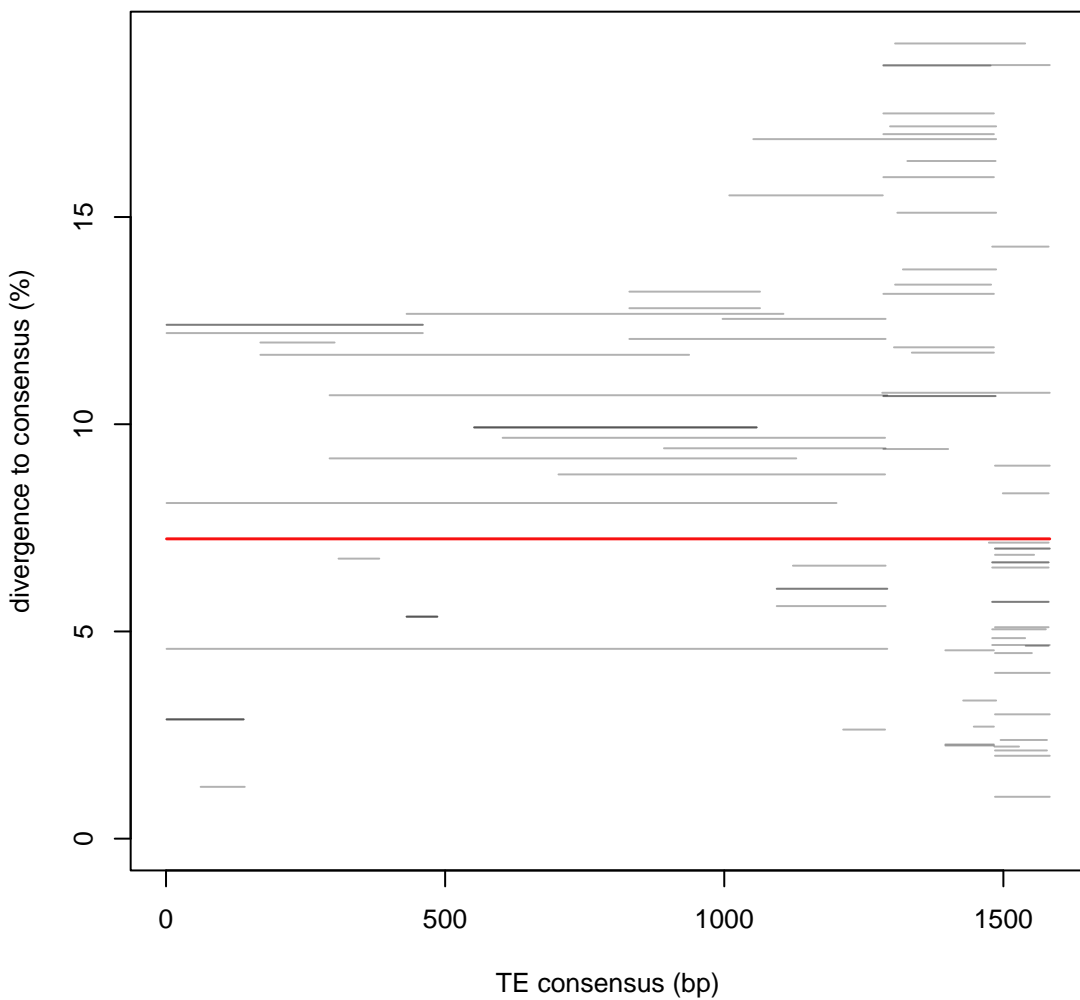
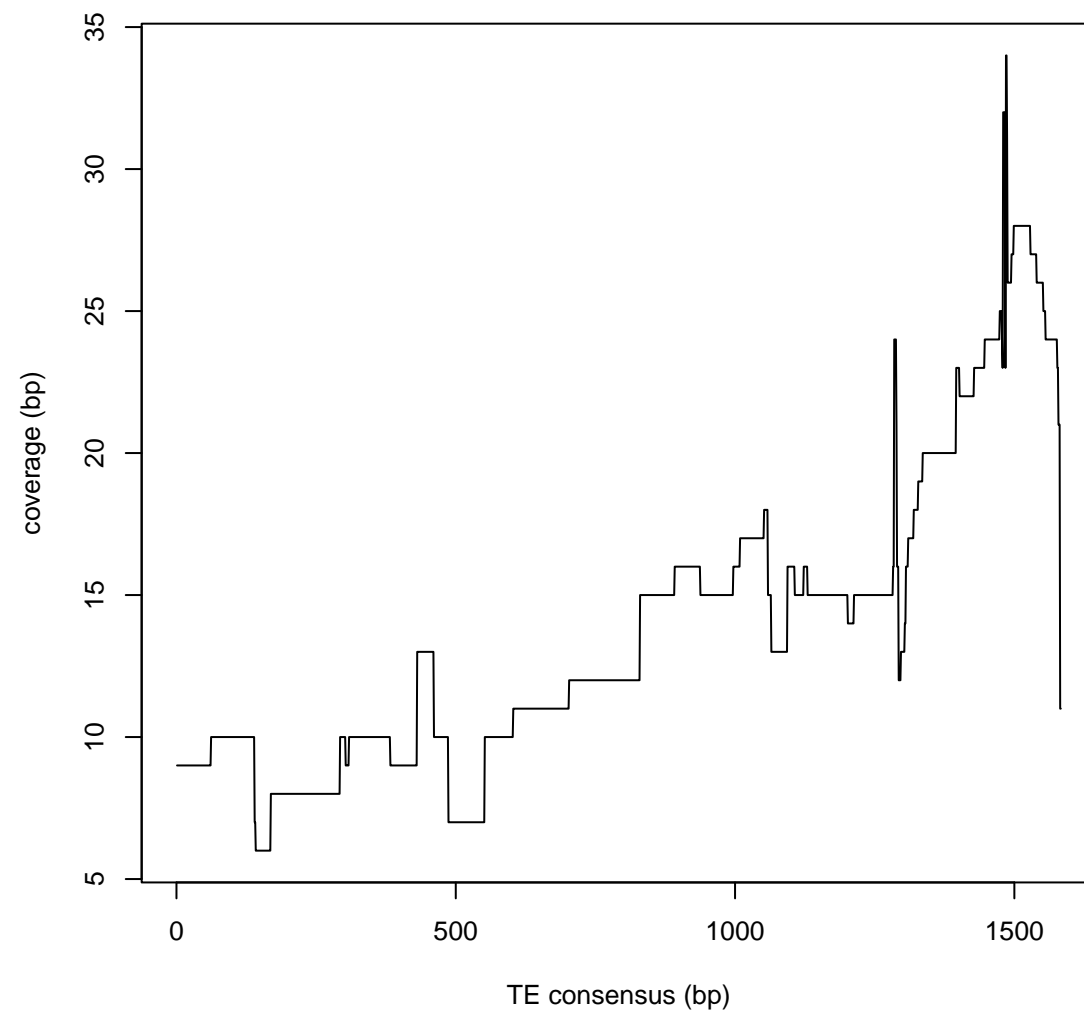


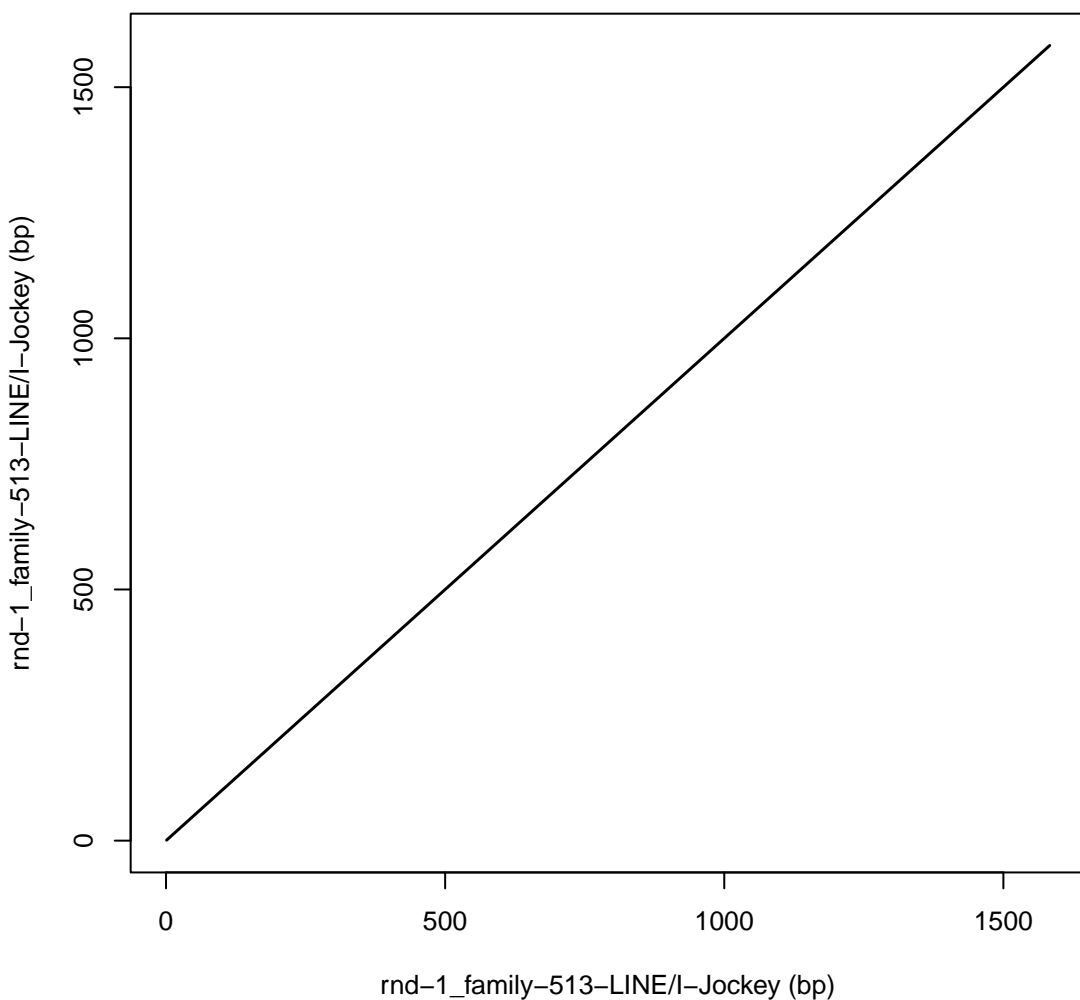
TE: rnd-1_family-513-LINE/I-Jockey
consensus size: 1583bp; fragments: 84; full length: 1 (≥ 1424.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

