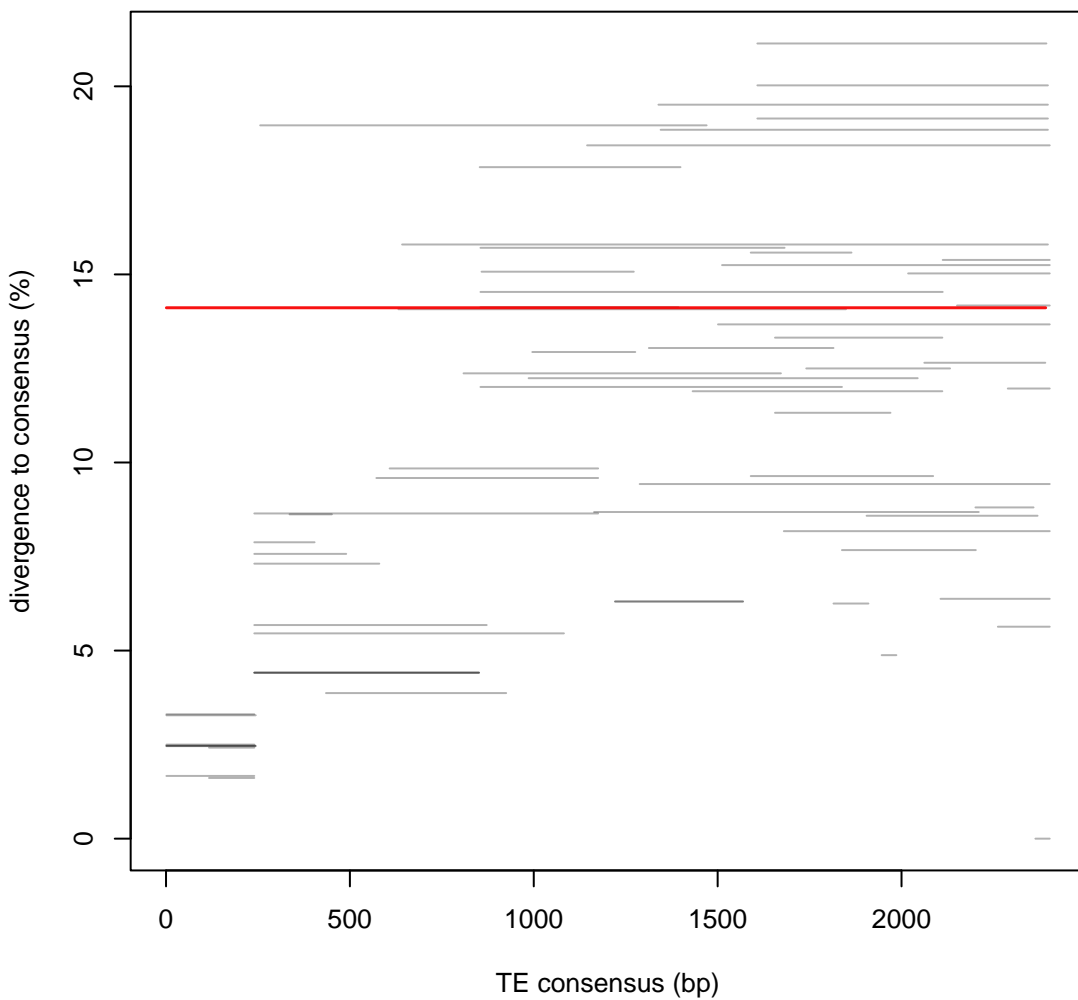
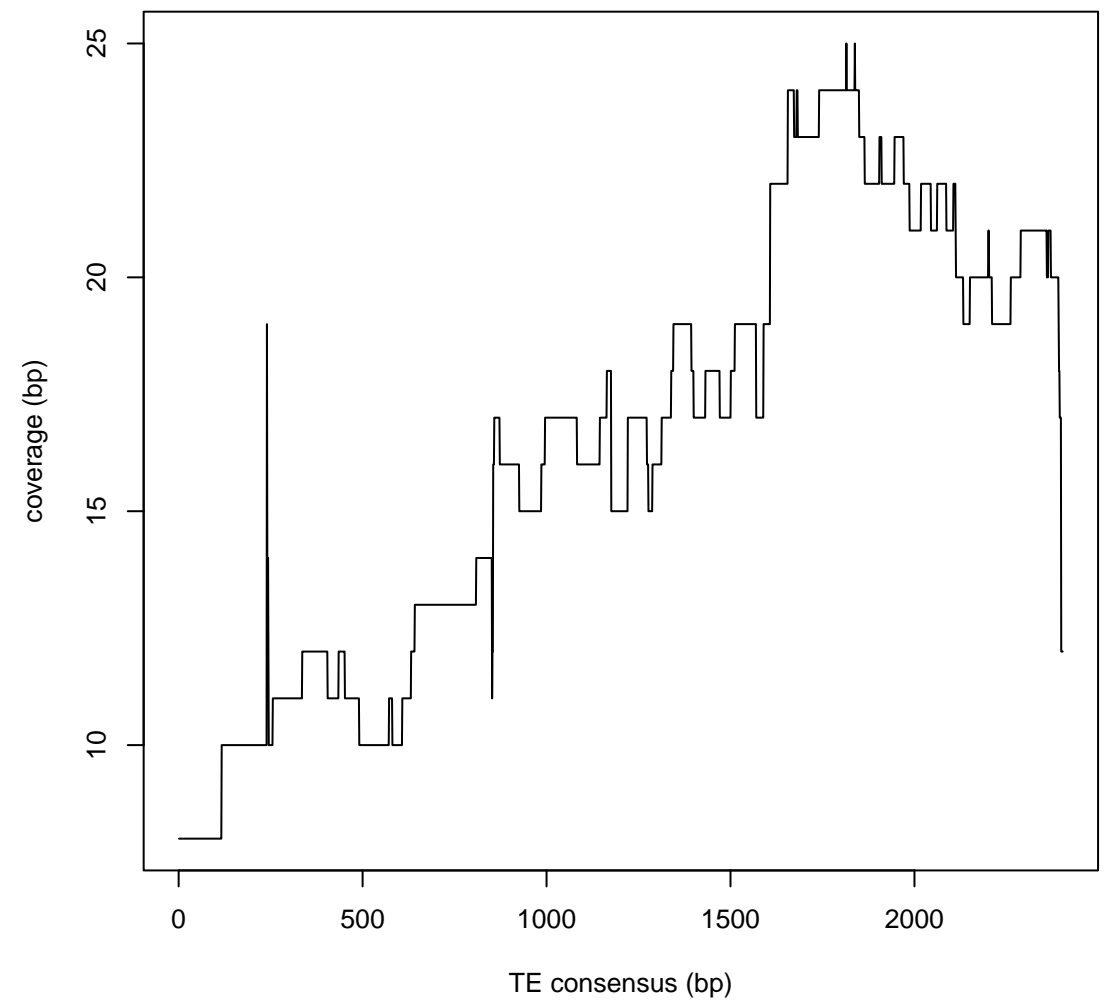


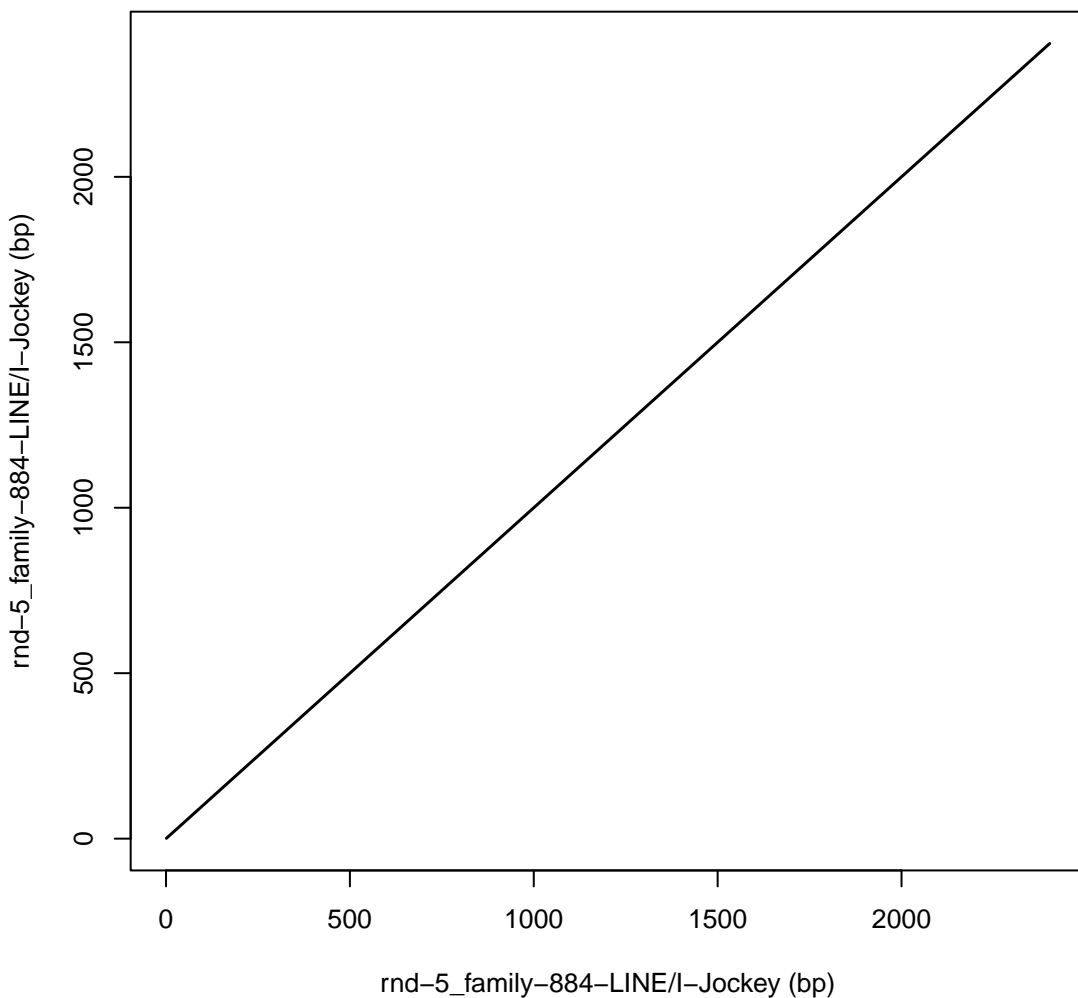
TE: rnd-5_family-884-LINE/I-Jockey
consensus size: 2403bp; fragments: 68; full length: 1 (>=2162.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

