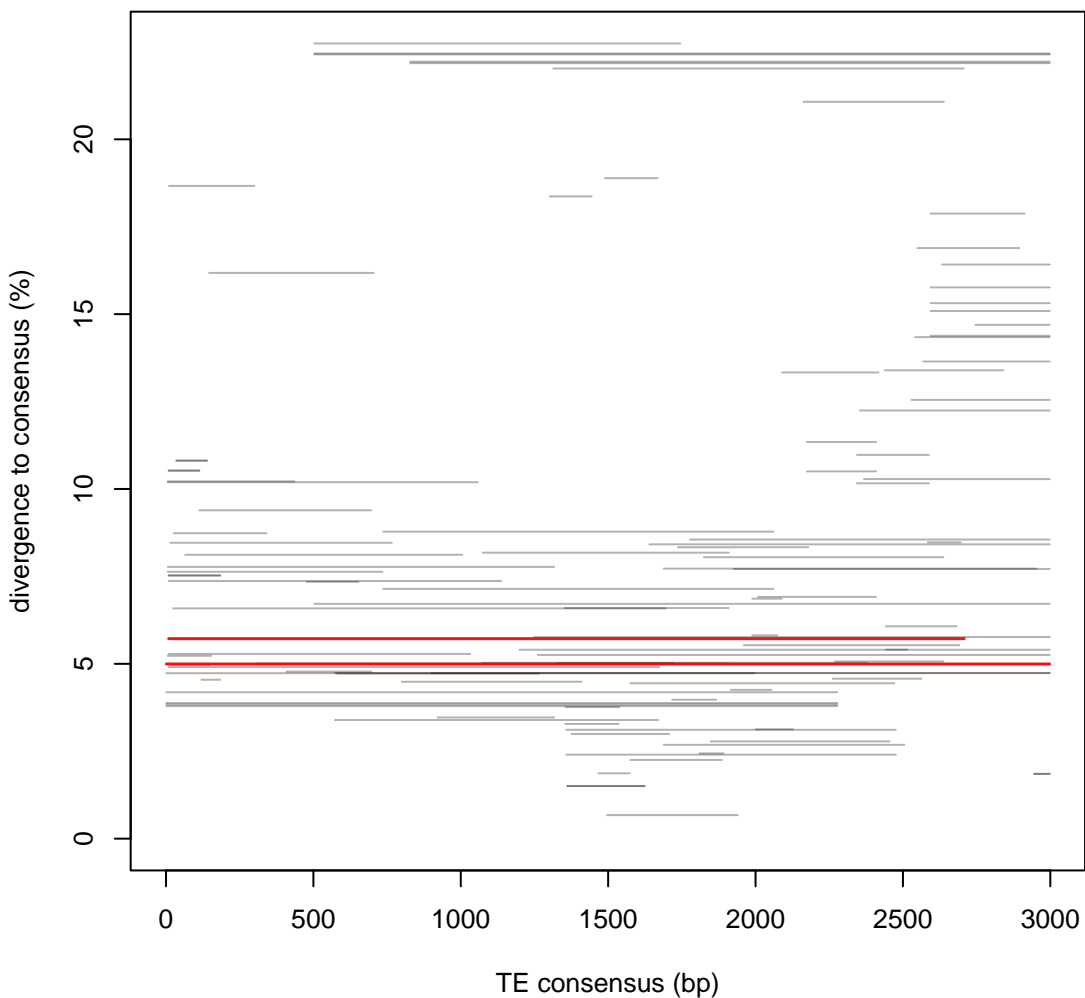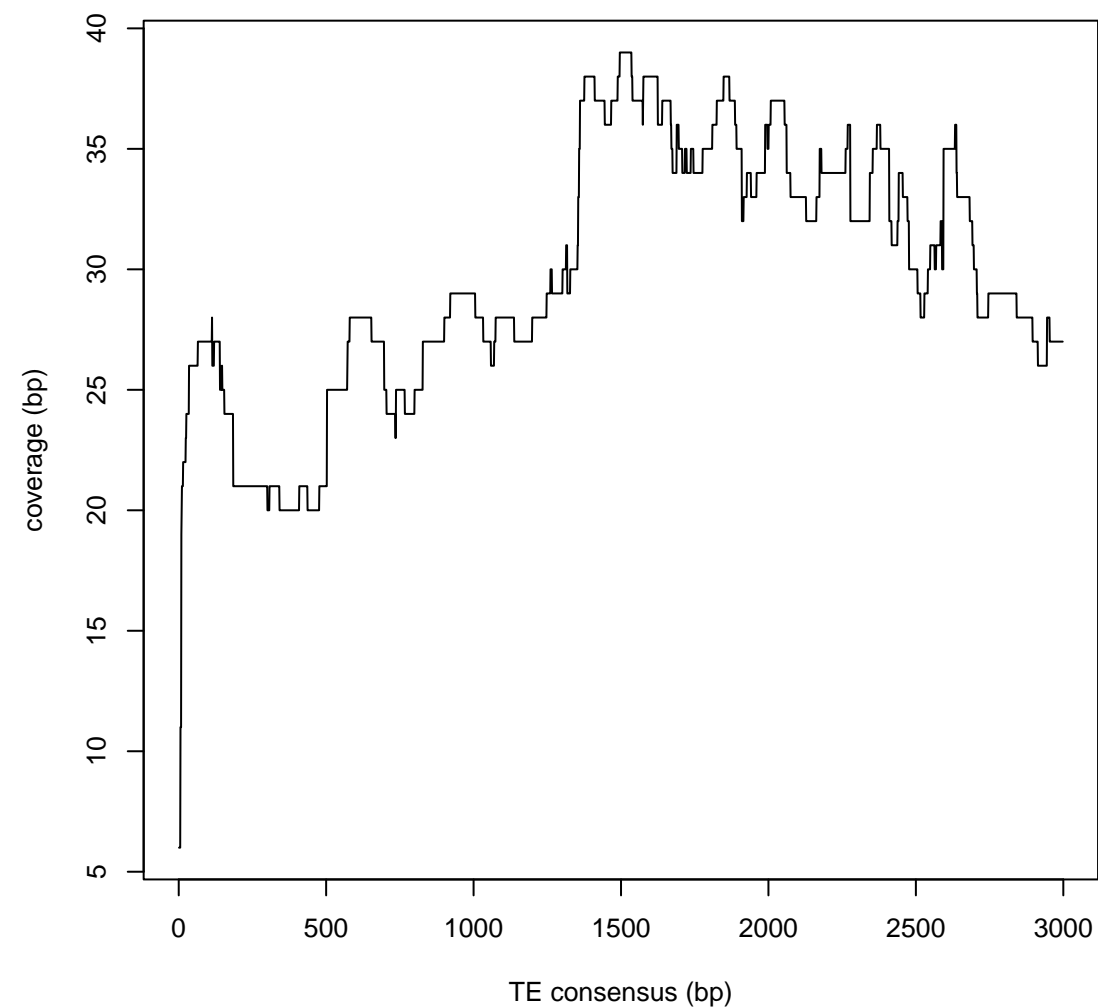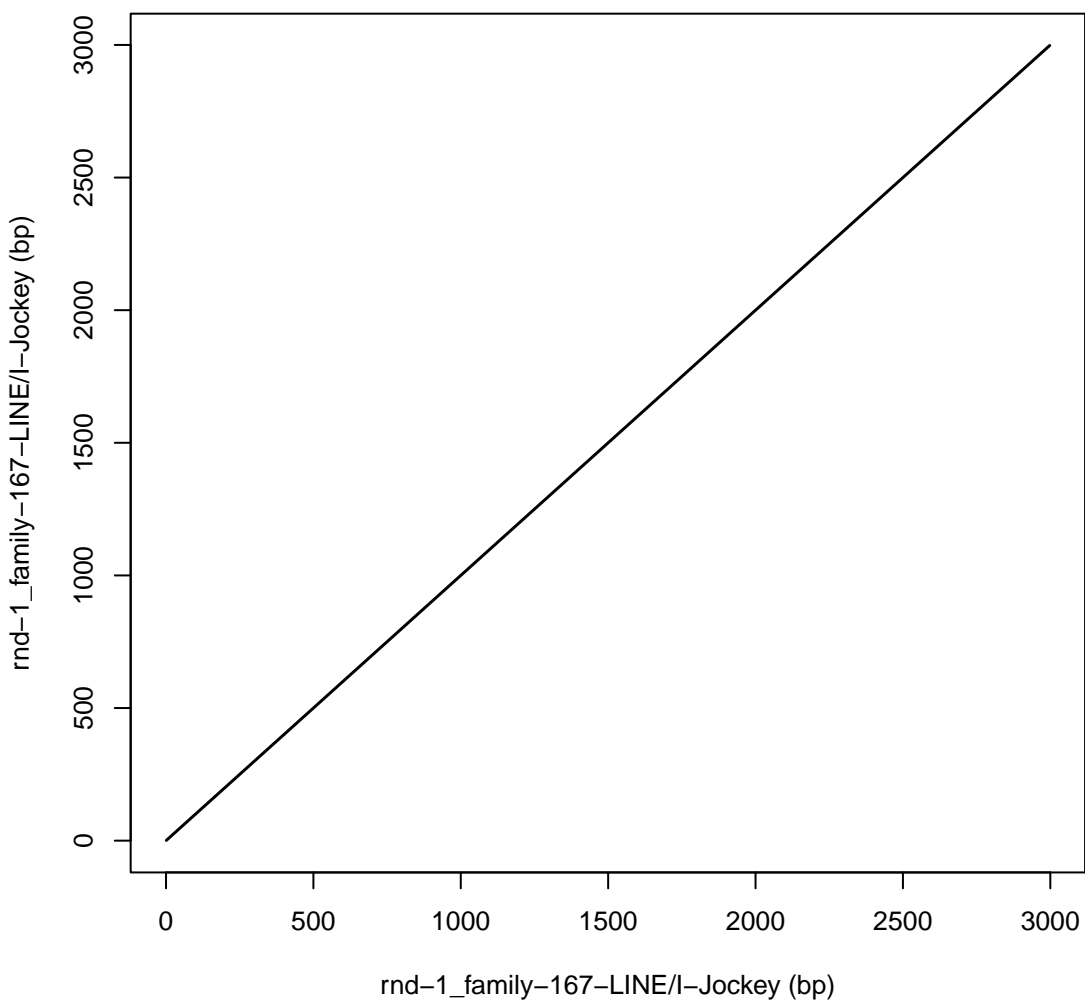**TE: rnd−1_family−167−LINE/I−Jockey**
consensus size: 2998bp; fragments: 111; full length: 2 (>=2698.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits