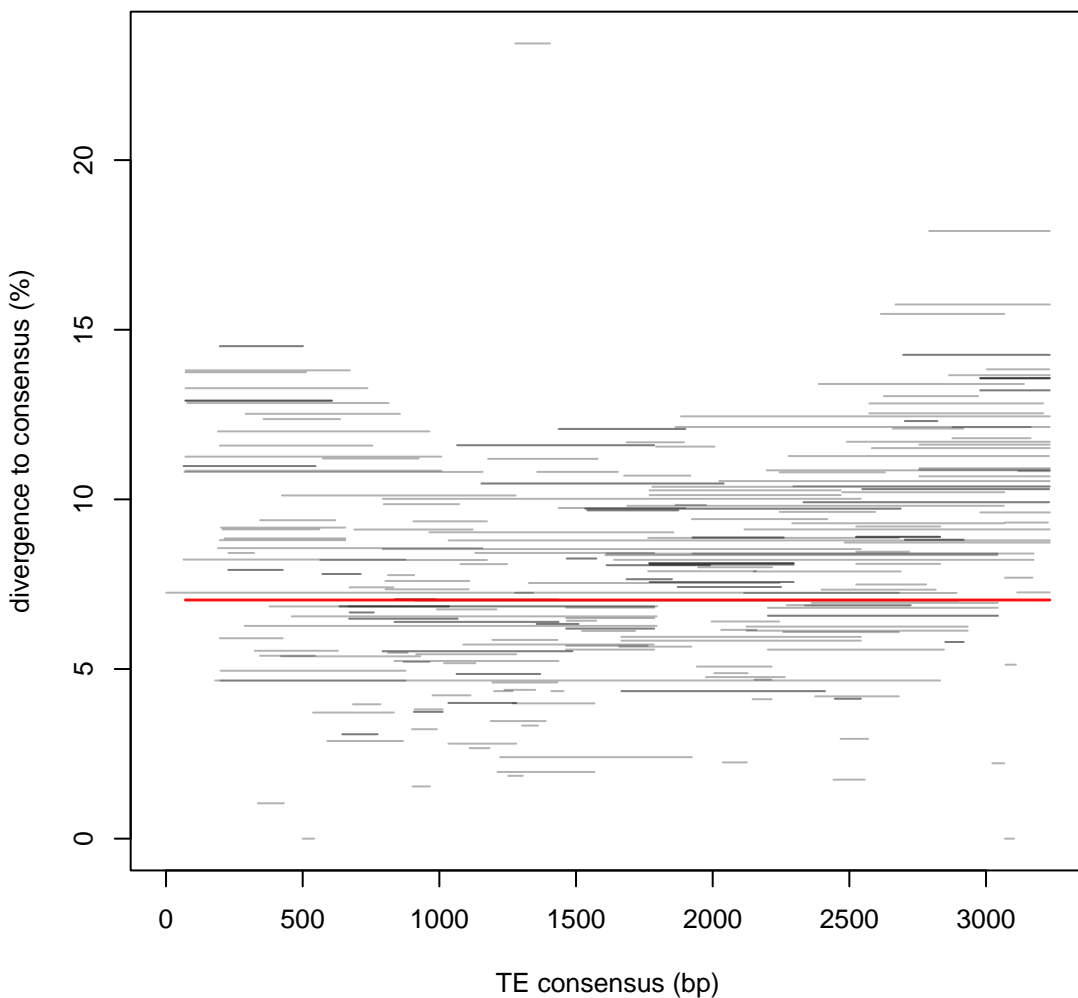
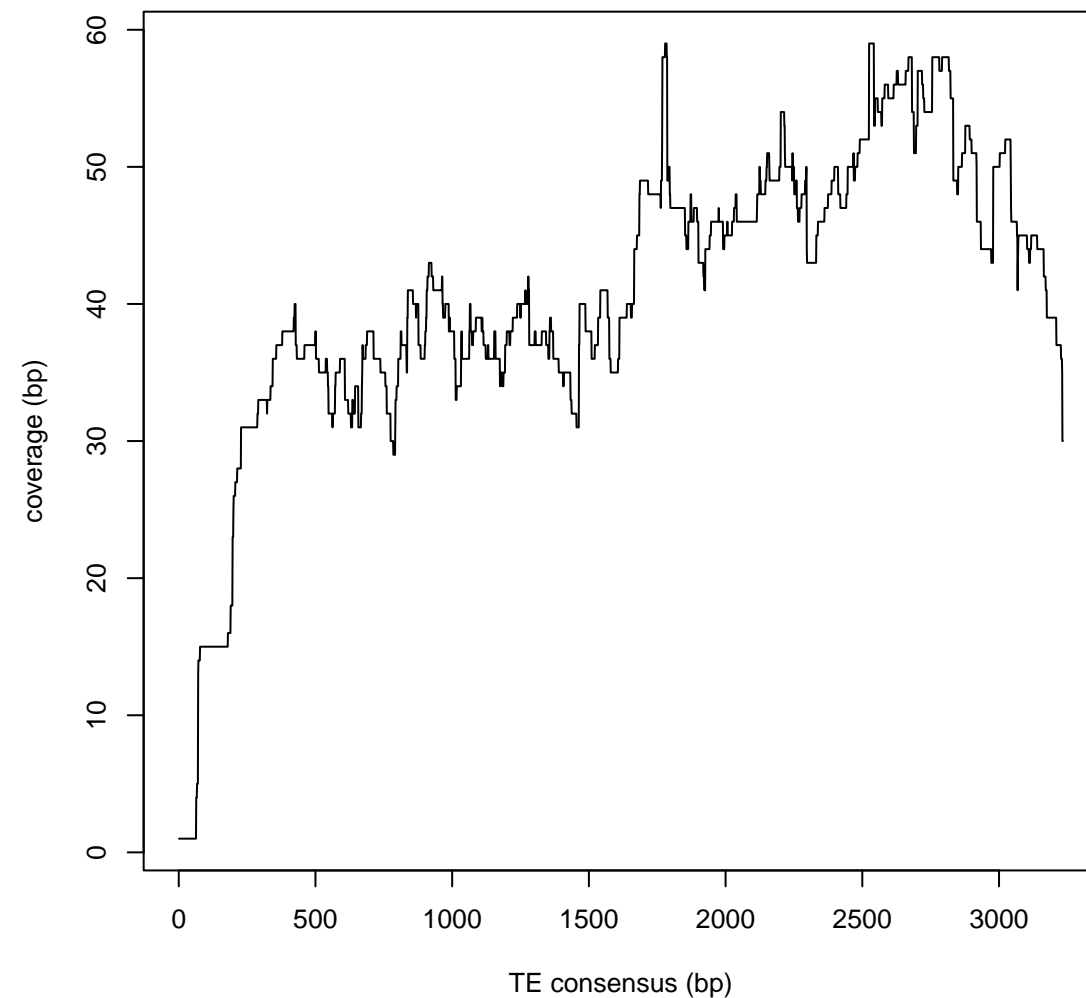


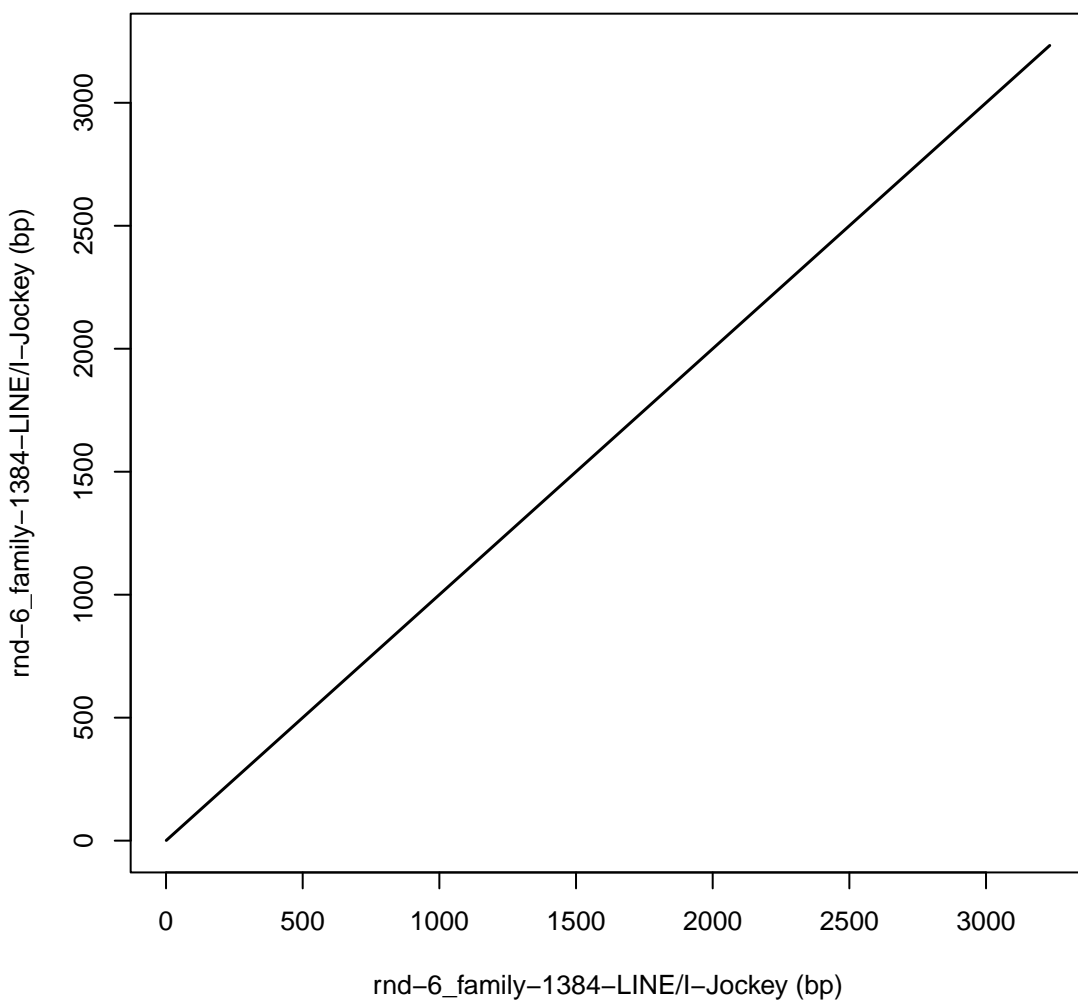
TE: rnd-6_family-1384-LINE/I-Jockey
consensus size: 3233bp; fragments: 263; full length: 1 (>=2909.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

