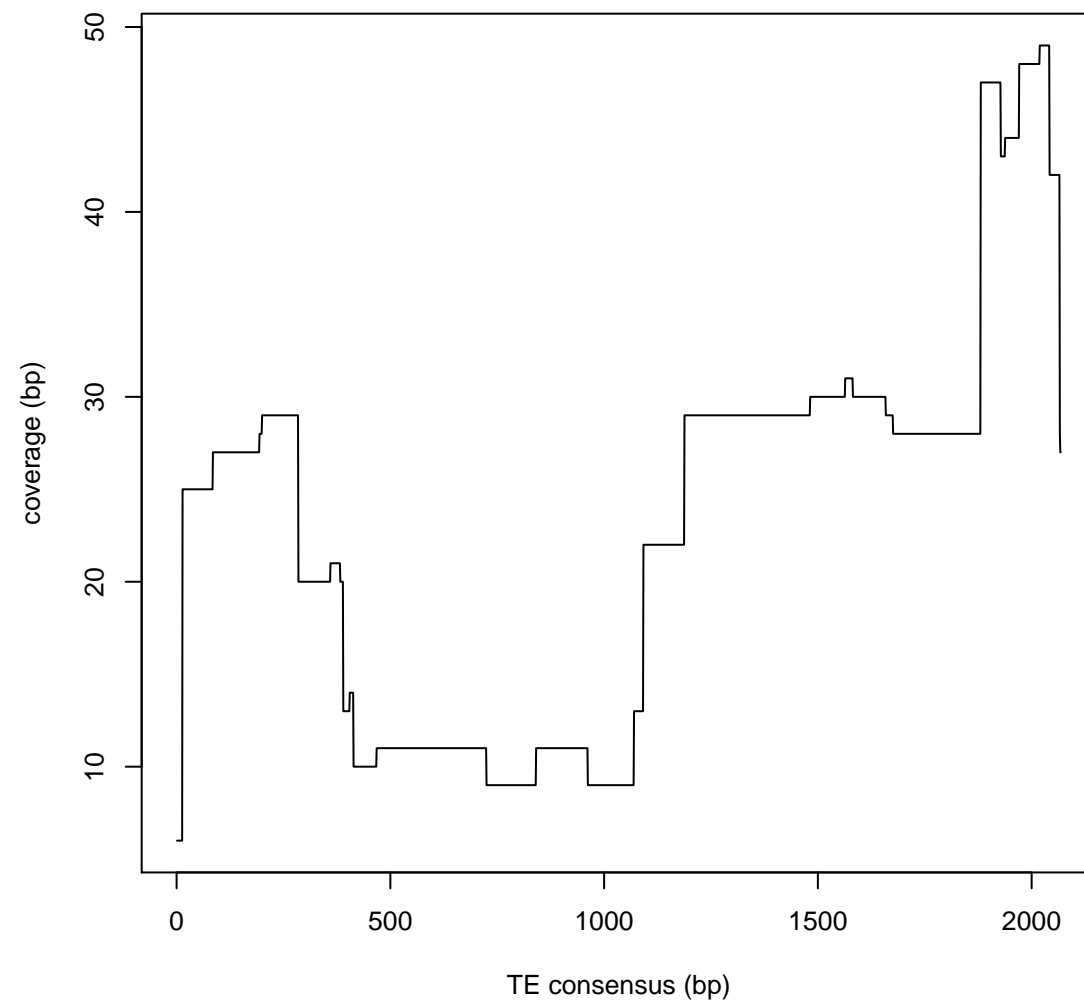


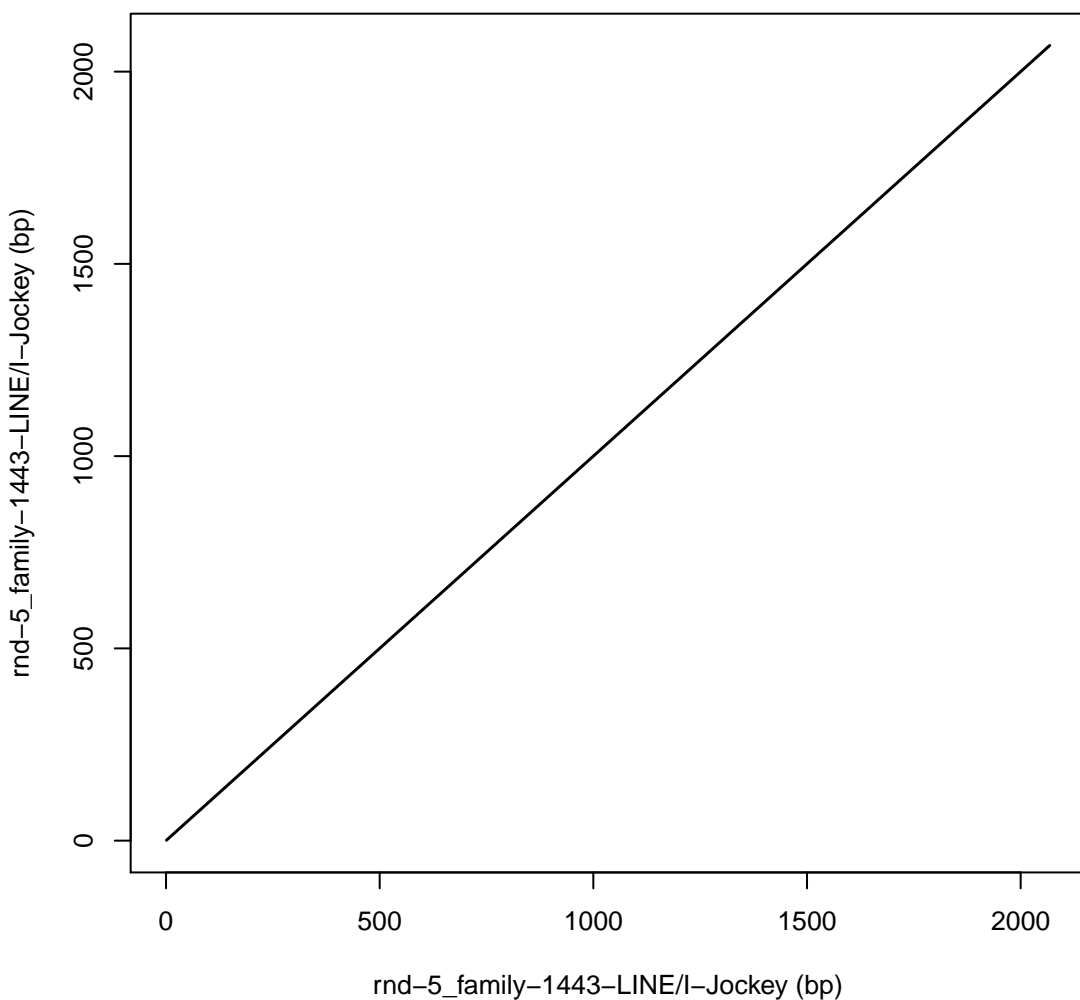
TE: rnd-5_family-1443-LINE/I-Jockey
consensus size: 2068bp; fragments: 81; full length: 6 (≥ 1861.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

