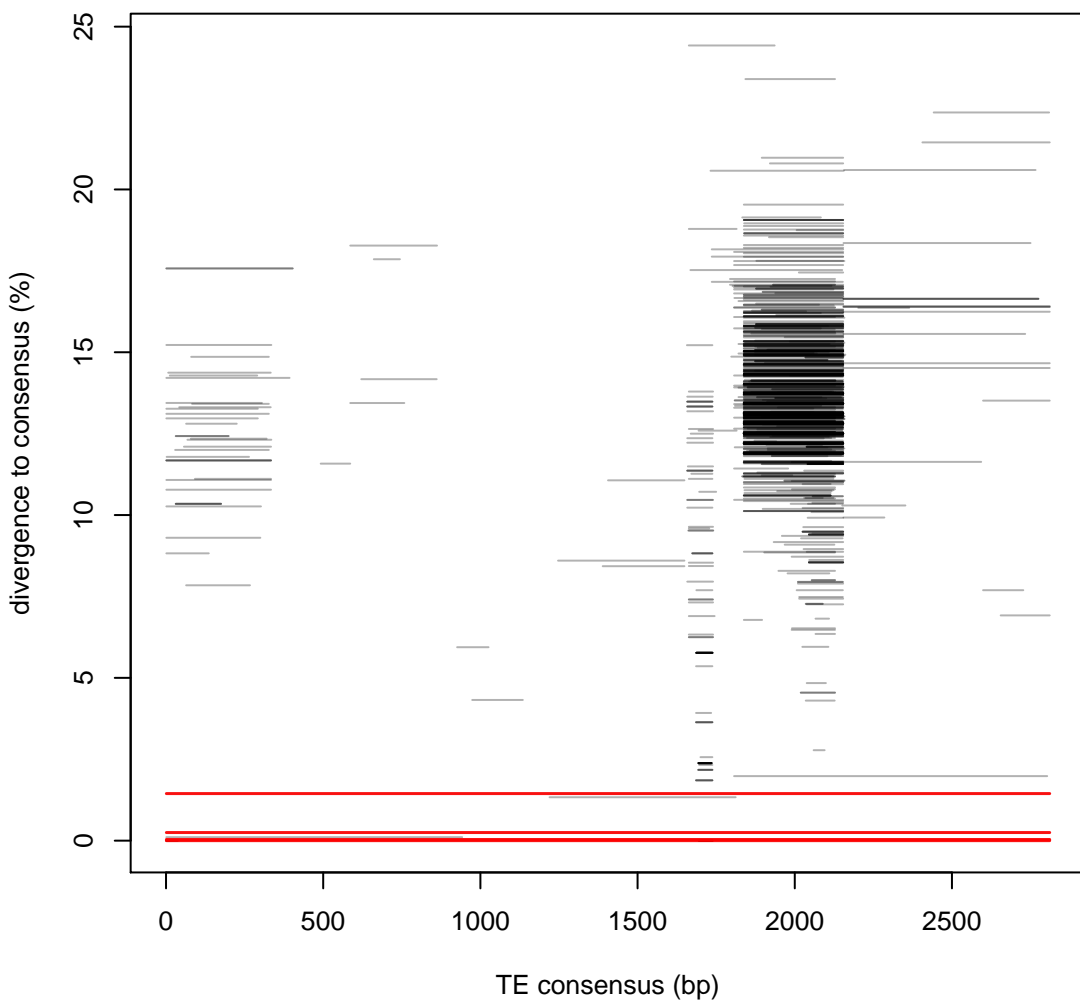
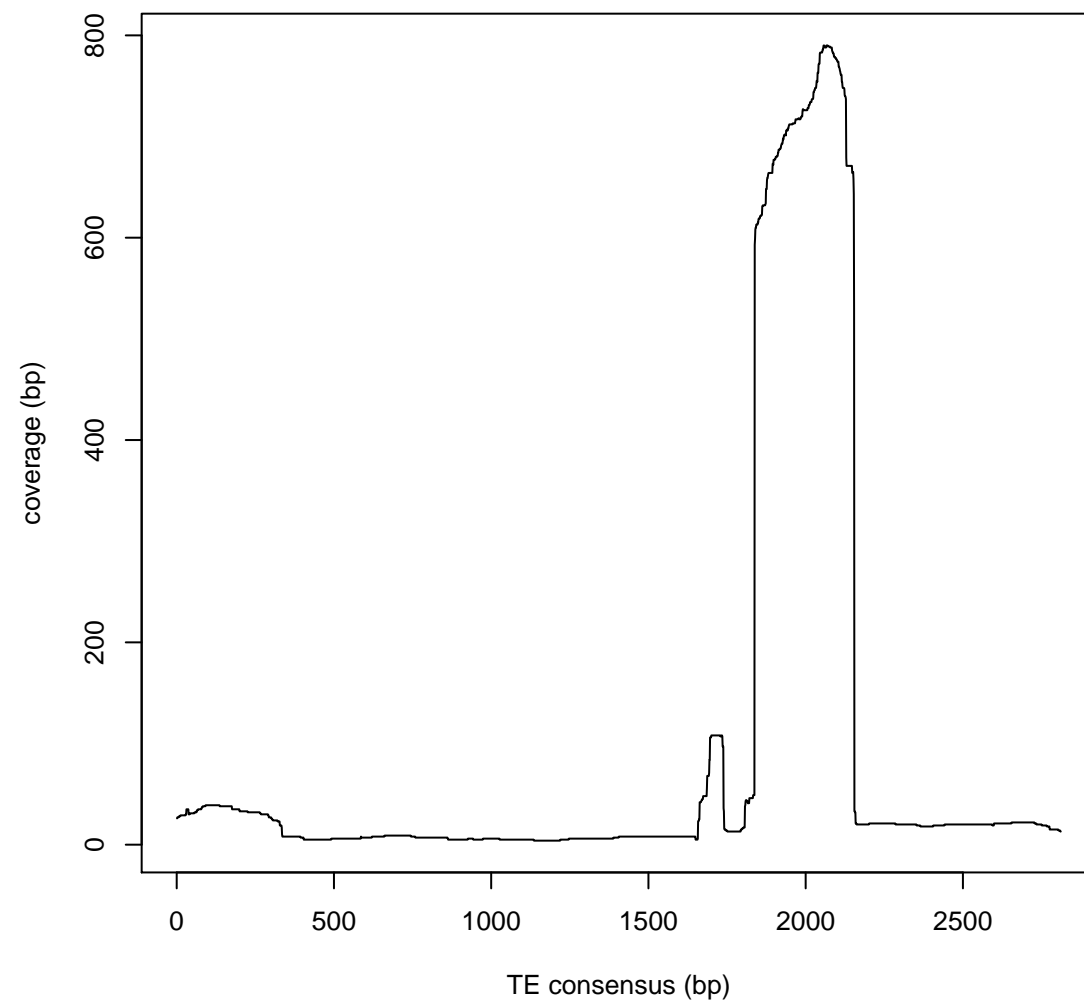


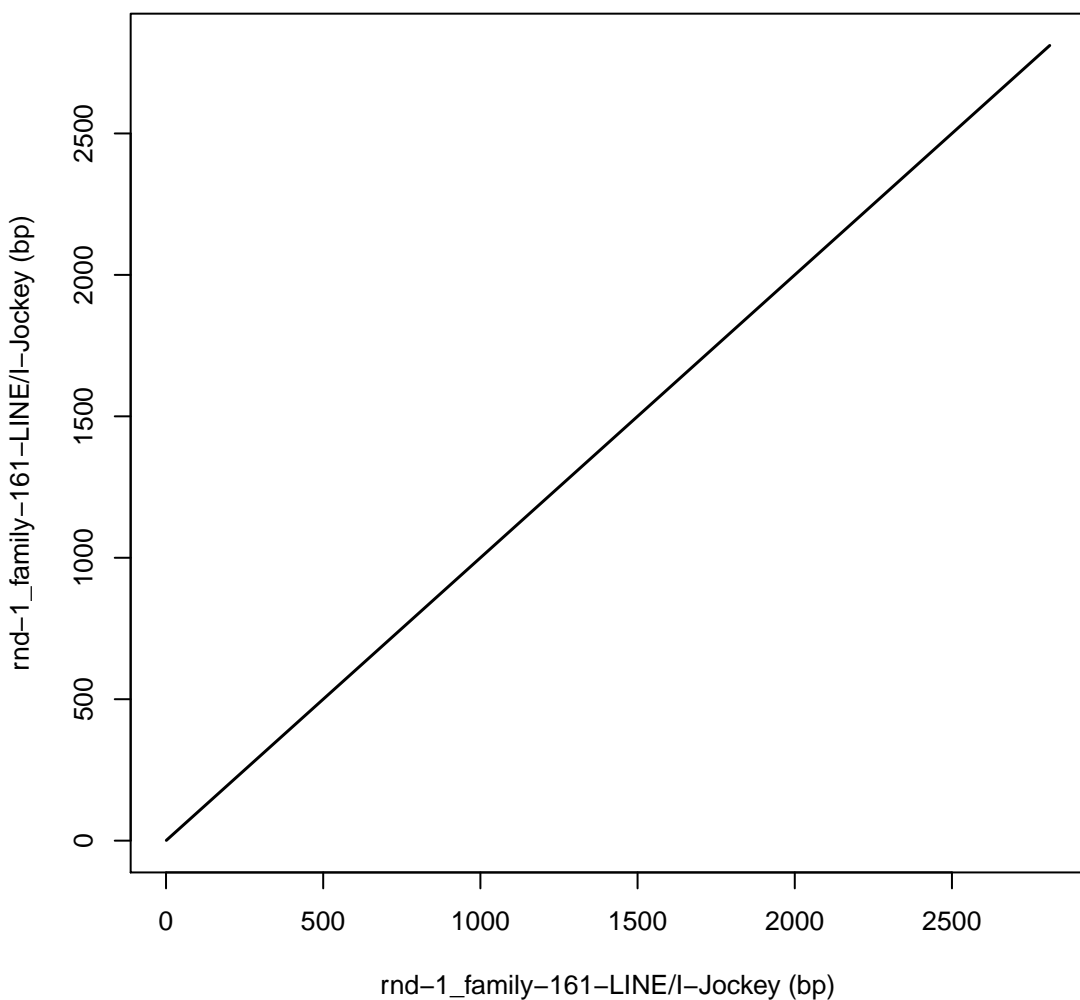
TE: rnd-1_family-161-LINE/I-Jockey
consensus size: 2811bp; fragments: 973; full length: 4 (>=2529.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

