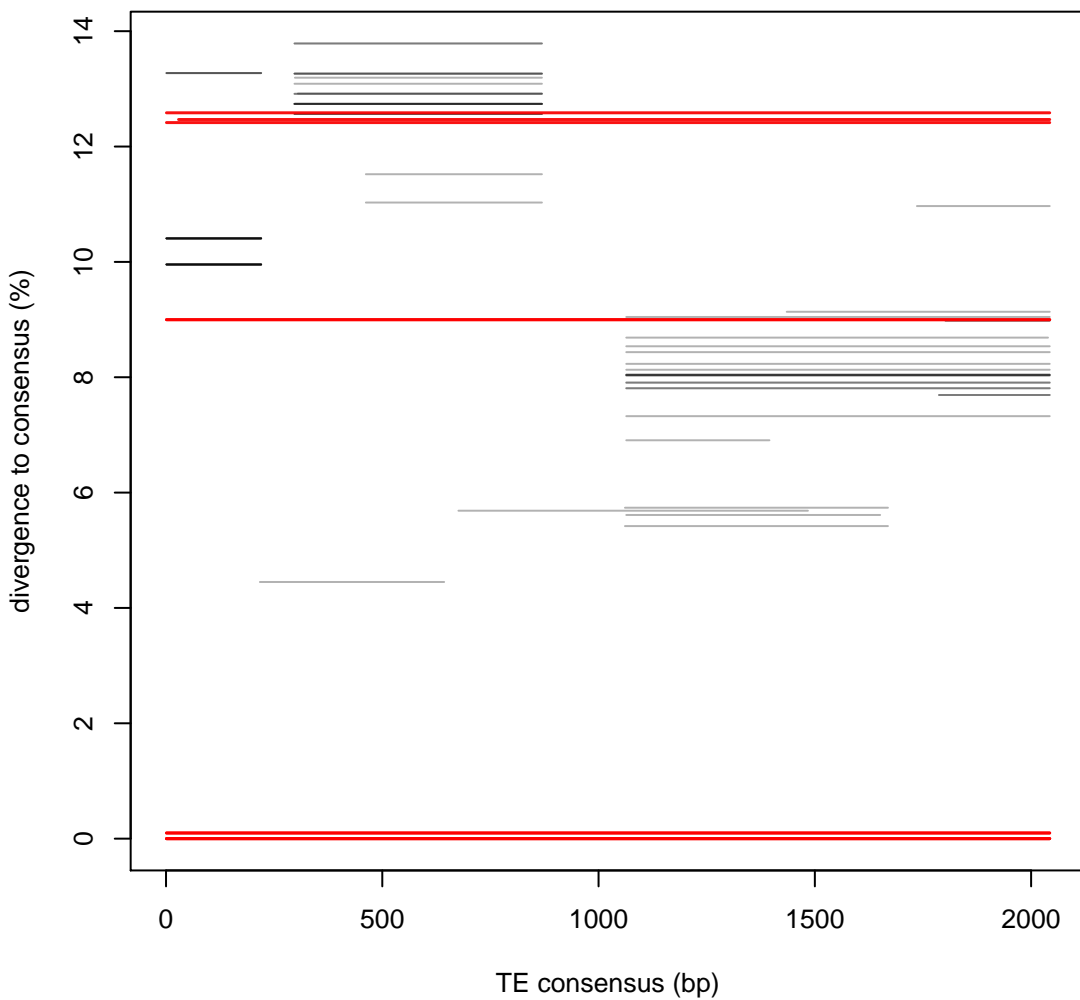
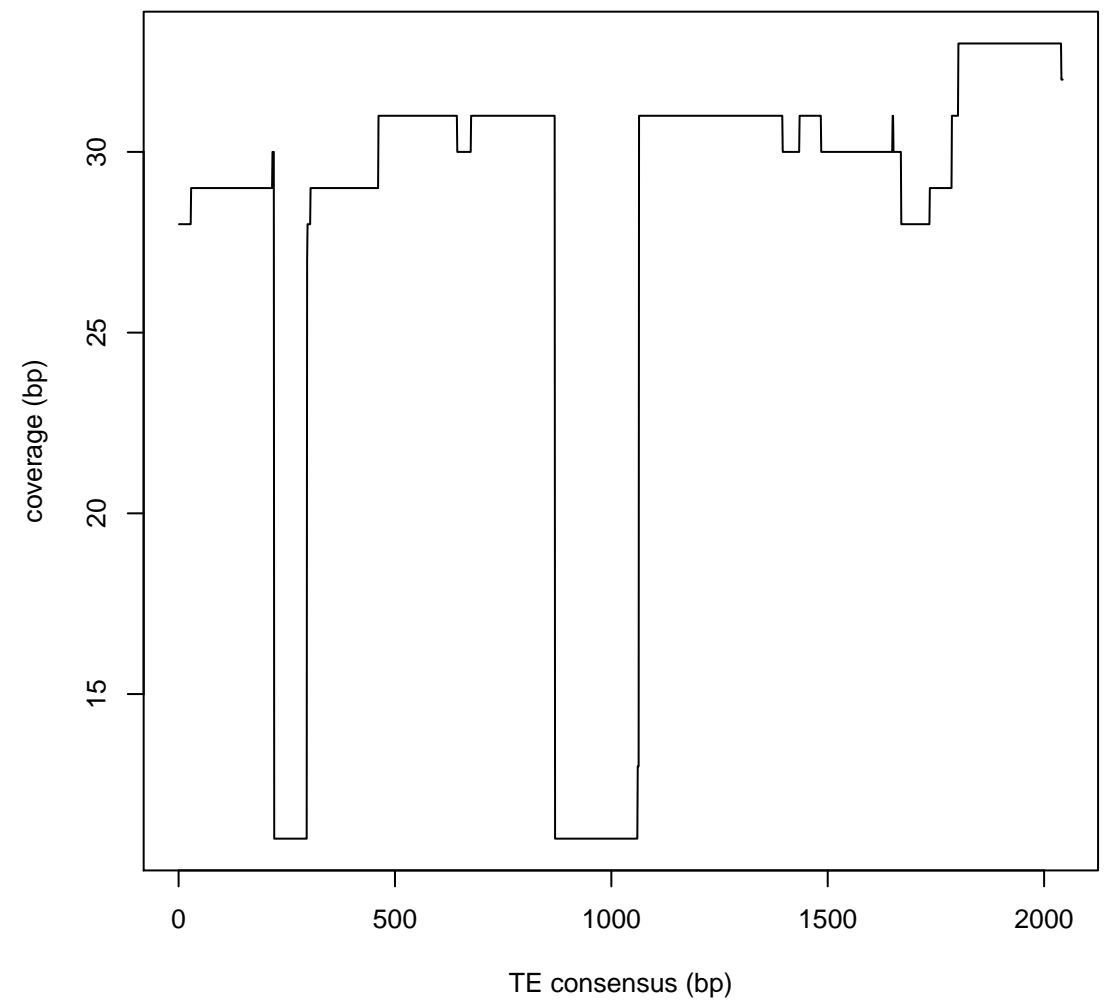


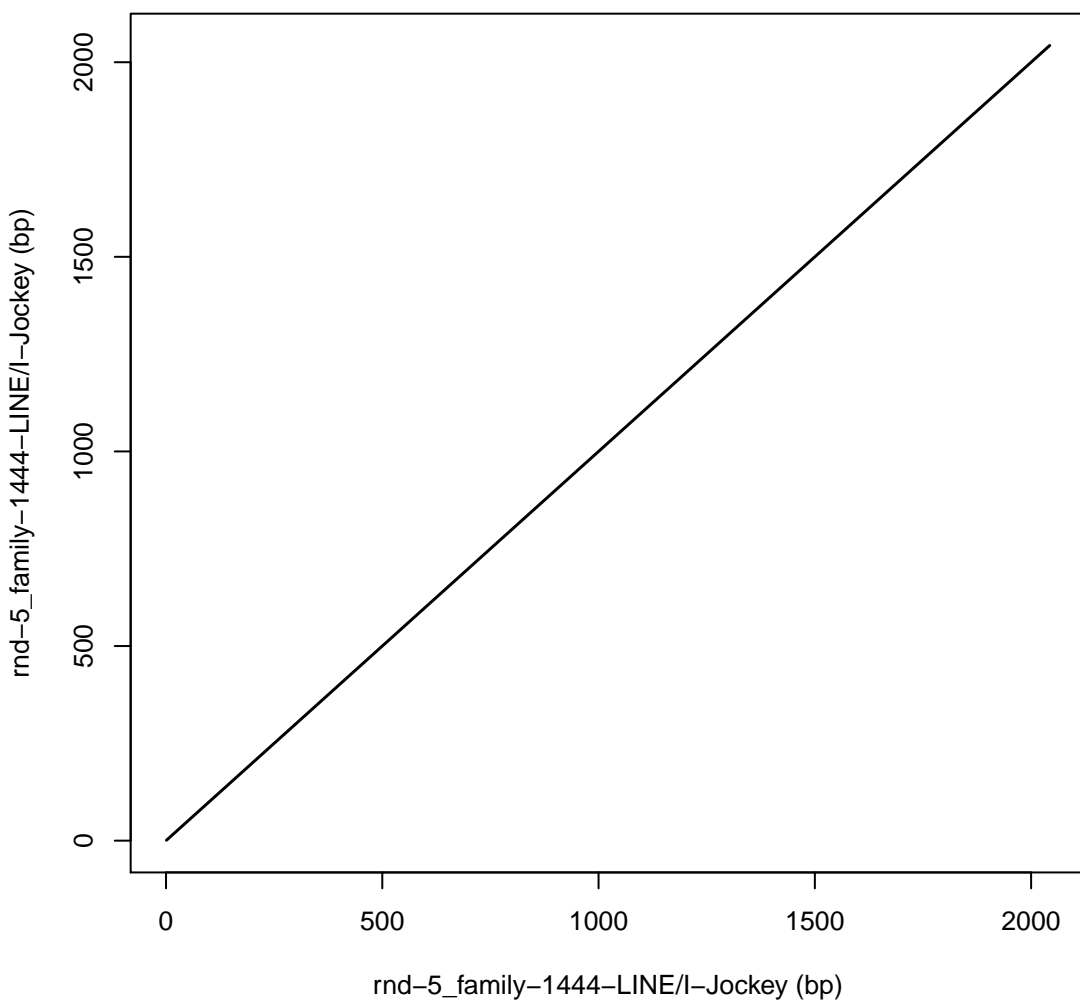
TE: rnd-5_family-1444-LINE/I-Jockey
consensus size: 2043bp; fragments: 78; full length: 10 (≥ 1838.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

