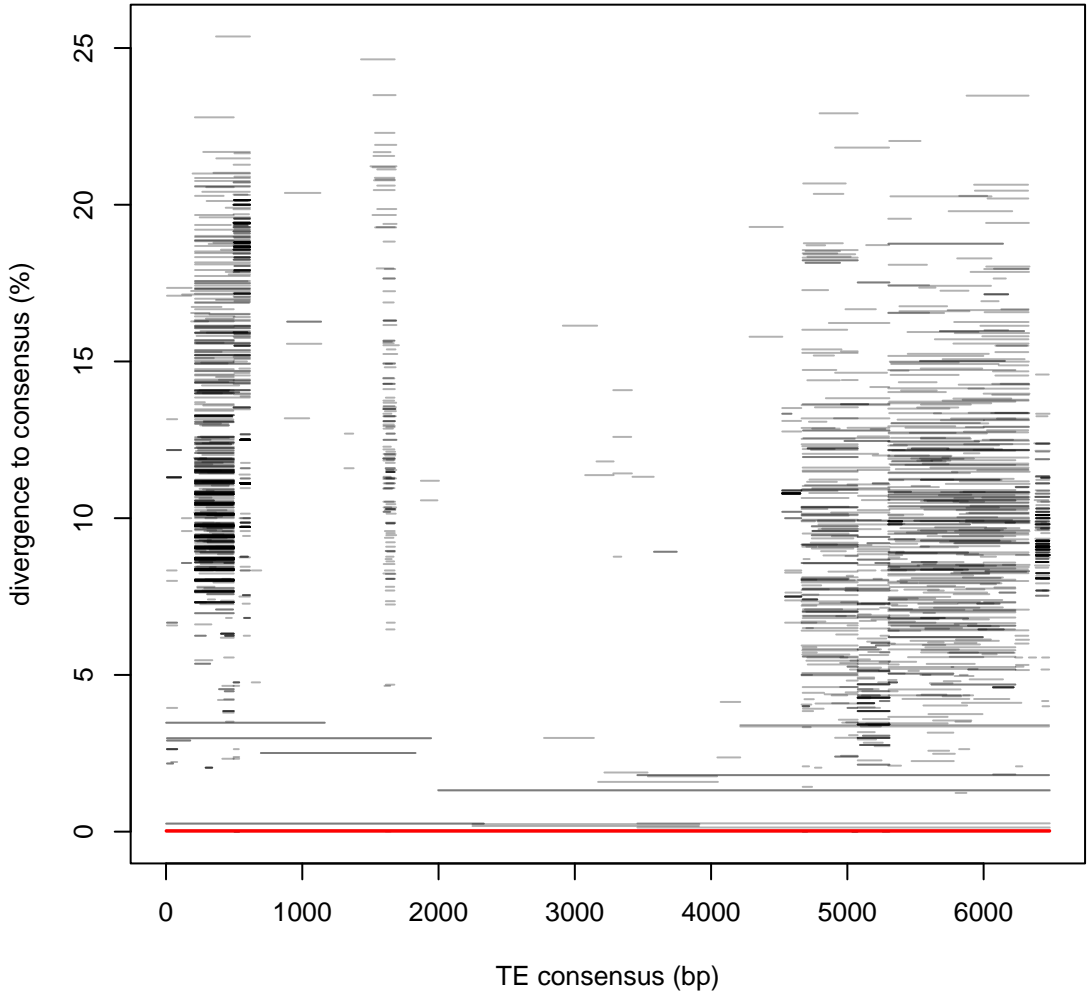
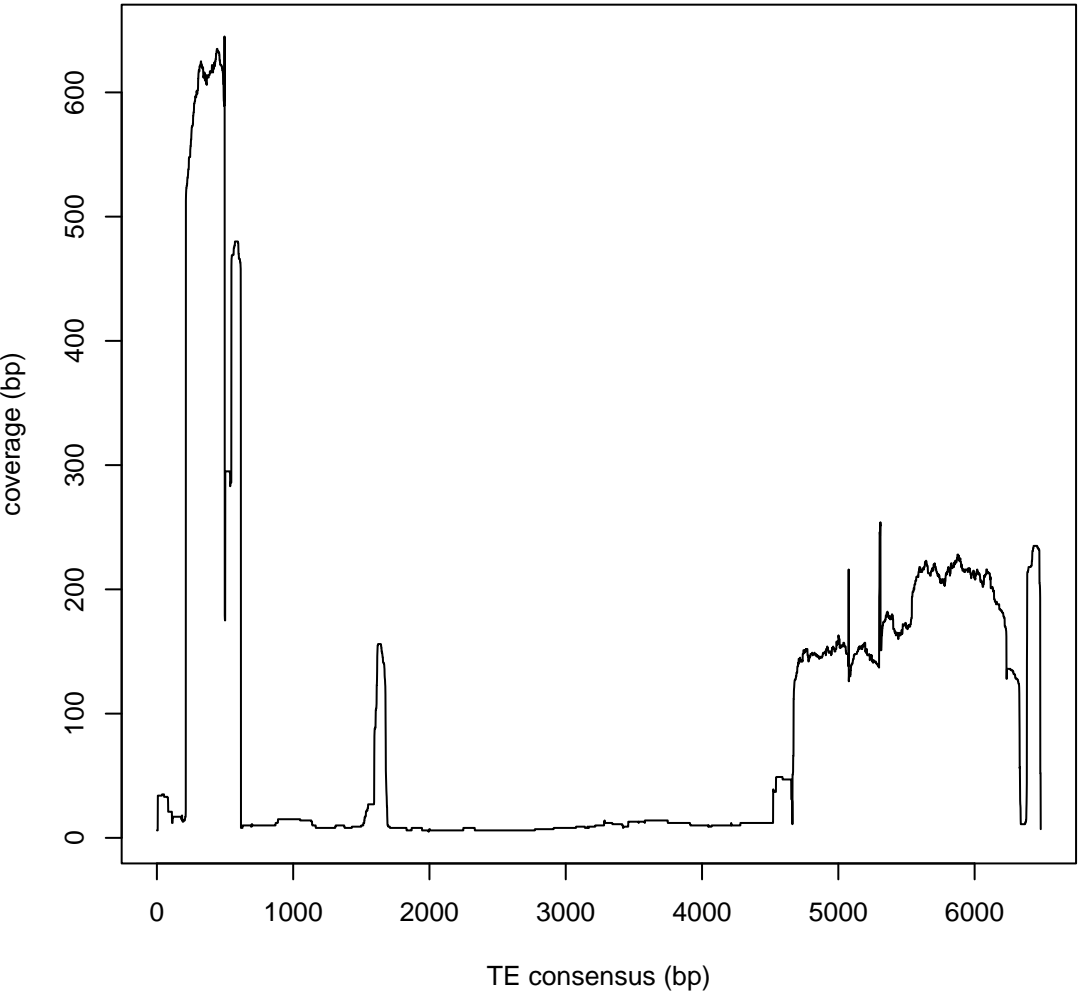


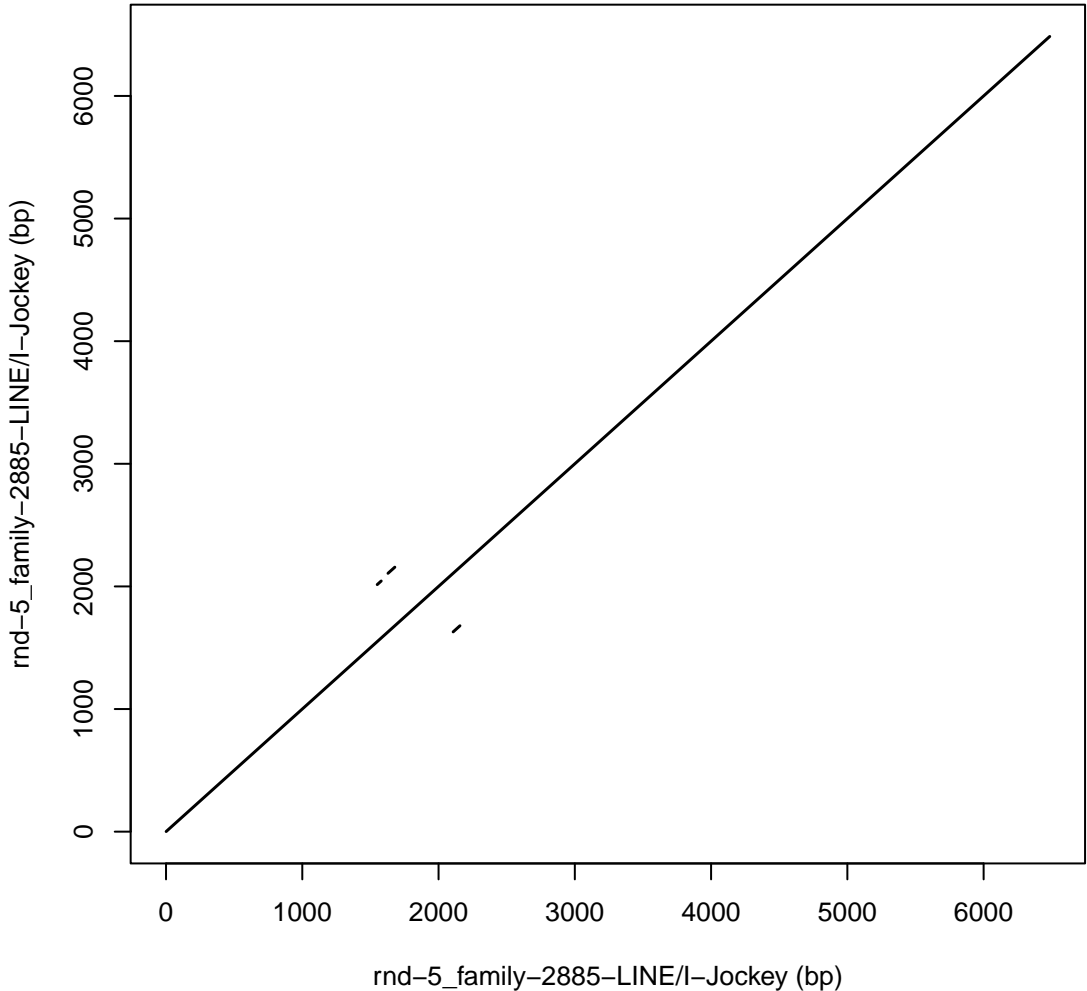
TE: rnd-5_family-2885-LINE/I-Jockey
consensus size: 6485bp; fragments: 2377; full length: 2 (>=5836.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

